

	phosphopeptide	protein
1	TSSTCSNESLSVGGTSVTPR202000000000000000	sp 060343 TBCD4_HUMAN
2	LQDSSSEEDVTEETDHR222000000000000000	sp Q76FK4 NOL8_HUMAN
3	SQSSEGVSSLSSSPNSLETQSQSLSR2200000000000000	sp 095155 UBE4B_HUMAN
4	EPVADEEEEDSDDDVPEITEFR2000000000000000	sp P54105 ICLN_HUMAN
5	GTGQSDSDIWDDTALIK200200000000000000	sp Q16637 SMN_HUMAN
6	GVDFESSEDDDDDPFMTSSLR220000000000000000	sp P49959 MRE11_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
7	GGSYSQAASSDSAQGSVSLTACK2E+20	sp P01892 1A02_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
8	LSEDEEEDEAEDDQSEASGK2. 2E+20	sp Q8NEJ9 NGDN_HUMAN
9	KLSSSDAPAQDTGSSAAAVETDASR2E+23	sp Q7Z4S6 KI21A_HUMAN
10	AEQSEEEEGEGEEEEEGGESK200000000000000000	sp Q8NE71 ABCF1_HUMAN
11	ASTASPCNNNINAATAVALQEPR2E+19	sp Q71RC2 LARP4_HUMAN
12	TSSDDESEDEDDLLQR200020000000000000	sp Q9Y5J1 UTP18_HUMAN
13	MESEGGADDSAEGLDLDNEDR200000000000000000	sp P07910 HNRPC_HUMAN
14	MLAESDESGDEESVSQTDK200200000000000000	sp P22059 OSBP1_HUMAN
15	SASSDTSEELNSQDSPPK200000000000000000	sp 014745 NHRF1_HUMAN
16	KRSSSEDDDLVDESDFDASINSYSVSDGSTR2. 2000000002E+30	sp Q14839 CHD4_HUMAN
17	LPSPTSPFSSLSQDQAATSK200000000000000000	sp Q8WWI1 LMO7_HUMAN
18	INSSGESGDESDFLQSR220020000000000000	sp 060841 IF2P_HUMAN
19	GTGQSDSDIWDDTALIK200000200000000000	sp Q16637 SMN_HUMAN
20	LAEDGESEPEAVGQSR200000000000000000	sp Q9UIG0 BAZ1B_HUMAN
21	HSLDSEEEEDDDGGSSK200000000000000000	sp 095400 CD2B2_HUMAN
22	GDVEGSQSQDEGEGSGESER2000000000000000	sp Q9UQE7 SMC3_HUMAN
23	SASPDDLGSNWAAADLGNEER2. 02E+23	sp 000193 SMAP_HUMAN
24	NENTEGSPQEDGVELEGLK2000000000000000	sp P11388 TOP2A_HUMAN
25	SPPGSGASTPVGPWDQAVQR2E+20	sp Q8TE77 SSH3_HUMAN
26	TVSQSFDGVS LDSSGPEDR200000000000000000	sp Q6BDS2 URFB1_HUMAN
27	SDAEDGGTVSQEEEDR200000000000000000	sp P27824 CALX_HUMAN
28	TADSSSEDEEEYVVEK222200000000000000	sp P45973 CBX5_HUMAN
29	KTSSDDESEDEDDLLQR200020000000000000	sp Q9Y5J1 UTP18_HUMAN
30	SEAAAPHTDAGGLSSDEEEGTSSQAEEAR2200000000000000	sp Q01831 XPC_HUMAN
31	LQELES CSGLGSTSDTDVR2000000000000000	sp Q14C86 GAPD1_HUMAN
32	MLAESDESGDEESVSQTDK1. 0002002E+19	sp P22059 OSBP1_HUMAN
33	ALDISLSSGEDEGDEEDSTAGTTK2. 2E+19	sp Q8IY81 SPB1_HUMAN
34	GDSESEDEDELEVPVPSR202000000000000000	sp P13861 KAP2_HUMAN
35	DSGSDDFLMEDDDSDYGSSK2. 02000001E+21	sp Q9H1E3 NUCKS_HUMAN
36	TSSTCSNESLSVGGTSVTPR220000000000000000	sp 060343 TBCD4_HUMAN
37	SPVFSDESDLDLDFDISK2000200000000000	sp 000566 MPP10_HUMAN
38	ELNEDDSTDCDDVQR220000000000000000	sp Q8WU17 RN139_HUMAN
39	AGGLQSDTEDECWSDTEAVPR202000002000000000	sp Q9H6H4 REEP4_HUMAN
		sp P05534 1A24_HUMAN
		sp Q09160 1A80_HUMAN
40	GGSYSQAASSDSAQGSVSLTACK2E+20	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
		sp Q6NXS1 IPP2M_HUMAN
41	IQQEESGGEEDSDLSPEER2200000000000000	sp P41236 IPP2_HUMAN
42	SSGSPYGGGYGSGGGSGGYGSR2E+21	sp P51991 ROA3_HUMAN

43	GSDSEDFEFEIQAEDDAR202000000000000000	sp Q8TDD1 DDX54_HUMAN
44	EELMSSDLEETAGSTSIPK2200000000000000	sp 000567 NOP56_HUMAN
45	GVDFESSEDDDDDPFMNTSSLR220000000010000000	sp P49959 MRE11_HUMAN
46	TPEELDDSDFETEDFDVR200000000000	sp P35221 CTNA1_HUMAN
47	LEDDSEESDAEFDEGFK20020000000000	sp Q03468 ERCC6_HUMAN
48	SSGSPYGGGYGSGGGSGGYGSR2E+19	sp P51991 ROA3_HUMAN
49	TVSQQSFDGVS LDSSGPEDR200200000000000000	sp Q6BDS2 URFB1_HUMAN
50	KTSSDDESEEDDLLQR220002000000000000	sp Q9Y5J1 UTP18_HUMAN
51	KAEQGSEEEGEGEEEEEGGESK200000000000000000	sp Q8NE71 ABCF1_HUMAN
52	FNDSEGDDTEETEDYR2000000000000000	sp Q9NYF8 BCLF1_HUMAN
53	SVASNQSEMEFSSLQDMPK2.002002E+19	sp Q8N163 CCAR2_HUMAN
54	EFSGPSTPTGTLEFEGGEVSLEGGK2E+23	sp Q09666 AHNK_HUMAN
55	SVAVSDEEEVEEEAER200020000000000000	sp Q9Y6X9 MORC2_HUMAN
56	VGGSDEEASGIPSR200000000000	sp Q9NQ55 SSF1_HUMAN
57	SASPDDDLGSSNWEAADLGNEER2E+21	sp 000193 SMAP_HUMAN
58	YLSADSGDADSDADLGS AVK2.002000002E+19	sp Q15361 TTF1_HUMAN
59	SQSPAASDCSSSSSASLPSSGR2E+21	sp 095817 BAG3_HUMAN
60	SQSSEGVSSLS SPSNSLETQS QLSLR220000000000000000	sp 095155 UBE4B_HUMAN
61	YLSADSGDADSDADLGS AVK2.002E+19	sp Q15361 TTF1_HUMAN
62	GLVAAYSGDSNNEELVER20020000000000	sp P52756 RBM5_HUMAN
63	FEEESKEPVADEEEEDSDDDV EPI TEFR20000000000000	sp P54105 ICLN_HUMAN
64	ALDSNSLENDL SAPGR200000000000	sp Q32MZ4 LRRF1_HUMAN
65	GGPEGVAAQAVASAASAGPADAEMEEIFDDASPGK20000	sp 060264 SMCA5_HUMAN
66	GTGQSDSDIWD TALIK200000000000000000	sp Q16637 SMN_HUMAN
67	SASPDDDLGSSNWEAADLGNEER2E+23	sp 000193 SMAP_HUMAN
68	EEASDDMMEGDEAVVR20001000000000	sp 075643 U520_HUMAN
69	SQASMV INSDT DDEEEVSAALTLAHLK2.02E+19	sp Q14676 MDC1_HUMAN
70	TASQEGCQELTQEQR2000000000000000	sp 043933 PEX1_HUMAN
71	GTLSDSEIETNSATSTIFGK200000000000000000	sp Q8WXG6 MADD_HUMAN
72	SLAALDALNTDDEENDEEEYEAWK2000000000000000	sp P55081 MFAP1_HUMAN
73	SEDEDSLEEAGSPAPGPCPR2.00002E+20	sp Q8TBB5 KLDC4_HUMAN
74	KLSSDAPAQDTGSSAAAVETDASR2E+21	sp Q7Z4S6 KI21A_HUMAN
75	SFNYS PNSTSEVSSTSASK200000000000000000	sp Q5VT52 RPRD2_HUMAN
76	SQSSDTEQQSPTS GGGK200000000000000000	sp P23588 IF4B_HUMAN
77	EEQTDTSDGESVTHHIR2022000000000000	sp P36915 GNL1_HUMAN
78	DESCYD NAEAAFS DDEEDLNSK200000000000	sp Q68E01 INT3_HUMAN
79	EGEELGEGERPEEDAAALELSSDEAVEVEEVIEESR220000000000000000	sp Q6NZI2 PTRF_HUMAN
80	SASSDTSEELNSQDSPPK200000000000000000	sp 014745 NHRF1_HUMAN
81	SSSVGSSSSYPI SPAVSR200000000000000000	sp Q15149 PLEC_HUMAN
82	SVASNQSEMEFSSLQDMPK200200000000000000	sp Q8N163 CCAR2_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
83	GGSYSQAASSDSAQGS DVSLTACK200000000000000000	sp P01892 1A02_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
84	MVIQGPSSPQGEAMVTDVLEDQK200000000000000000	sp Q12888 TP53B_HUMAN
85	QKSDAEEDGGTVS QEEEDR200000000000000000	sp P27824 CALX_HUMAN
86	NQGGYGGSSSSSYGSGR20000000	sp P09651 ROA1_HUMAN
87	GQESSDQE QVDVESIDFSK220000000000000000	sp Q96JM3 CHAP1_HUMAN
88	SPTMEQAVQTASAHL PAPA AVGR2.1E+21	sp Q8ND56 LS14A_HUMAN
89	NHSVNEEEQEEQEGESEDWEQV GPR20000000000000	sp Q14694 UBP10_HUMAN
90	LFDEEEDSSEKLFDDSDER2200000000000000	sp 043719 HTSF1_HUMAN

		sp P05534 1A24_HUMAN
		sp Q09160 1A80_HUMAN
91	GGSYSQAASSDSAQGSVDVSLTACK20000000000000	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
92	SVASNQSEMEFSSLQDMPK2.00200201E+19	sp Q8N163 CCAR2_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
93	LNQSGTSGVTDEESDVTQEEER2002000000000000	sp A6NKT7 RGPD3_HUMAN
		sp P0DJ0 RGPD1_HUMAN
		sp P0DJ1 RGPD2_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
		sp P01892 1A02_HUMAN
94	GGSYSQAASSDSAQGSVDVSLTACK2002000000	sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
95	SDAEDGGTVSQEEEDR200000000020000000	sp P27824 CALX_HUMAN
96	LLEDSEESSEETVSR200200000000	sp O60231 DHX16_HUMAN
97	EEENADSDDEGELQDLLSQDWR2000000000000000	sp Q96NB3 ZN830_HUMAN
98	DYEIESQNPLASPTNTLLGSAK200000000000	sp Q7Z417 NUFP2_HUMAN
99	EEQTDTSDGESVTHHIR2002000000000000	sp P36915 GNL1_HUMAN
100	QKSDAEEDGGTVSQEEEDR200000000020000000	sp P27824 CALX_HUMAN
101	VPVRPPQQYSDDDDYEDDEEDDVQNTNSALR2E+23	sp Q92560 BAP1_HUMAN
102	SQLQALHIGLDSSSIGSGPGDAEADDFPESR2.0002E+20	sp O75116 ROCK2_HUMAN
103	SSSEDAESLAPR220000000000	sp Q4G0J3 LARP7_HUMAN
104	DGGSGNSTIIVSR200000000000	sp Q7Z6Z7 HUWE1_HUMAN
105	AESPESAIESTQSTPQK2000000000000000	sp Q9NTI5 PDS5B_HUMAN
106	QSPNSSAAASVSR2000000000000000	sp Q9H1B7 I2BPL_HUMAN
107	KPVLPALTINPTIAEGSPSTSEGASEANLVDLQK2000000000000000	sp P36507 MP2K2_HUMAN
108	ETEEQSDSAEQGDPAGEGK2000000000000000	sp O95361 TRI16_HUMAN
109	TPLSSLSATSNNSTVACIDR20000000000000	sp P00533 EGFR_HUMAN
110	TGSNIGASSDISLDEQYK2.00002E+19	sp P22059 OSBP1_HUMAN
111	SSLGQSASETEEDTVSVSK2020000000000000	sp Q13435 SF3B2_HUMAN
112	TGSNIGASSDISLDEQYK200200000000000000	sp P22059 OSBP1_HUMAN
113	TAAGEYDSVSESEDEEMLEIR202020000100000	sp Q9BZE4 NOG1_HUMAN
114	SSSNDVDEETAESDTPVLEK200002002000000000	sp Q99590 SCAFB_HUMAN
115	AADSDDGAVSAPAASDGGVSK200000000000000000	sp Q96GM8 TOE1_HUMAN
116	SPSDSSTASTPVAEQIER2000000000000000	sp Q16643 DREB_HUMAN
117	LSEGSQPAEEEEEDQETPSR200200000000000000	sp Q9UKV3 ACINU_HUMAN
118	TPLSSLSATSNNSTVACIDR22000000000000	sp P00533 EGFR_HUMAN
119	AESSDGAESEEEEEAQEEVK2000200000000000	sp Q5T8D3 ACBD5_HUMAN
120	TASGSSVTSLDGTR2020000000000000	sp Q92597 NDRG1_HUMAN
121	SSSVGSSSSYPIPAVSR200200000000000000	sp Q15149 PLEC_HUMAN
122	NQGGYGGSSSSSYGSGR200000000000	sp P09651 ROA1_HUMAN
123	VVDYSQFQESDDADEDYGR200000000000	sp Q9H1E3 NUCKS_HUMAN
124	TINNQHVSVEQDNDQPDYDSVASDEDTDLETTASK200200000000000000	sp Q14161 GIT2_HUMAN
125	SSGSETEQVVDVSDR2200000000000000	sp O95251 KAT7_HUMAN
126	SVASNQSEMEFSSLQDMPK2.002E+19	sp Q8N163 CCAR2_HUMAN
127	TAAGEYDSVSESEDEEMLEIR2020200000000000	sp Q9BZE4 NOG1_HUMAN
128	SDSSGGYNLSDI IQSPSSTGLLK2E+20	sp Q9P2D0 IBTK_HUMAN
129	GQESSDQEQVDVESIDFSK220000000000000000	sp Q96JM3 CHAP1_HUMAN

130	NQGGYGGSSSSSYGSGR20000000000	sp P09651 ROA1_HUMAN
131	NSLLAGDDDTMSVISGISSR2002002000	sp Q8N3U4 STAG2_HUMAN
132	VTNDISPESSPGVGR20000000000	sp Q15154 PCM1_HUMAN
		sp Q641Q2 FA21A_HUMAN
133	LTDEDFSPFGSGGLFSGGK20000000000000	sp Q5SNT6 FA21B_HUMAN
		sp Q9Y4E1 FA21C_HUMAN
134	DWEDDSDEDMSNFDR20000000000	sp Q15185 TEBP_HUMAN
135	ALSSDSILSPAPDAR22000000000000	sp Q8IVT2 MISP_HUMAN
136	KAEDSDSEPEPEDNVR2020000000000	sp Q9H0D6 XRN2_HUMAN
137	HDSIPAADTFEDLSDVEGGGSEPTQR20000000000000	sp O95671 ASML_HUMAN
		sp P08238 HS90B_HUMAN
138	IEDVGSDEEDDSGKDK200000000000	sp Q58FF8 H90B2_HUMAN
		sp Q16643 DREB_HUMAN
139	SPSDSSTASTPVAEQIER20000000000000000000	sp Q8N163 CCAR2_HUMAN
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141	DLFDLNSSEEDDTGFSER2000000000000000	sp O00566 MPP10_HUMAN
142	KDDNDEEEEDIDFFEDIDSDEDEGGLFGSK20000000000000	sp Q99543 DNJC2_HUMAN
143	ELSEESEDEELQLEEFMLK20020000000000000000	sp P51114 FXR1_HUMAN
144	KDELSDWSLAGEDDR20020000000000	sp Q6PKG0 LARP1_HUMAN
145	NTFTAWSDEESDYEIDDR2000000000000000	sp P23588 IF4B_HUMAN
146	TGSESSQTGTSTTSSR22000000000000	sp Q9UHB6 LIMA1_HUMAN
147	EGHSLEMENENLVENGADSDEDDNSFLK200000000000	sp O75448 MED24_HUMAN
148	LLSSNEDDANILSSPTDR200000000000000000	sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
		sp P01892 1A02_HUMAN
149	GGSYSQAASSDSAQGSVSLTACK2000000000000000	sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
150	EELMSSDLEETAGSTSIPK12200000000000000000	sp O00567 NOP56_HUMAN
151	DWEDDSDEDMSNFDR20001000000	sp Q15185 TEBP_HUMAN
152	AFVEDSEDEDGAGEGSSLLQK20000000000000000000	sp Q8N9T8 KRI1_HUMAN
153	LPSGGAASPTGSAVDIR20000020000000000000	sp Q09666 AHNK_HUMAN
154	DYLLSESEDEGDNDGER2020000000000000	sp Q9BVJ6 UT14A_HUMAN
155	ESPRPLQLPGAEGPAISDGEEGGEPGAGGGAAGAAGAGR2E+39	sp Q6PKG0 LARP1_HUMAN
156	SSSPAPADIAQTVQEDLR22000000000000000000	sp Q13283 G3BP1_HUMAN
157	GESAADSDGWSAPSDLR20000200000000	sp Q9BUL5 PHF23_HUMAN
158	VDSEGDSENDDAAGDFR20000200000000000000	sp O43823 AKAP8_HUMAN
159	FASDDEHDEHDENGATGPVK20000000000000000000	sp P05455 LA_HUMAN
160	AQSTDSLGTSGSLQSK20000000000000000000	sp Q15276 RABE1_HUMAN
161	YGLLAN TEDPTEMASLDSDEETVFESR2002000000000000	sp Q9BVV8 CSO24_HUMAN
162	TPAAAAAMNLASPR2000	sp Q9UQ35 SRRM2_HUMAN
163	EEQTDTSDGESVTHHIR2200000000000000	sp P36915 GNL1_HUMAN
164	FIIGSVSEDNSEDEISNLVK20000000000000	sp Q13085 ACACA_HUMAN
165	NWEDEDFYSDDDTFLDR20000000000000	sp Q9BWU0 NADAP_HUMAN
166	LSEGSQPAEEEEEDQETPSR20000000000000000000	sp Q9UKV3 ACINU_HUMAN
167	SSSPAPADIAQTVQEDLR20000000000000000000	sp Q13283 G3BP1_HUMAN
		sp E9PAV3 NACAM_HUMAN
168	VQGEAVSNIQENTQTPTVQEESEEEVEDETGVEVK200000000000000000	sp Q13765 NACA_HUMAN
		sp Q9Y6R9 CCD61_HUMAN
169	LGSGGSGDGPSVSWSR20020000000000000000	sp Q3YEC7 RABL6_HUMAN
170	DDPSDV TDEDEGPAEPPPPPK20020000000000000000	sp Q8ND56 LS14A_HUMAN
171	SPTMEQAVQTASAHL PAPA AVGR2E+21	sp O60841 IF2P_HUMAN
172	KQSFDDNDSEELDKDSK20000020000000000000	sp Q8N163 CCAR2_HUMAN
173	SVASNQSEMEFSSLQDMPK2.00200201E+19	sp Q6KC79 NIPBL_HUMAN
174	YAEISSDEDNDSDEAFESSR22000000000000000000	

175	TSDSDQQAYLVQR200000000000	sp Q96CV9 OPTN_HUMAN
176	DSGSEDFLMEDDDSDYGSSK2.02E+21	sp Q9H1E3 NUCKS_HUMAN
177	SSGSETEQVVFSDR20200000000000	sp 095251 KAT7_HUMAN
178	GTLDEEDEEADSDTDDIDHR2000000000	sp Q9HCN4 GPN1_HUMAN
179	DSSGQEDETQSSNDDPSQSVASQDAQEIIRPR2E+30	sp Q8N108 MIER1_HUMAN
180	GLVAAYSGSDNNEELVER200000000000	sp P52756 RBM5_HUMAN
181	SPSVSPSKQPVSTSSK20200000000000	sp Q96JG6 CC132_HUMAN
182	LLSSNEDDANILSSPTDR2000000000000000	sp 075448 MED24_HUMAN
183	VAGEAAETDSEPEPEPEPTAAPR2020000000000000	sp Q6QNY0 BL1S3_HUMAN
184	SSGSETEQVVFSDR2202000000000000	sp 095251 KAT7_HUMAN
185	GTLSDSEIETNSATSTIFGK2000000000000000	sp Q8WXG6 MADD_HUMAN
186	SSSVGSSSSYPISPAISR220020000000000000	sp Q15149 PLEC_HUMAN
187	TGVTSTSDSEEGDDQEGEK2020000000000000	sp 075475 PSIP1_HUMAN
188	DDSPDEVSTFTVQPPPSR200000000000000000	sp Q8N328 PGBD3_HUMAN
189	SPSPGNHTSNSSNASNATVVPQNSSAR2.02E+28	sp Q9BTA9 WAC_HUMAN
190	APSVANVGSCHDLK2000000000000000	sp P21333 FLNA_HUMAN
191	STGSATSLASQGER200000000000	sp Q5SW79 CE170_HUMAN
192	TGSESSQTGTSTTSSR2000000000000000	sp P23588 IF4B_HUMAN
193	LNSGGGLSEELGSAR2000000000000000	sp Q96KQ7 EHMT2_HUMAN
194	SSGTASSVAFTPLQGLEIVNPQAAEK2.0000002E+23	sp Q8WWY3 PRP31_HUMAN
195	KPATPAEDDEDDIDLFGSDNEEEDKEAQLR2000000000000000	sp P29692 EF1D_HUMAN
196	IQQESSGEEDSDLSPEER2200000000000000	sp P41236 IPP2_HUMAN
197	EESSEDENEVSNILR2200000000000000	sp Q6NXS1 IPP2M_HUMAN
198	SLDSDESEDEEDDYQK2002000000000000	sp Q01804 OTUD4_HUMAN
199	FSDSEGEETVPEPR2020000000000000	sp Q13442 HAP28_HUMAN
200	SPGSPVGEFTGSPPK2000000000000000	sp Q13286 CLN3_HUMAN
201	LKSEDGVEGDLGETQSR2000000000000000	sp P35611 ADDA_HUMAN
202	SPTPPSSAGLGSNSAPPIDSR2.02E+22	sp Q09666 AHNK_HUMAN
203	FSLCSDNLEGISEGPSNR200200000000000000	sp Q8IWX8 CHERP_HUMAN
204	NSLLAGGDDTMSVISGISSR2002000	sp Q14C86 GAPD1_HUMAN
205	DSHSSEDEASSQTDLSQTISK2.2E+19	sp Q8N3U4 STAG2_HUMAN
206	AADSSSDGAEESPAEPGAPR220200000000000000	sp Q5JTV8 TOIP1_HUMAN
207	GQVIIISDSDDDDDER20200000000	sp P16383 GCFC2_HUMAN
208	NPWSDDESKSESLEETEPVVIPR2.00020202E+21	sp Q7Z333 SETX_HUMAN
209	DDDDIDLFGSDDEESEEAK200000000000	sp Q02880 TOP2B_HUMAN
210	VAAETQSPSLFGSTK2000000000	sp P24534 EF1B_HUMAN
211	SPVGLSSDGISSSSSSSR200000000000000000	sp Q9UKX7 NUP50_HUMAN
212	AGGASPAASSTAQPPTQHR2000000000000000	sp 060583 CCNT2_HUMAN
213	KPATPAEDDEDDIDLFGSDNEEEDK2E+23	sp Q8IU81 I2BP1_HUMAN
214	FTMDLSDSDFDFDEK1000200002000000	sp P29692 EF1D_HUMAN
215	YSQGDDGSSSSGSSVAGSQSTLTK2E+25	sp P11388 TOP2A_HUMAN
216	GTGSGGQLQDLDCSSDDEGAAQNSTKPSATK2.0000000022E+29	sp Q9UH99 SUN2_HUMAN
217	EKEISDDEAEEKGEK200000000000	sp P08240 SRPR_HUMAN
218	NKPGPNIESGNEDDASFK200000000000	sp P08238 HS90B_HUMAN
219	MPCESPPEADTPTSTR2000000000000000	sp 060841 IF2P_HUMAN
220	TGSESSQTGTSTTSSR2000000000000000	sp P46013 KI67_HUMAN
221	TVSSPPTSPRPGSAATVSASTSNIIPR2.2E+22	sp P23588 IF4B_HUMAN
222	LAAAETAVSPR2000	sp 076094 SRP72_HUMAN
223	IEDVGSDEEDDSGK2000000000	sp Q9BUH6 PAXX_HUMAN
224	ESSANNSVSPESLR20000000	sp P08238 HS90B_HUMAN
225	LSVPTSDEEDEVPAKPR2000000000000000	sp Q58FF8 H90B2_HUMAN
226	EESSEEEDEDEEEEEEEK2000000000000000	sp Q04726 TLE3_HUMAN
		sp Q8NE71 ABCF1_HUMAN
		sp P35659 DEK_HUMAN

		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
227	LNQSGTSVGTDEESDVTQEEER220020000000000000	sp A6NKT7 RGPD3_HUMAN
		sp P0DJD0 RGPD1_HUMAN
		sp P0DJD1 RGPD2_HUMAN
228	EGHSLEMENENLVENGADSDEDDNSFLK1.00000000002E+22	sp Q9UHB6 LIMA1_HUMAN
229	DNLLDTYSADQGDSSEGGTLAR20000000000	sp Q6ZRP7 QSOX2_HUMAN
		sp Q641Q2 FA21A_HUMAN
230	TPSDDEEDNLFAPPK2020000000000000	sp Q9Y4E1 FA21C_HUMAN
		sp Q14745 NHRF1_HUMAN
231	SASSDTSEELNSQDSPPK2022000000000000	sp P35251 RFC1_HUMAN
232	I IYDSDESEETLQVK2020000000000	sp P00264 PGRC1_HUMAN
233	GDQPAASGSDDDDEPPPLPR200000000000	sp Q9Y4E1 FA21C_HUMAN
234	GLFSDEEDSEDLFSSQSASNLK2E+19	sp P49768 PSN1_HUMAN
235	AAVQELSSSILAGEDPEER2200000000000	sp Q14134 TRI29_HUMAN
236	SKSGSEEVLCDSICGNK2000000000000000	sp Q8IYB3 SRRM1_HUMAN
237	KETESAEADNLDLEK2020000000000000	sp P54725 RD23A_HUMAN
238	EDKSPSEESAPTTSPESVSGSVPSSGSSGR2E+27	sp Q02447 SP3_HUMAN
239	IGPPSPGDDEEEAAAAAGAPAAAGATGLASALGAPNR2E+36	sp P46937 YAP1_HUMAN
240	QASTDAGTAGALTPQHVR2000000000000000	sp Q9Y3B9 RRP15_HUMAN
241	DHFYSDDAIEADSEGAEPCK2.00000002E+19	sp Q08J23 NSUN2_HUMAN
242	ESTQLSPADLTEGKPTDPSK2000000000000000	sp Q9UK58 CCNL1_HUMAN
243	LSPTPSMQDGLDLPSETDLR2E+19	sp P29692 EF1D_HUMAN
244	KPATPAEDDEDDIDILFGSDNEEDK200000000	sp P02545 LMNA_HUMAN
245	SVGGSGGSGFDNLVTR200000000000000000	sp Q9Y4C8 RBM19_HUMAN
246	SKPASDYLNFDSDSGQSEEEGAGEDLEEEASLEPK2.020002E+25	sp Q12888 TP53B_HUMAN
247	NSPEDLGLSLTGDSCK2000000000000000	sp P46100 ATRX_HUMAN
248	RPTETNPVTSNSDEECNETVK20000000000	sp Q13442 HAP28_HUMAN
249	SLDSDESEDEEDDYQQK200000200000000000	sp Q8N163 CCAR2_HUMAN
250	SVASNQSEMEFSSLQDMPK20000100000000000	sp P51858 HDGF_HUMAN
251	KGNAEGSSDEEGKLVIDEPAK2200000000000000	sp Q12888 TP53B_HUMAN
252	MVIQGPSSPQGEAMVTDVLEDQK1.0000002E+23	sp Q32MZ4 LRRF1_HUMAN
253	ALDSNSLENDLAPGR2000000000000000	sp Q9H1E3 NUCKS_HUMAN
254	DSGSDDFLMEDDDSDYGSSK2.02000001000002E+21	sp Q5SW79 CE170_HUMAN
255	STGSATSLASQGER2020000000000000	sp Q14134 TRI29_HUMAN
256	SKSGSEEVLCDSICGNK200000000000000000	sp Q9P2I0 CPSF2_HUMAN
257	EADIDSSDESIEEDIDQPSAHK220020000000000000	sp Q7Z6Z7 HUWE1_HUMAN
258	AGSSTPGDAPPAVAEVQGR22000000000000000	sp P05783 K1C18_HUMAN
259	LLEDGEDFNLGDALDSSNSMQTIQK2000000000	sp Q9BPX3 CND3_HUMAN
260	TLHCEGTEINSDEQESK200000000	sp Q9UHW9 S12A6_HUMAN
261	LTSIGSDEDEETETYQEK200020000000000000	sp Q4G0J3 LARP7_HUMAN
262	SSSEDAESLAPR2200000000000000	sp Q13283 G3BP1_HUMAN
263	SSSPAPADIAQTVQEDLR200000000000000000	sp Q9NTJ3 SMC4_HUMAN
264	REEGPPPPSPDGASSDAEPEPPSGR200002000000000000	sp Q14976 GAK_HUMAN
265	DESEVSDEGGSPISSEGEPR2.002E+19	sp Q3YEC7 RABL6_HUMAN
266	AAQQDSDSGEALGGNPMVAGFQDDVDLEDQPR2.02E+28	sp P50750 CDK9_HUMAN
267	KGSQITQQSTNQRSR2000000000000000	sp Q8N163 CCAR2_HUMAN
268	SVASNQSEMEFSSLQDMPK200000000000000000	sp Q9Y5J1 UTP18_HUMAN
269	KTSSDDESEEDDLLQR200002000000000000	sp Q8N163 CCAR2_HUMAN
270	SVASNQSEMEFSSLQDMPK2.00200001E+19	sp Q15021 CND1_HUMAN
271	YQPLASTASDNDFVTPEPR2000000000000	sp Q96ST2 IWS1_HUMAN
272	NQASDSENEELPKPR2020000000000	sp Q75475 PSIP1_HUMAN
273	TGVTSTSDSEEGDDQEGEK2.00000202E+20	sp P13861 KAP2_HUMAN
274	VADAKGDSSEEDLEVPVPSR202000000000000000	sp Q9UQR1 ZN148_HUMAN
275	QPLEQNQTI SPLSTYEESK2000000000000	

276	VENSPQVDGSPPGLEGLGGIGEK2000000000000000	sp Q8WVB6	CTF18_HUMAN
277	QIDSSPVGGETDETTVSQNYR200000000000000000	sp O15027	SC16A_HUMAN
278	VALSDDETKETENMR20000000000000	sp Q15054	DPOD3_HUMAN
279	TINNQHVESQDNDQPDYDSVASDEDTDLETTASK30000200000000000000	sp Q14161	GIT2_HUMAN
280	ILQSQDFSLDSSAEEGLR2200000000	sp Q5T8I3	F102B_HUMAN
281	YYSDSDELTVQR20200000000000	sp Q8NFC6	BD1L1_HUMAN
282	TGSNISGASSDISLDEQYK20020020000000000000	sp P22059	OSBP1_HUMAN
283	GTGSGGQLQDLDCSSSDDEGAAQNSTKPSATK2.0000000002E+29	sp P08240	SRPR_HUMAN
284	FIIGSVSEDNSEDEISNLVK20000020000000000000	sp Q13085	ACACA_HUMAN
285	APEPLGPADQSELGPEQLEAEVGESSDEEPVESR220000000000	sp Q969G5	PRDBP_HUMAN
286	TEFLDLNSPLSPSPR2000	sp Q8NCF5	NF2IP_HUMAN
287	ASSHSSQTQGGGSVTK20000000000000000000	sp P02545	LMNA_HUMAN
288	GDSIEEILADSEDEEDNEEEER2000000000000000	sp Q5JTH9	RRP12_HUMAN
289	TSSGTSLSAMHSSGSSGK20000000000000000000	sp Q12888	TP53B_HUMAN
290	GLFSDEEDSEDLFSSQSASK20000000000000000000	sp Q641Q2	FA21A_HUMAN
		sp Q5SNT6	FA21B_HUMAN
291	VDENENCSSLGSPSEPPQTLDLVR2000000000000000	sp Q96P16	RPR1A_HUMAN
292	SLLSHEFQDETDTTEETLYSSK20200000000000	sp O75976	CBPD_HUMAN
293	LLEDSEESSEETVSR200220000000	sp O60231	DHX16_HUMAN
294	SPPSTGSTYGSSQK2000000000000000	sp Q9Y2W1	TR150_HUMAN
295	NTFTAWSDEESDYEIDDR20002000000000	sp Q6PKG0	LARP1_HUMAN
296	GGPEGVAAQAVASAASAGPADAEMEEIFDDASPGK1000000020000	sp O60264	SMCA5_HUMAN
297	TSSTCSNESLSVGGTSVTPR2.2E+19	sp O60343	TBCD4_HUMAN
298	TPLSSLSATSNNSTVACIDR20000000000000	sp P00533	EGFR_HUMAN
299	TSSGTSLSAMHSSGSSGK20000000000000000000	sp Q12888	TP53B_HUMAN
300	LPSPTSPFSSLSQDQAATSK22000000000000000000	sp Q8WWI1	LMO7_HUMAN
301	ATGGLCLLGAYADSDDDNDVSEK20000000000000	sp Q8N3X1	FNBP4_HUMAN
302	AETSEGSAPSAPVPEASAPK2000	sp P51608	MECP2_HUMAN
303	MESEGGADDSAEGLDLLDDDDNEDRGDDQLELIK1.000000002E+34	sp P07910	HNRPC_HUMAN
304	SQLQALHIGLDSSSIGSGPGAADDFPESR2.002E+19	sp O75116	ROCK2_HUMAN
305	INSSGESGDESDEFLQSR220000002000000000	sp O60841	IF2P_HUMAN
306	GGGSCSGSAGGGGSGSLPSQR20000000000000000000	sp O15234	CASC3_HUMAN
307	FTMDLSDSDFSDFDKTDDEDFVPSDASPPK1.000200002E+30	sp P11388	TOP2A_HUMAN
308	MQSLQLDCVAVPSSR2000000000000000	sp Q13546	RIPK1_HUMAN
309	REEGPPPPSPDGASSDAEPEPPSGR20000000000000	sp Q9NTJ3	SMC4_HUMAN
310	MNEEISSDSESESLAPR22020000000000	sp O43818	U3IP2_HUMAN
311	SPVGKSPSTGSTYGSSQK2.00002E+19	sp Q9Y2W1	TR150_HUMAN
312	SPAGPAATPAQAQAASTPR2E+19	sp Q13428	TCOF_HUMAN
313	SSLGQSASETEEDTVSVSK2000000000000000	sp Q13435	SF3B2_HUMAN
314	TTSSANNPNLMYQDECDR20000000000000000000	sp Q92841	DDX17_HUMAN
315	SIGSAVDQGNESIVAK20000000000000000000	sp Q9H0H5	RGAP1_HUMAN
316	IEDVGSDEEDDSGKDKK20000000000000	sp P08238	HS90B_HUMAN
		sp Q58FF8	H90B2_HUMAN
		sp Q6PEY2	TBA3E_HUMAN
		sp Q9BQE3	TBA1C_HUMAN
317	TIGGGDDSFNTFFSETGAGK2000000000000000	sp Q71U36	TBA1A_HUMAN
		sp Q13748	TBA3C_HUMAN
		sp P68363	TBA1B_HUMAN
318	EEASDDDMEGDEAVVR2000000000000000	sp O75643	U520_HUMAN
319	TSAAACAVTDLSDSDDFDEK2002000000	sp Q9NQZ2	SAS10_HUMAN
320	TASISSPSEGTPTVGSYGCTPQSLPK2E+21	sp Q6PKG0	LARP1_HUMAN
321	INSSGESGDESDEFLQSR220020002000000000	sp O60841	IF2P_HUMAN
322	GQDTVAIEGFTDEEDTESGGEGQYR2000000200000000	sp Q2KHR3	QSER1_HUMAN

323	EAESESSGGEEEDENIEVVYSK2. 2E+19	sp P55010 IF5_HUMAN
324	SAGEEEDGPVLTDEQK2000000000000000	sp Q66PJ3 AR6P4_HUMAN
325	NSQEDSEDSKDKVK20000000000	sp Q9H1E3 NUCKS_HUMAN
326	TGVTSTSDSEEGDDQEGEK2202000000000000	sp 075475 PSIP1_HUMAN
327	TVSSPPTSPRPGSAATVSASTSNIIPPR2. 00002E+21	sp 076094 SRP72_HUMAN
328	EETESSEGEAAAGGGAK202200000000000000	sp Q96G23 CERS2_HUMAN
329	GTGQSDSDIWDITALIK2000000000000000	sp Q16637 SMN_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
330	GGSYSQAASSDSAQGSVSLTACK2002000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
331	CGSGPVHISGQHLVAVEEDAESDEEEEDVK20000000000	sp P06748 NPM_HUMAN
332	NEGSESAPEGQAQQR2020000000000000	sp P67809 YBOX1_HUMAN
333	MADHLEGLSSDDEETSTDITNFNLEK220000000000000000	sp Q9Y5B6 PAXB1_HUMAN
334	REEGPPPPSPDGASSDAEPEPPSGR20000000000000	sp Q9NTJ3 SMC4_HUMAN
335	KVEEDLKADEPSSEESDLEIDK200200000000	sp P50502 F10A1_HUMAN
336	ENEEDGTGNWYSSDEDEGGSSVTSILK220000000000000000	sp Q9UPT8 ZC3H4_HUMAN
337	NLEDDSDDEEDFFLR20000000000	sp 075379 VAMP4_HUMAN
338	YSEFTSTSGTGHNQTR20000000000	sp P48960 CD97_HUMAN
339	SGSPAPETTNEVPFAQHSSLSR2. 02E+24	sp O15047 SET1A_HUMAN
340	NEGSESAPEGQAQQR2000000000000000	sp P67809 YBOX1_HUMAN
341	SSVAAPEKSSNDSVDEETAESDTSPVLEK2. 2002E+21	sp Q99590 SCAFB_HUMAN
342	NFSCGSVEFDEADLESRLR20020000000000000000	sp Q86SQ0 PHLB2_HUMAN
343	SEAEDDEDYVPYVPLR20000000000000000000	sp Q9UJV9 DDX41_HUMAN
344	FNSESESGSEASSPDYFGPPAK2. 0202002E+20	sp Q9BW71 HIRP3_HUMAN
345	TGSTSSKEDDYESDAATIVQK2E+21	sp Q5T4S7 UBR4_HUMAN
346	LGAGEGGEASVSPEK20000	sp Q13428 TCOF_HUMAN
347	SESLIDASEDSQLEAAIR200000002000000000	sp O94888 UBXN7_HUMAN
348	KPPAPPSPVQSQSPSTNWSPAVPVK2. 000002E+19	sp Q8IYB3 SRRM1_HUMAN
349	EVLYDSEGLSGEER2000200000	sp Q9H7N4 SFR19_HUMAN
350	SSSISEEKGSDDEKPR200000002000000000	sp Q9UKV3 ACINU_HUMAN
351	TESPATAAETASEELDNR20000000000000000000	sp Q9NTJ3 SMC4_HUMAN
352	KVEEDLKADEPSSEESDLEIDK2200200000000	sp P50502 F10A1_HUMAN
353	AEDSDSEPEPEDNVR2020000000000000	sp Q9H0D6 XRN2_HUMAN
354	SPEPEATLTFPFLDK20000000000000000000	sp Q9UPQ0 LIMC1_HUMAN
355	NVLSDSVSSDDEFFDAR220000000000	sp Q9BZ71 PITM3_HUMAN
356	GPSPSPPTPPAAAAPAEQAPR22000000000000000000	sp Q86VQ1 GLC11_HUMAN
357	SDGAPASDSKPGSSEAAPSSK2E+21	sp P80723 BASP1_HUMAN
358	SESTDLSGLSPSEVTAIQCK20200000000000000000	sp O60318 GANP_HUMAN
359	SVTSNQSDGTQESCESPDVLR20000000000000000000	sp Q9UGV2 NDRG3_HUMAN
360	DIEVLSEDTDYEEDEVTK2002000000000000	sp Q4LE39 ARI4B_HUMAN
361	LEGSDVDSELEDR200020000000	sp Q92614 MY18A_HUMAN
362	DSELSDTDSGCCLGQSESDK20000000000000000000	sp Q96T88 UHRF1_HUMAN
363	TQQLHPVLQKPSVFGNDSDDDETSVSESLQR20000000000000000000	sp Q9H0G5 NSRP1_HUMAN
364	EYDSGDTDEIIAMK20020000000000	sp Q76FK4 NOL8_HUMAN
365	EEGSSDEISSGVGSESEGLNSPVK2. 200022E+22	sp P52701 MSH6_HUMAN
366	SSQSEGIFLGESEDESVR202000000000	sp Q9BYW2 SETD2_HUMAN
367	YEDEVFGDSEPELDEESEDEVEEQEER2. 00000002E+20	sp Q9UN86 G3BP2_HUMAN
368	VEEQEADEEDVSEEAESK2000000000	sp Q9H3N1 TMX1_HUMAN
369	FTDKDQPSGSEGEDDDAEALK20200000000000000000	sp Q9NXG2 THUM1_HUMAN
370	KQSSSEISLAVER2000000000000000	sp P48634 PRC2A_HUMAN
371	LPSTSDDCPAIGTPLR20000000000000000000	sp Q15648 MED1_HUMAN



372	IQQFDDGGSDDEEDIWEEK2000000000	sp Q5H9R7 PP6R3_HUMAN
373	GSPSGGSTAEASDTLSIR200000000000000000	sp Q9P206 K1522_HUMAN
374	SDSSGGYNLSDIIQSPSSTGLLK2. 2E+21	sp Q9P2D0 IBTK_HUMAN
375	ENSGPVENGVSdqEGEEQAR20000000000	sp Q8TEQ6 GEMI5_HUMAN
376	GDQVLNFSDAEDLIDDSK200000000000	sp Q96EZ8 MCRS1_HUMAN
377	IVEPEVVGESDSEVEGDWR202000000000	sp P55081 MFAP1_HUMAN
378	KSPVFSDESDLDLFDISK200020002000000000	sp 000566 MPP10_HUMAN
379	SLSNSNPDISGTPTSPDDEV2. 02E+19	sp Q96N67 DOCK7_HUMAN
380	KTSAVSSPLLDQQR20002000000000	sp Q5T200 ZC3HD_HUMAN
381	FNDSEGDDTEETEDYR200002000000000	sp Q9NYF8 BCLF1_HUMAN
382	KGSSGNASEVSVACLTER200000000000000000	sp Q69YQ0 CYTSA_HUMAN
383	SLSNSNPDISGTPTSPDDEV200000000000000000	sp Q96N67 DOCK7_HUMAN
384	EEVGAGYNSEDEYEAAR200000000000	sp Q96G74 OTUD5_HUMAN
385	KTSSDDESEEDDLLQR20200020000000000000	sp Q9Y5J1 UTP18_HUMAN
386	SPTAALNESLVECPK2000000000000000	sp Q53EZ4 CEP55_HUMAN
387	VLNTGSDVEEAVADALK20000000000000	sp 075122 CLAP2_HUMAN
388	SDSFENPVLQQR2000000000000000	sp P12956 XRCC6_HUMAN
389	GNAEGSSDEEGKLVIDEPK220000000000000000	sp P51858 HDGF_HUMAN
390	KSSADTEFSDECTTAER220000000000000000	sp Q9H6S0 YTDC2_HUMAN
391	DELNEELIQEESSEDEGEYEEVR22000000000000	sp 060524 NEMF_HUMAN
392	YQDEVFGGFVTEPQEESEEEVEEPEER200000000000	sp Q13283 G3BP1_HUMAN
393	LSSSATLLNSPDR20000	sp Q14244 MAP7_HUMAN
394	TELNSSAESEQPLDK2200000000000000	sp Q9Y5B6 PAXB1_HUMAN
395	IGDEYAEDSSDEEDIR220000000	sp Q14137 BOP1_HUMAN
396	DEEQSEADAGSGPPTPGPTTLGPK2000200000000000	sp Q6ZRS2 SRCAP_HUMAN
397	EKPDSDDLDIASLVTA200000000000000000	sp Q86V48 LUZP1_HUMAN
398	ELHENVLDKELEENDSENSEFEDDGSEK20020000002000	sp 043719 HTSF1_HUMAN
399	SISLGDSEGPVATLAQPLR20000000000000000000	sp 043379 WDR62_HUMAN
400	ANSGVDLDSSGEFASIEK2000000220000000000	sp Q92766 RREB1_HUMAN
401	SDVQEESEGSDDDNKDSAAFEDNEVQDEFLEK2. 000002E+22	sp Q9UIG0 BAZ1B_HUMAN
402	GSPDGLQTKGPSAPK20000000000000000000	sp Q8IZL8 PELP1_HUMAN
403	ALSSDSILSPAPDAR20000200000000	sp Q8IVT2 MISP_HUMAN
404	AESSDSGAEESEEEEAQEEVK22020002000000000000	sp Q5T8D3 ACBD5_HUMAN
405	SIQDLTVTGTEPGQVSSR20000000000000000000	sp 043318 M3K7_HUMAN
406	MNEISSDSESESLAPR1000022020000000000	sp 043818 U3IP2_HUMAN
407	EDSSEEEEEIDDEEIER22000000000000000000	sp P55081 MFAP1_HUMAN
408	SGSEEEAELSLSLAR20200000000000000000	sp Q8IV50 LYSM2_HUMAN
409	DSGSDDFLMEDDDSDYGSSK2. 02000000000002E+21	sp Q9H1E3 NUCKS_HUMAN
410	YESQLSTNEEKVDTDDR2000000000000000	sp Q13433 S39A6_HUMAN
411	TSFVSGSDDELGPIR20020000000000	sp Q9NRY4 RHG35_HUMAN
412	GLVAAYSGESDSEEEQER2020000000	sp P98175 RBM10_HUMAN
413	TADSSSEDEEEYVVEK20022020000000000000	sp P45973 CBX5_HUMAN
414	QHYSSEDEPDDNPVLDLDR22000000000000000000	sp Q8NFC6 BD1L1_HUMAN
415	SDAEEDGGTVSQEEEDR200000000	sp P27824 CALX_HUMAN
416	GQESSDQEQVDVESIDFSK22200000000000000000	sp Q96JM3 CHAP1_HUMAN
417	LSVPTSDEEDEVPAKPR22000000000000000000	sp Q8NE71 ABCF1_HUMAN
418	EESDGEYDEFGR200000000000	sp 095218 ZRAB2_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
419	KGGYSQAASSDSAQSDVSLTACKV20020000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
420	IACEEFSDEEEGEGGR202000000000	sp Q13547 HDAC1_HUMAN

421	KGSVSHDTVQPR20000000000	sp Q96D71 REPS1_HUMAN
422	ENYSDSEEDDDVASSR2020000000000000	sp Q9Y2U8 MAN1_HUMAN
423	YDSGDGKSDDLVVDVSNEDPATPR2.00002E+22	sp Q04726 TLE3_HUMAN
424	SPVFSDESDLDFDISK200020002000000000	sp 000566 MPP10_HUMAN
425	VFPEPTESGDEGEELGLPLLSTR2000000000000000	sp Q9BUR4 WAP53_HUMAN
426	TADSSSEDEEEYVVEK200022200000000000	sp P45973 CBX5_HUMAN
427	EELMSSDLEETAGSTSIPK2000000000000000	sp 000567 NOP56_HUMAN
428	GESAPTLSTSPSPSPSPTSPSPTLGR2000000	sp Q9H330 TM245_HUMAN
429	DLGHPVEEEDLESGDQEDEDDESEDPGK2000000000000000	sp 060763 USO1_HUMAN
430	SYSSPDITQAIQEEEK2000000000000000	sp P40818 UBP8_HUMAN
431	TEDGGWEWSDEFDEESEEGK2000000000000000	sp 095747 OXSR1_HUMAN
432	RQSDLVQCGVTSPSSAEATGK2E+19	sp Q9HC52 CBX8_HUMAN
433	NQGGYGGSSSSSYGSGR200000000	sp P09651 ROA1_HUMAN
434	LCDFGSASHVADNDITPYLVSR300000	sp Q13523 PRP4B_HUMAN
435	TPAAAAAMNLASPR10002000	sp Q9UQ35 SRRM2_HUMAN
436	SKGSDISDEEAAQSK200020020000000000	sp Q5T200 ZC3HD_HUMAN
437	KPISDNSFSSDEEQSTGPIK22000000000000	sp 060293 ZC3H1_HUMAN
438	STSATDTHHVMAR2000000000000000	sp 014874 BCKD_HUMAN
439	KETESAEADNLDDLEK2000000000000000	sp Q8IYB3 SRRM1_HUMAN
440	KPEDVLDDDDAGSAPLK200000	sp P35613 BASI_HUMAN
441	VDPSLMEDSDDGSLPTK20000000000	sp 015258 RER1_HUMAN
442	QVAGDAPVEQATAETASVHR200000	sp Q92766 RREB1_HUMAN
443	ERDSELSDTDSGCCLGQSESDK2000200000000000	sp Q96T88 UHRF1_HUMAN
444	GPPDFSSDEEREPTPVLGSGAAAAGR2.2E+21	sp P42166 LAP2A_HUMAN
445	SLYASSPGGVYATR2000000000	sp P42167 LAP2B_HUMAN
446	NSPGSQVASNPR200000000000	sp P08670 VIME_HUMAN
447	RPDYAPMESSDEEDEFQFIK1022000000000000	sp Q9H0D6 XRN2_HUMAN
448	SPVFSDESDLDFDISK200020000000000000	sp P55081 MFAP1_HUMAN
449	LLSSESEDEEELIPLAQR2020000000000000	sp 000566 MPP10_HUMAN
450	SEDEDSLEEAGSPAPGPCPR2E+20	sp Q96AY2 EME1_HUMAN
451	NQGGYGGSSSSSYGSGR2000	sp Q8TBB5 KLDC4_HUMAN
452	LMHNASDSEVDQDDVVEWK2020000000000000	sp P09651 ROA1_HUMAN
453	GGSYSQAASSDSAQGSVSLTACK2E+20	sp Q9H2P0 ADNP_HUMAN
454	SAQGTGFELGQLQSIR2000000000000000	sp Q09160 1A80_HUMAN
455	SVSSPTSNTPTPTK20000000000000	sp P05534 1A24_HUMAN
456	ALSSDSILSPAPDAR20000000000000	sp P30447 1A23_HUMAN
457	LTSIGSDEDEETETYQEK2002000000000000	sp P01891 1A68_HUMAN
458	GLSGEEEDDEPDCCNDER2000000000000000	sp P01892 1A02_HUMAN
459	SPLSPTETFSWPDVR2002000000000000	sp P10316 1A69_HUMAN
460	SPVFSDESDLDFDISK200000002000000000	sp Q9C0B5 ZDHC5_HUMAN
461	KGAGDGSDEEVDGK200000000	sp Q5M775 CYTSB_HUMAN
462	TPHDILEDINASPEMR200000	sp Q8IVT2 MISP_HUMAN
463	TSSFTEQLDEGTPNR2000000000000000	sp Q9UHW9 S12A6_HUMAN
464	SPSSDLDTAEGDDFELLDQSELSQLDPASSR2.2E+30	sp Q6P158 DHX57_HUMAN
465	TSSTLDSEGTFSYR2000000000000000	sp A1L390 PKHG3_HUMAN
466	KKPSTDEQTSSAEEDVPTCYLNVLSNSR2.2E+20	sp 000566 MPP10_HUMAN
467	SSGNSSSSGSGSGSTSAGSSSPGAR200000	sp P35579 MYH9_HUMAN
		sp Q9Y6Q9 NCOA3_HUMAN
		sp Q13439 GOGA4_HUMAN
		sp Q86VR2 F134C_HUMAN
		sp Q92609 TBCD5_HUMAN
		sp Q8N556 AFAP1_HUMAN
		sp Q12797 ASPH_HUMAN

468	GPPDFSSDEEREPTPVLGSGAAAAGR2. 2E+21	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
469	DDDDIDLFGSDDEEESEEAKR200000000000	sp P24534 EF1B_HUMAN
470	SVTSNQSDGTQESCESPDVLR2000000000000	sp Q9UGV2 NDRG3_HUMAN
471	AASSSSPGSPVASSPSR2000000000	sp Q92538 GBF1_HUMAN
472	VLQSDSETEDTNASPEK20200000000000	sp Q9HAW4 CLSPN_HUMAN
473	YAEISSDEDNDSDEAFESSR22000002000000000	sp Q6KC79 NIPBL_HUMAN
474	AKPVVSDDDSEEEQEEDR20002000000000	sp P51532 SMCA4_HUMAN
475	LPSSPVYEDAASF200000000000	sp Q14247 SRC8_HUMAN
476	TSAACAVTDLSDSDFDEK2002002000000	sp Q9NQZ2 SAS10_HUMAN
477	VLGSEGEDEALSPAK200000000020000	sp P18858 DNLI1_HUMAN
478	TASGSSVSLDGT2000000000000	sp Q92597 NDRG1_HUMAN
479	VEMYSGSDDDDFNKLK1020200000000000	sp 060841 IF2P_HUMAN
480	VEMYSGSDDDDFNKLK20200000000000	sp 060841 IF2P_HUMAN
481	QDDSPSGASYGQDYDLSPSR2000000000000000	sp Q9NYV4 CDK12_HUMAN
482	TLSTDAAPQPLATQAYSLVGPEVR2. 2E+23	sp Q9H6B4 CLMP_HUMAN
483	SEEDNEIESEEEVQPK20000000200000000	sp P52701 MSH6_HUMAN
484	VEESTGDPFGFSDDESPLVSSK200000000000	sp Q7Z5K2 WAPL_HUMAN
		sp Q6NXS1 IPP2M_HUMAN
485	IQQESSGEEDSDLSPEER2000000000000	sp P41236 IPP2_HUMAN
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486	QDDSPSGASYGQDYDLSPSR200000000000200	sp P17812 PYRG1_HUMAN
487	SGSSPDSEITELK220000000000	sp Q02880 TOP2B_HUMAN
488	SESDLTEPVVIPR2020000000000000	sp Q7Z460 CLAP1_HUMAN
489	QSSGSATNVASTPDNR2000000000000000	sp Q9NZ63 CI078_HUMAN
490	RGDSESEEDQDSEEV2020000000000000	sp 095251 KAT7_HUMAN
491	SSGSETEQVDFSDR202000000002000	sp Q86U70 LDB1_HUMAN
492	KMSGGSTMSSGGGNTNNSK2E+19	sp 060832 DKC1_HUMAN
493	AGLESGAEPGDGSDTTK20000000200000	sp Q6ZRS2 SRCAP_HUMAN
494	DEEQSEADAGSPPTPGPTTLGPK2. 0000020002E+20	sp P16989 YBOX3_HUMAN
495	NYAGEEEEEEGSGSSEGFDPATDR202200000000000	sp Q9H8G2 CAAP1_HUMAN
496	STDSSVSGSLQQETK2000000000000000	sp Q9ULX3 NOB1_HUMAN
497	KDSDDDGGGWITPSNIK2000000000000000	sp Q9NZI8 IF2B1_HUMAN
498	QGSPVAAGAPAK200000000000	sp Q9P1Y6 PHRF1_HUMAN
499	SGSESEDDGETLLEVAGTQK2. 02002E+22	sp Q8NE71 ABCF1_HUMAN
500	KLSVPTSDEEDEVPAKPR200220000000000000	sp Q9UFC0 LRWD1_HUMAN
501	ACASPSAQVEGSPVAGSDGSQPAVK2E+22	sp Q96AY2 EME1_HUMAN
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503	GSAPHSEDLPEQEEIILGSDDEQEDPNDYCK20000000000000	sp Q2KHR3 QSER1_HUMAN
504	GQDTVAIEGFTDEEDTESGEGQYR2. 0000000000002E+22	sp Q9NYF8 BCLF1_HUMAN
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506	SPVGSQAPQAAAPAPAAHVAGNPGGDAAPAATGTAASLATAAGSEDAEK2E+47	sp P16989 YBOX3_HUMAN
507	EEGPPPPSPDGASSDAEPEPPSGR200000200000000000	sp Q9NTJ3 SMC4_HUMAN
508	SASDTEELNSQDSPPK20000000000000	sp 014745 NHRF1_HUMAN
509	HCDSINDFGSESGGCGDSSPGPSASQGPR200000000000	sp Q8TD19 NEK9_HUMAN
510	THSTSSSLGSGESPF2000000000000000	sp Q9UGV2 NDRG3_HUMAN
511	DSPSKSSAEQTPEDTPNK20000000000000	sp 043493 TGON2_HUMAN
512	AEGEWEDQEALDYFSDKESGK20000000	sp Q9NYF8 BCLF1_HUMAN
513	DLFDLNSSEEDTEGFSER22000000000000	sp Q9Y3T9 NOC2L_HUMAN
514	GGGGQDNGLEGLGNSR200	sp P08621 RU17_HUMAN
515	VPVRPPQYSDDEDDYEDDEDDVQNTNSALR3000000000000000	sp Q92560 BAP1_HUMAN
516	LQELESCSLGSDTDVR20002000000000	sp Q14C86 GAPD1_HUMAN
517	IHDLEDDLEMSDASDASGEEGGR1220020020000000	sp P35269 T2FA_HUMAN
518	GLNTSQESDDDILDESSPEGTQK2. 002E+20	sp Q5VV41 ARHGG_HUMAN

519	GVPAGNSDTEGGQPGR20000000000	sp Q9UKV3 ACINU_HUMAN
520	EAEQAASEAAGDTPGSSPSSLYEEPLGQPPR2. 2E+21	sp Q86TC9 MYPN_HUMAN
521	IACDEEFSDEDEGEGGR202000000000	sp Q92769 HDAC2_HUMAN
522	YQPLASTASDNDFVTPEPR200200000000000	sp Q15021 CND1_HUMAN
523	REEGPPPPSPDGASSDAEPEPPSGR200000200000000000	sp Q9NTJ3 SMC4_HUMAN
524	GFSDSGGGPPAK200000000000	sp Q9H307 PININ_HUMAN
525	VEEDLKADEPSSEESDLEIDK200200000000	sp P50502 F10A1_HUMAN
526	KVELSESEEDKGGK202000000000	sp Q8IZP2 ST134_HUMAN
527	RPDHSGGSPSPTSEPAR200000000000	sp Q8IYB3 SRRM1_HUMAN
528	REEGPPPPSPDGASSDAEPEPPSGR200002200000000000	sp Q8TAD8 SNIP1_HUMAN
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		sp P10316 1A69_HUMAN
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		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
		sp P01892 1A02_HUMAN
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533	SSSSPELQTLQDILGDPGDK2000000000000000000	sp P49815 TSC2_HUMAN
534	YVDEENSdGETSNHR2000000000	sp Q8N9T8 KRI1_HUMAN
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536	AETPTESVSEPEVATK2000002000000000	sp P18858 DNLI1_HUMAN
537	TSSTLDSEGTFNSYR220000000000000	sp Q92609 TBCD5_HUMAN
538	SPSEESAPTTSPESVSGVPSSGSSGR200000000000000000	sp P54725 RD23A_HUMAN
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542	SEDESETEDEEEKSQEDQEQK2020000000000000000	sp P25205 MCM3_HUMAN
543	REESPMdVDQPSPSAQDTQSIASDGTpQGEK2. 00000002E+28	sp Q7Z6Z7 HUWE1_HUMAN
544	ENSPSSQSAGLSSINK2000000000000000	sp Q9H2Y7 ZN106_HUMAN
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546	GESENAGTNQETR2000000000000	sp Q08170 SRSF4_HUMAN
547	SPDLGEYDPLTQADSDEEDDLVLNLQK2002000000000000	sp A2RU67 K1467_HUMAN
548	SQSLPNSLDYtQTSdPGR200000000000000000	sp Q96TC7 RMD3_HUMAN
549	WAHDKfSGEEGEIEDDESgTENR200000000000000000	sp Q9Y2W1 TR150_HUMAN
550	NLGSINTELQDVQR2000000000000	sp 075396 SC22B_HUMAN
551	KGSGSEQEGEDEEGGER202000000000000000	sp Q6PD62 CTR9_HUMAN
552	SGSSSPDSEITELK2022200000000000	sp P17812 PYRG1_HUMAN
553	TEESPASDEAGEK200200000000	sp P05114 HMGn1_HUMAN
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555	FSGEEGEIEDDESgTENR2000000	sp Q9Y2W1 TR150_HUMAN
556	VLSDSEDEEKDADVPGTSTR20200000000000000000	sp Q8WWQ0 PHIP_HUMAN
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		sp Q58FF8 H90B2_HUMAN
558	NQGGYGGSSSSSYGSGR2000000	sp P09651 ROA1_HUMAN
559	VQMTSPSSTGSPMFK2000000000000	sp P49790 NU153_HUMAN
560	DHSPTPSVFNSDEER2000000000000000	sp Q6UN15 FIP1_HUMAN
561	VTTEIQLPSQSPVEEQSPASLSSLR200000000000000000	sp Q8WWI1 LMO7_HUMAN
562	VTNDISPESSPGVGR20020000000	sp Q15154 PCM1_HUMAN
563	TDVSNFDEEFTGEAPTLSPPR20000	sp Q16512 PKN1_HUMAN
564	GLRDSHSSEEDeASSQTdLSQTISK2. 022E+21	sp Q5JTV8 TOIP1_HUMAN
565	GSAPHSEDLPEQEEIILGSDDDEQEDPNdYCK2E+32	sp Q96SB4 SRPK1_HUMAN

566	QVAEQGGDLSPAANR2000000	sp 094804 STK10_HUMAN
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570	AAVGQESPGGLEAGNAK20000000000	sp Q29RF7 PDS5A_HUMAN
571	AQGSDEDKLQNSDDDEK200000002000000	sp Q8WVC0 LEO1_HUMAN
572	AGMSSNQSISSPVLDVPR22000000000	sp Q9UQ35 SRRM2_HUMAN
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575	WAHDKFSGEEGEIEDDESGTENR200000000002000000	sp Q9Y2W1 TR150_HUMAN
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583	AGEPNSPDAEEANSPDVTAGCDPAGVHPPR200000000000000000	sp Q08J23 NSUN2_HUMAN
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588	ESEDKPEIEDVGSDEEEEEK20000000	sp P07900 HS90A_HUMAN
589	GAVENEEDLPELSDSGDEAAWEDEDDADLPHGK2. 02E+21	sp 060678 ANM3_HUMAN
590	LENEGSEDEDIETDVLVSPQALK20000000000000000000	sp 060502 OGA_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
591	GGSYSQAASSDSAQGS DVSLTACK2E+20	sp P05534 1A24_HUMAN
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		sp P01892 1A02_HUMAN
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593	LQEEGGGSDEEETGSPSEDGMQSAR20000000000000000000	sp Q13610 PWP1_HUMAN
594	VPSSDEEVVEEPQSR22000000000000	sp Q9C0C2 TB182_HUMAN
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596	NEEDSEGSSDEDEDEDGVSAAATFLKK2. 0022E+22	sp Q99613 EIF3C_HUMAN
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599	SFEVEEVETPNSTPPR2000000000000000	sp Q9Y2W1 TR150_HUMAN
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		sp P01892 1A02_HUMAN
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605	SPPGAGGSPPGA AVAPSALQLSPAAAAAAHR2. 0000002E+32	sp P07910 HNRPC_HUMAN
606	MESEGGADDSAEGLDLLDDDDNEDRGDDQLELIK2E+25	sp Q92522 H1X_HUMAN
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648	SPSDSSTASTPVAEQIER2020000000000000000	sp Q16643 DREB_HUMAN
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653	SRSPLDKDTYPPSASVVGASVGGHR2.0000003E+23	sp Q96T37 RBM15_HUMAN
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681	NQKPSQVNGAPGSPTEPAGQK2000000000	sp Q9H4L4 SEN3_HUMAN
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685	SSPGQTPEEGAQALAEFAALHGPALR2E+25	sp Q15459 SF3A1_HUMAN
686	KKASSDSEDSSEEEEEVQGPPAK2.22020022E+21	sp Q14166 TTL12_HUMAN
687	AGMSSNQSISSPVLDVPR2002000000000	sp Q14978 NOLC1_HUMAN
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		sp Q29960 1C16_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
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		sp P30510 1C14_HUMAN
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		sp Q29865 1C18_HUMAN
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708	DLGSTEDGDGTDDFLTDKEDEK2000020000000	sp Q15527 SURF2_HUMAN
709	SYSAGNASQLEQLSR20000000000000	sp Q14678 KANK1_HUMAN
710	DSHSSEDEASSQTDLSQTISK2.02E+21	sp Q5JTV8 TOIP1_HUMAN
711	GSGTASDDEFENLR2000000000	sp Q7Z6Z7 HUWE1_HUMAN
712	EKEPSYPMPVQETQAPESPGENSEQALQTLSPR20000000000000000	sp Q7Z434 MAVS_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
713	KGGSYSQAASSDSAQGSVDVSLTACK20002002000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
714	SNSAPLIHGLSDTSPVFQAEAPSAR2E+23	sp Q96E09 F122A_HUMAN
715	FYDLSDSNSLGEDSK20202002000000	sp Q9H501 ESF1_HUMAN
716	LDSPPSPITEASEAAEAAEAGNLAVSSR2.0000002E+27	sp Q9C0C2 TB182_HUMAN
717	TPDLSNQSSDQANEWETASESSDFNER202020000000	sp Q9Y520 PRC2C_HUMAN
718	GESAPTLSTSPSPSSPSPTSPSPTLGR200000000000	sp Q9H330 TM245_HUMAN
719	TIQNSSVSPTSSSSSSSSTGETQTQSSSR2000000000000000	sp Q7Z417 NUFP2_HUMAN
720	EELMSSDLEETAGSTSIPK12000000000000000	sp O00567 NOP56_HUMAN
721	STDSPGADAELPESAR20200000000000000	sp Q9Y4K1 AIM1_HUMAN
722	LPQDHADSCVVSDDDEELSR2200000000	sp P78317 RNF4_HUMAN
723	QLHLEGASLELSDDDTESK2000000000	sp P35580 MYH10_HUMAN
724	STSATDTHHVMAR20000000000000	sp O14874 BCKD_HUMAN
725	AGMSSNQSISSPVLDVPR2000000000	sp Q9UQ35 SRRM2_HUMAN
726	EAEDSDSDNIKR2020000000	sp Q96ST2 IWS1_HUMAN
727	ISSTPTTETIR2000000000	sp Q96QE3 ATAD5_HUMAN
728	VGDGDLAEEIPENEVSLR200000000000000	sp Q14684 RRP1B_HUMAN



729	AASAAAAASAAAASASGSPGPEGGSAGGEK2. 000000002E+23	sp Q13263 TIF1B_HUMAN
730	NSSEASSGDFLDLK200000000000	sp Q9UK76 HN1_HUMAN
731	HQEVQDQDPVFGSDSSGYQSDHKK2022000200000	sp Q3B726 RPA43_HUMAN
732	SGGSGHAVAEPASPEQELDQNK20000000000	sp P46013 KI67_HUMAN
733	SLSSPTVTLAPLEGAK2200000000000000	sp Q96PU5 NED4L_HUMAN
734	GGVTGSPEASISGSK200000000000	sp Q09666 AHNK_HUMAN
735	MSPMGTASGSNSPTSDSASVQR2E+21	sp 076080 ZFAN5_HUMAN
736	SGSSQELDVKPSASPQER2000000000000000	sp Q9UQ35 SRRM2_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
737	KGGSYSQAASSDSAQGSVSLTACK2E+22	sp P01892 1A02_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
738	LSSGFDDIDLPSAVK2200000000000000	sp Q9Y446 PKP3_HUMAN
739	LPSGGAASPTGSAVDIR2000000000000000	sp Q09666 AHNK_HUMAN
740	SRSEDLSPESDEEGYALSGR2. 000000002E+20	sp Q15751 HERC1_HUMAN
741	VDSTTCLFPVEEK200000000000	sp Q06210 GFPT1_HUMAN
742	LLLAEDSEEEVDFLSER200000000000	sp Q14676 MDC1_HUMAN
743	SSSLGSTPHEELER22002000000000	sp Q9NRA8 4ET_HUMAN
744	HLDGEEDGSSDQSQASGTTGGR20020020000000	sp Q9UNF1 MAGD2_HUMAN
745	DVDVSEDSPPPLPER2000000000	sp Q05209 PTN12_HUMAN
746	NDSLSSLDFFFFDLSR202200000000000000	sp P25054 APC_HUMAN
747	ALDISLSSGEEDEGDEEDSTAGTTK2. 02E+21	sp Q8IY81 SPB1_HUMAN
748	SLSSPTDNLELSLR200000000000	sp Q9Y2X7 GIT1_HUMAN
749	VSSPLPSAMTDAANSQAAAK2. 00002E+21	sp Q8WXI9 P66B_HUMAN
750	ASTPDWVSEGPQPGLR2000000000000000	sp Q8IVT2 MISP_HUMAN
751	VVEAVNSDSDSEFGIPK202000000000	sp Q02880 TOP2B_HUMAN
752	DQPDGSSLSPAQSPSQSPPAASSLR200020000000000000	sp Q6WCQ1 MPRIP_HUMAN
753	GSPSSPTPPAAAAPAEQAPR202000000000000000	sp Q86VQ1 GLC11_HUMAN
754	TISAQDTLAYATALLNEK200000000000000000	sp Q9UPU5 UBP24_HUMAN
755	TPLSSLSATSNNSTVACIDR200200000000	sp P00533 EGFR_HUMAN
756	GPGQVPTATSALSLELQVEVEPLGLPQASPSR20000	sp Q6P2E9 EDC4_HUMAN
757	GGEFDEFVNDTDDDLPI SK2000000000	sp Q6PD62 CTR9_HUMAN
758	EVEDKESEGEEDDEDL SK2000000000000000	sp 095218 ZRAB2_HUMAN
759	GFGDGYNGYGGGPGGNFGGSPGYGGGR200000000	sp P22626 ROA2_HUMAN
760	SQSDLDDQHDYDSVASDEDTDQEPLR2002000000000000	sp Q9Y2X7 GIT1_HUMAN
761	KYSEVDDSLPSGGEKPSK200000000000000000	sp Q05D32 CTSL2_HUMAN
762	DDAESSSEPEEENVGNDTK2200000000000000	sp Q13823 NOG2_HUMAN
763	MESEGGADDSAEGLDLDLDDDDNEDR1. 000000002E+25	sp P07910 HNRPC_HUMAN
764	SPGEYINIDFGEPGAR200000000000000000	sp Q9Y4H2 IRS2_HUMAN
765	QSFDDNDSELEDK20000000	sp 060841 IF2P_HUMAN
766	NALFPEVFSPTPDENSQNSR2000000000000000	sp Q8IVT2 MISP_HUMAN
767	SSSISEEKGSDDEKPR20000000	sp Q9UKV3 ACINU_HUMAN
768	MEVGPFSTGQESPTAENAR200000000	sp Q9Y4W2 LAS1L_HUMAN
769	SKFDSDEEEDTENVEAASSGK2000000000000020000	sp Q8TF01 PNISR_HUMAN
770	SGSSSPDSEITELK20200000000000	sp P17812 PYRG1_HUMAN
		sp Q9BQE3 TBA1C_HUMAN
		sp Q6PEY2 TBA3E_HUMAN
771	TIGGGDDSFNTFFSETGAGK20000000000000	sp Q71U36 TBA1A_HUMAN
		sp Q13748 TBA3C_HUMAN
		sp P68363 TBA1B_HUMAN
772	EYVSNDAAQSDDEEK2000000	sp Q5JSH3 WDR44_HUMAN
773	ASHASSDWTTPRPR20022000000000	sp Q96EB6 SIR1_HUMAN

774	NTSSDNSDVEVMPAQSPR220020000000000000	sp Q9HCK8 CHD8_HUMAN
775	ENPPVEDSSDEDDKR220000000	sp Q8TCJ2 STT3B_HUMAN
776	DELAELSEAESGDEKPK20002000000000	sp Q3L8U1 CHD9_HUMAN
777	YESQLSTNEEKVDTDDR20000000000000	sp Q13433 S39A6_HUMAN
778	LASDSAESDSR20200000000	sp Q8WXG6 MADD_HUMAN
779	TAVQYIESSDSEEIETSELPQK2202000000000000	sp Q14527 HLTF_HUMAN
780	SSSLIQLTSQNSSPNQR200000000000000000	sp O95639 CPSF4_HUMAN
781	KTSSVSSISQVSPER2000000000000000	sp P42356 PI4KA_HUMAN
782	LFDEEEDSSEKLFDDSDER2000000020000	sp O43719 HTSF1_HUMAN
783	ATGDGSSPELPSLER22000000000	sp Q9H2Y7 ZN106_HUMAN
784	FNSESESGSEASSPDYFGPPAK202000000000000000	sp Q9BW71 HIRP3_HUMAN
785	LGASNSPGQPNSVK2000000000	sp P49756 RBM25_HUMAN
786	AGEPNSPDAEEANSPDVTAGCDPAGVHPPR2.00000002E+25	sp Q08J23 NSUN2_HUMAN
787	RPTETNPVTSNSDEECNETVK2202000000000000	sp P46100 ATRX_HUMAN
788	AYACHAEHENDSDDDDEAEDDDETEELGSDEDDIDEDGQEYLEILAK2E+36	sp O95373 IPO7_HUMAN
789	EGSEFSFSDGEVAEK200202000000000	sp Q9BVS4 RIOK2_HUMAN
790	DAHVDSPTSTDTEAQLTVER200000000000000000	sp Q8IVF2 AHNK2_HUMAN
791	QASTDAGTAGALTPQHVR200000000000000000	sp P46937 YAP1_HUMAN
792	DGGSGNSTIIVSR20002000000	sp Q7Z6Z7 HUWE1_HUMAN
793	SRPTSEGSDIESTEPQK20000000000	sp Q4G0J3 LARP7_HUMAN
794	NSSEASSGDFLDLK2000000000000000	sp Q9UK76 HN1_HUMAN
795	ESYTEEIVSEAESHVSGISR2000000200000	sp Q5QJE6 TDIF2_HUMAN
796	EAEQAASEAAGDTPGSSPSSLYEEPLGQPPR22000000000000000000	sp Q86TC9 MYPN_HUMAN
797	VQEKPDSPGGSTQIQR20000000000	sp Q13459 MYO9B_HUMAN
798	VAAAAGSGSPPGSPGHDR20000002000000	sp Q9BTU6 P4K2A_HUMAN
799	GPSPSPTPPAAAAPAEQAPR2.002E+19	sp Q86VQ1 GLC11_HUMAN
800	VIKDEALSDGDDL20000000	sp Q01831 XPC_HUMAN
801	GTGGGQLQDLDCSSSDDEGAAQNSTKPSATK2.22E+19	sp P08240 SRPR_HUMAN
802	EADDDEEVDDNIPEMSPK2000	sp P26358 DNMT1_HUMAN
803	SGSPSDNSGAEEMEVSLAKPK2.0000200001E+19	sp P31749 AKT1_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
804	KGGSYSQAASSDSAQGSVSLTACKV200020020000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
805	AGDRNSEDGVMTFSSVK2000000000000000	sp Q9NY61 AATF_HUMAN
806	DNQESSDAELSSSEYIK2000002000000	sp Q96JM3 CHAP1_HUMAN
807	YGLQDSDEEEEEHPSK200000000000	sp P52948 NUP98_HUMAN
808	SPSFGEDYGPSR2020000000000000	sp P49761 CLK3_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
809	LNQSGTSVGTDEESDVTQEEER2000000000000000	sp A6NKT7 RGPD3_HUMAN
		sp P0DJ0 RGPD1_HUMAN
		sp P0JD1 RGPD2_HUMAN
810	QQASASTDGVDKLSGK2000	sp P46100 ATRX_HUMAN
811	DPTSLLGVLQAEADSTSEGLEDAVHSR20000000000000	sp Q27J81 INF2_HUMAN
812	SYLEGSSDNQLKDESTPVDDR2.000002E+22	sp Q9UHI6 DDX20_HUMAN
813	KDSLDPADLASCGLSQSSASFFTPR2E+24	sp Q86SQ0 PHLB2_HUMAN
814	GPSLDIDTPDVNIEGPEGK20000000000000	sp Q09666 AHNK_HUMAN
815	GSSDGRGSDSEDLPHR200002000000000000	sp Q9UPQ0 LIMC1_HUMAN
816	SKSDGEAKPEPSPSPR202000000000000000	sp P26358 DNMT1_HUMAN

817	DIEVLSEDTDYEEDEVTKK2002000000000000	sp Q4LE39 ARI4B_HUMAN
818	AASPPASASDLIEQQQK2000000000000000	sp Q5VSL9 STRP1_HUMAN
819	SQSESSDEVTELDLSHGK2022000000000000	sp Q86YS7 C2CD5_HUMAN
820	NESEDNKFSDSDDDFVQPR200000200200000000	sp O15164 TIF1A_HUMAN
821	MESEGGADDSAEGLDLDLDDDDNEDR2.0000002E+23	sp P07910 HNRPC_HUMAN
822	NPWSDDESSESLEETEPVVIPR2020200000000000	sp Q02880 TOP2B_HUMAN
823	SPTMEQAVQTASAHLPAPAAVGR2E+23	sp Q8ND56 LS14A_HUMAN
824	HDYDDSSSEEQAEDSYEASPGSETQR2.2E+21	sp P98175 RBM10_HUMAN
825	FAQSAPGSPLSSQPVLITVQR2000000000000000	sp Q01167 FOXK2_HUMAN
826	DDDLVEFSDLESEDDERPR20002000000000	sp Q9HCK8 CHD8_HUMAN
827	FDSNEEDSASVFSFGLK200000000000000000	sp Q02880 TOP2B_HUMAN
828	KPVLPALTINPTIAEGPSPTSEGASEANLVDLQK200000000000000000	sp P36507 MP2K2_HUMAN
829	EVDSSDDTTEEGPENSQGAQELSLAPHPEVSQSVAPK2.2E+34	sp Q6P0Q8 MAST2_HUMAN
830	NSISDDDTSEDEL20002020000000	sp Q76FK4 NOL8_HUMAN
831	DLLSDLQDISDSER200000200000	sp P21127 CD11B_HUMAN
		sp Q9UQ88 CD11A_HUMAN
832	EVSDSEAGGGPQGER2020000000000000	sp Q9BW71 HIRP3_HUMAN
833	KGSDDDGGDSPVQDIDTPEVDLYQLQVNTLR2.0000002E+29	sp O75446 SAP30_HUMAN
834	KSPVGKSPSTGSTYGSSQK2.00002E+19	sp Q9Y2W1 TR150_HUMAN
835	EEPLSEEPCTSTAIASPEK200000000000000000	sp Q9Y2X3 NOP58_HUMAN
836	VSPAAAQLADTLFSK2000000000000000	sp O14686 KMT2D_HUMAN
837	GKEELAEAEIKDSPDSPEPPNK20000000	sp Q9NVM9 ASUN_HUMAN
838	SPAGPAATPAQAQAASPR2E+19	sp Q13428 TCOF_HUMAN
839	ILSQSTDSLNR2000000000	sp Q92974 ARHG2_HUMAN
840	SPDEVTLTSIVPTR2000000000000000	sp Q9UKS6 PACN3_HUMAN
841	KQSFDDNDSELEDK20000020000000	sp O60841 IF2P_HUMAN
842	SPVGS GAPQAAAPAPAAHVAGNPGGDAAPAATGTAASLATAAGSEDAEK2E+51	sp P16989 YBOX3_HUMAN
843	GAGDGSDEEVDGK2000000000	sp P35579 MYH9_HUMAN
844	SDVQEESEGSDDDNKDSAAFEDNEVQDEFLEK2.0000000002E+27	sp Q9UIG0 BAZ1B_HUMAN
845	LPSGSGAASPTGSAVDIR2000000000	sp Q09666 AHNK_HUMAN
846	KEDTAFSDWSEDEVPDR200200000000	sp Q5T200 ZC3HD_HUMAN
847	GISSSNEGVEEPSK20000000000000	sp Q9NXV6 CARF_HUMAN
848	KPPAPSPVQSQSPSTNWSPAVPVK2.000200000002E+19	sp Q8IYB3 SRRM1_HUMAN
849	EDQEAELLSEPEEESER2000000000	sp Q63HN8 RN213_HUMAN
850	ELVGDGTSGEQGDHEPSGSETEEDTSSSPHR2.0000000202E+23	sp PODJ93 SIM13_HUMAN
851	NSPSNLSSSDTGSVGGTYR200000000000000000	sp O43166 SIL1_HUMAN
852	TPSYSISSTLNPAPEFILGCTASK2E+23	sp Q14694 UBP10_HUMAN
853	SASPDLDLSSNWEADLGNEER2.00000002E+21	sp O00193 SMAP_HUMAN
854	RSSTVAPAQPDGAESEWTDVETR2E+22	sp Q02241 KIF23_HUMAN
855	GNSRPGTPSAEGGSTSSTLR200020000000000000	sp P35269 T2FA_HUMAN
856	ANSGVDLDSSGEFASIEK200000002000000000	sp Q92766 RREB1_HUMAN
857	SSVAAPEKSSNSVDEETAESDTSPVLEK2.22002E+22	sp Q99590 SCAFB_HUMAN
858	SPSAPAATSSASSSTSSYK2E+19	sp Q15345 LRC41_HUMAN
859	EEGPPPPSPDGASSDAEPEPPSGR200002000000000000	sp Q9NTJ3 SMC4_HUMAN
860	KQSLGELIGTLNAAK2000000000000000	sp P60174 TPIS_HUMAN
861	ELKSDASSDQGPASPEK2002200000000000	sp Q6BDS2 URFB1_HUMAN
862	AKPAMPQDSVSPR2000	sp P53396 ACLY_HUMAN
		sp P08238 HS90B_HUMAN
863	IEDVGSDEEDDSGKDK200000200000	sp Q58FF8 H90B2_HUMAN
864	SGSSSPDSEITELK200000000000	sp P17812 PYRG1_HUMAN
865	MEQQYESSSDGTEKLPER200000000000	sp P46100 ATRX_HUMAN
866	NGFPHPEDCNPSEAASEESNSEIEQEIPEVQK2000000000000000	sp Q9NR30 DDX21_HUMAN

867	SSLSGDEEDELFK202000000000	sp Q5T1M5 FKB15_HUMAN
868	LSEDEEEDEAEDDQSEASGKK2. 2E+21	sp Q8NEJ9 NGDN_HUMAN
869	EVADIFNAPSDDEEFVGF20000000000	sp Q96GN5 CDA7L_HUMAN
870	SGSMDPSGAHPSVR200000000000	sp Q07666 KHDR1_HUMAN
871	GLQLTPGIGGMQHHFFDDED2000000000000000	sp P12270 TPR_HUMAN
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873	KLSVPTSDEEDEVPAKPR2200000000000000	sp Q8NE71 ABCF1_HUMAN
874	SSSEDAESLAPR200000000000	sp Q4G0J3 LARP7_HUMAN
875	YMAENPTAGVVQEEEDNLEYSDGNPIAPTK20000000000	sp Q86XP3 DDX42_HUMAN
876	HSSTGDSADAGPPAAGSAR200000000000000000	sp Q6ZU35 K1211_HUMAN
877	ADDFPVRDDPSDVTDEDEGPAEPPPPK200200000000000000	sp Q3YEC7 RABL6_HUMAN
878	EKEISDDEAEEK2000000000	sp P08238 HS90B_HUMAN
879	ASGQAFELILSPR2000	sp P16949 STMN1_HUMAN
880	KTSDANETEDHLESLICK200000000000000000	sp Q09161 NCBP1_HUMAN
881	SYSSPDITQAIQEEK20000000000000	sp P40818 UBP8_HUMAN
882	VQGEAVSNIQENTQTPTVQEESEEEVDETGVK2E+21	sp E9PAV3 NACAM_HUMAN
883	EYVSNDAAQSDDEEKLQSQPTD200000000000000000	sp Q13765 NACA_HUMAN
884	SEDESETEDEEEK2000000000	sp Q5JSH3 WDR44_HUMAN
885	EEGSSDEISSGVGSESEGLNSPVK2. 2000200002E+22	sp P25205 MCM3_HUMAN
		sp P52701 MSH6_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
886	GGSYSQAASSDSAQGSVSLTACK2E+22	sp P05534 1A24_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
		sp P30447 1A23_HUMAN
887	TSSFAEPGGGGGGGGGGPGGSASGPGGTGGGK2E+30	sp P49840 GSK3A_HUMAN
888	TESPATAAETASEELDNR20000000000000	sp Q9NTJ3 SMC4_HUMAN
889	SYDNLTTACDNTVPLASR200000000000000000	sp Q13615 MTMR3_HUMAN
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891	EMEHNTVCAAGTSPVGEIGEEK200000000000	sp P18583 SON_HUMAN
892	TAGSSVTSLDGTR2000000	sp Q92597 NDRG1_HUMAN
893	HIISATSLSTSPTELGSR200000000000	sp Q5JSZ5 PRC2B_HUMAN
894	SPGIISQASAPR20000000000000	sp Q08495 DEMA_HUMAN
895	TSPADHGGVSGSESGSAVDSVAGEHSVSGR200200000000000000	sp Q5T4S7 UBR4_HUMAN
896	DQTASAPATPLVNK20000000000	sp Q53SF7 COBL1_HUMAN
897	NHSDSSTSESEVSSVPLK20000	sp Q9NY27 PP4R2_HUMAN
898	SRSPLDKDYPPSASVVGASVGGHR2. 000002E+23	sp Q96T37 RBM15_HUMAN
899	RPSQEQSASASSGQPQAPLNR2E+19	sp Q5T1M5 FKB15_HUMAN
900	KVEEQEADEEDVSEEAESK200000000	sp Q9H3N1 TMX1_HUMAN
901	SSSVGSSSSYPISPAISR200000000000000000	sp Q15149 PLEC_HUMAN
902	GSSGVDETFLFALPAASEPVIR2. 02E+20	sp Q9NQC3 RTN4_HUMAN
903	AGLESGAEPGDGSDTTK200000	sp Q060832 DKC1_HUMAN
904	KPPAPSPVQSQSPSTNWSPAVPK2. 0002E+19	sp Q8IYB3 SRRM1_HUMAN
905	DPSVEVTQDLIDESEER200000	sp Q5MIZ7 P4R3B_HUMAN
906	SESELIDELSEDFDR2000000000000000	sp P20810 ICAL_HUMAN
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908	KEESEESDDMGFLFD2000000000000000	sp P05387 RLA2_HUMAN
		sp P05386 RLA1_HUMAN
909	SNSLPHSAVSNAGSK20000000000000	sp Q8TBZ3 WDR20_HUMAN
910	SSTVTEAPIAVVTSR20000000000000	sp Q8TD19 NEK9_HUMAN
911	SVGGSGGGSFGDNLVTR2000000000	sp P02545 LMNA_HUMAN
912	TKDQPDGSSLSPAQSPSQSPPAASSLR200020000000000000	sp Q6WCQ1 MPRIP_HUMAN
913	KASGSENEGDPGR20000000000000	sp Q02880 TOP2B_HUMAN

914	VSDPISTSESSEEEEEAEATAK222022000000000000	sp Q13428 TCOF_HUMAN
915	INSSGESGDESDEFLQSR2002000200000000	sp O60841 IF2P_HUMAN
916	SRSPTPPSSAGLGSNSAPPIDSR2. 02002E+22	sp Q8IWX8 CHERP_HUMAN
917	SVSSPTSSNTPTPTK22000000000000	sp Q5M775 CYTSB_HUMAN
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		sp P10316 1A69_HUMAN
918	GGSYSQAASSDSAQGSVDVSLTACK20000000000000	sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
		sp P01892 1A02_HUMAN
919	YTPTSPSYSPSSPEYTPSPK3. 00000002E+21	sp P24928 RPB1_HUMAN
920	EMDESLANLSEDEYYSEEER1. 00000002000002E+19	sp O60341 KDM1A_HUMAN
921	SLSTSGESLYHVLGLDK2000000000000000	sp Q9H3Z4 DNJC5_HUMAN
922	DAPTSPASVASSSSTPSSK2000000000000000	sp Q04726 TLE3_HUMAN
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924	IACKSPPPEVDTPSTK2000000000000000	sp P46013 KI67_HUMAN
925	APVSSSTSVIQSNTPTPPSQPLNETAEESR2. 02E+19	sp Q9HC35 EMAL4_HUMAN
926	TCEERPAEDGSDEEDPDSMEAPTR20000000000000	sp Q9H6Y2 WDR55_HUMAN
927	EFDEDSDEKEEEEDTYEK20000000000000	sp O43719 HTSF1_HUMAN
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929	KDSLTLQAQEQGNLLN20000000000000	sp Q5JTD0 TJAP1_HUMAN
930	EGMNPSYDEYADSDQHDAYLER1. 000000002E+22	sp Q08945 SSRP1_HUMAN
931	STGSATSLASQGER200200000000	sp Q5SW79 CE170_HUMAN
932	TRSPDVISSASTALSQDIPEIASEALSR2E+26	sp Q6P2E9 EDC4_HUMAN
		sp Q09160 1A80_HUMAN
		sp P05534 1A24_HUMAN
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		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
934	ENSPAVSPTTNSTAPFGLKPR2. 0002E+19	sp Q9Y2L9 LRCH1_HUMAN
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936	MLAESDESGDEESVSQTDKTELQNTLR2. 002E+23	sp P22059 OSBP1_HUMAN
937	APSSSSLNSATSSQSK2000000000000000	sp Q5F1R6 DJC21_HUMAN
938	METVSNASSSSNPSSPGR20000	sp Q9NTI5 PDS5B_HUMAN
939	SQDVAVSPQQQCSK2000000000	sp Q96F86 EDC3_HUMAN
940	EENPESDGEPVVEDGTSVK20000000000000	sp Q13439 GOGA4_HUMAN
941	SAYQEYDSDSDVPEELKR200000000000	sp Q96JG6 CC132_HUMAN
942	ESESEAEAGAAELR2000000000000000	sp Q8NAF0 ZN579_HUMAN
943	SGSSQELDVKPSASPQER2200000000000000	sp Q9UQ35 SRRM2_HUMAN
944	NKSNEQSMGNWQIK20000000000000	sp P02545 LMNA_HUMAN
945	SGSIVELIAGGGSSCSPVLSR2. 000000000002E+19	sp P42858 HD_HUMAN
946	SSSLGSTEPHEELER2200020000000000	sp Q9NRA8 4ET_HUMAN
947	AEGEWEDQEALDYFSDKESGK20002000	sp Q9NYF8 BCLF1_HUMAN
948	KPGDASSLPDAGLSPGSQVDSK2000000000	sp Q13459 MYO9B_HUMAN
949	SPPEGDTTLFLSR2000000000000000	sp Q9BTC0 DIDO1_HUMAN
950	DSHSSEDEASSQTDLSQTISKK2. 2E+20	sp Q5JTV8 TOIP1_HUMAN
951	SPTMEQAVQTASAHLPAVAVGR2. 001E+23	sp Q8ND56 LS14A_HUMAN
952	TASESISNLSEAGSIK2000	sp P30622 CLIP1_HUMAN
953	HQASDSENEELPKPR20200000000000	sp Q96ST2 IWS1_HUMAN
954	LPSDITDSDASEDPGGAER202002000000000000	sp Q9UJK0 TSR3_HUMAN
955	SGVSSDNEDDDDEEDGNYLHPSLFASK2E+23	sp Q12872 SFSWA_HUMAN
956	DQQPSGSEGEDDDAEALK2020000000000000	sp Q9NXG2 THUM1_HUMAN
957	EPTPSIASDISLPIATQELR20020000000000	sp A0FGR8 ESYT2_HUMAN

958	AASPESASSTPESLQAR2022000000000	sp Q96BT3 CENPT_HUMAN
959	ASGEMASAQYITAALR2000000000000000	sp P54578 UBP14_HUMAN
960	DGESDGFADDEESGEEGEDQEEGREPGAGR2.000000002E+28	sp P48681 NEST_HUMAN
961	DWQDDQSDNQSDYSVASEEGDEDFDER2.0002030002E+21	sp Q14839 CHD4_HUMAN
962	KLSSSSEPYEEDFNDDQSIK2.202E+19	sp Q9H4L7 SMRCD_HUMAN
963	MHDGELEEQEEDDEKSDSEGGDLK20200000000	sp Q9NU22 MDN1_HUMAN
964	TSDSDQAYLVQR20000000000	sp Q96CV9 OPTN_HUMAN
965	DLESCSDDDNQGSK202000000000	sp Q5T3I0 GPTC4_HUMAN
966	LASPSGSTSSGLEVVAPGTSAPGGGPGTLDDSATICR2E+29	sp Q13263 TIF1B_HUMAN
967	MHNLHGTKPPPSEGSDEEEEEDEEDEEER1.00000000002002E+30	sp Q9BQ67 GRWD1_HUMAN
968	VVSDADSDSDAVSDK2000020000000000	sp Q96ST2 IWS1_HUMAN
969	AQAVSEEEEEEGK20000000000	sp Q9GZR7 DDX24_HUMAN
970	AGMSSNQSISSPVLDAVPR100000022000000000	sp Q9UQ35 SRRM2_HUMAN
971	SSSSPELQTLQDILGDPGDK2200000000000000000	sp P49815 TSC2_HUMAN
972	SEPVKEESSELEQPFADTSSVGPDR2000000000000000000	sp Q15424 SAFB1_HUMAN
973	GPAGEAGASPPVR200000	sp Q14151 SAFB2_HUMAN
974	VFDDESDEKEDEEYADEK2000000000000000	sp Q13425 SNTB2_HUMAN
975	DVEDMELSDVEDDGSK2000000000	sp 043719 HTSF1_HUMAN
976	YFGFDLSESEDEDDDCQVER2020000000000000	sp Q5VT52 RPRD2_HUMAN
977	TPASTPVSGTPQASPMIER20002000000	sp Q7Z5K2 WAPL_HUMAN
978	SQEADVQDWEFR20000000000000	sp Q9UHR4 BI2L1_HUMAN
979	TAAGEYDSVSESEDEEMLEIR30002020000000000	sp Q9COC2 TB182_HUMAN
980	GTEASSGTEAATGLEGEEK2000000000000000	sp Q9BZE4 NOG1_HUMAN
981	VEMYSGSDDDDFNK202000000000	sp Q8IY81 SPB1_HUMAN
982	SISNEGLTLNNSHVSK2000000000000000	sp 060841 IF2P_HUMAN
983	ESYTEEIVSEAEHVSGISR20002000000000	sp Q08AD1 CAMP2_HUMAN
984	LNDLSDSNLDEESEDEFK2020000002000000	sp Q5QJE6 TDIF2_HUMAN
985	GNAEGSSDEEGKLVIDEPAK2000000000000000	sp Q96T23 RSF1_HUMAN
986	TSTTGVAATQSPTPR200000	sp P51858 HDGF_HUMAN
987	VSPLEGSEGDGDTD200000200	sp Q8WWI1 LM07_HUMAN
988	TQEDEEISTSPGVSEFVSDAFDACNLNQEDLR2E+23	sp Q9Y606 TRUA_HUMAN
989	ESEVCPVPTNSPPLPPPLPK2020000000000000	sp Q96A49 SYAP1_HUMAN
990	VDSTTCLFPVEEK200000000000	sp Q9ULJ3 ZBT21_HUMAN
991	SPVIGSEVFLPNSNHVASGAGEAEER2E+26	sp Q06210 GFPT1_HUMAN
992	TGSCSELDACPSK200000000000	sp P29590 PML_HUMAN
993	VPAEDETQSIDSEDSFVPGR20020000000000	sp Q9H1H9 KI13A_HUMAN
994	SVSGTDVQECCR20000000000000	sp Q8WZ73 RFFL_HUMAN
995	SVSEINSDELKSGK2000000000	sp P49321 NASP_HUMAN
996	IDEPSTPYHSMGDEDEDACSDTEATEAMPDILAR1.10000000002E+25	sp P82094 TMF1_HUMAN
997	EVAATEEDVTRLPSPTSPFSSLSQDQAATSK22000000000000000	sp P41236 IPP2_HUMAN
998	LPSTSDDCPAIGTPLR200000000020000	sp Q8WWI1 LM07_HUMAN
999	SLYGDVDSPLPTGEAGPPSTR2000000000000000	sp Q15648 MED1_HUMAN
1000	AASADSTTEGTPADGFTVLSTK2.002E+20	sp Q9Y4B5 MTCL1_HUMAN
1001	VKLDDSDDEESK2000000000	sp 075367 H2AY_HUMAN
1002	SVEDDKEGHLVCR20000000000000	sp Q12872 SFSWA_HUMAN
1003	DLLSDLQDISDSER200000	sp P49761 CLK3_HUMAN
1004	GDIVAVNALLDHPEELQPSSESDDEGPELLQVTR202020000000000000	sp P21127 CD11B_HUMAN
1005	NEEDSEGSSDEDEDEDGVSAATFLK2.0022E+21	sp Q9UQ88 CD11A_HUMAN
1006	DSDSEEDLVSYGTGLEPLEEGERPK2.02E+24	sp Q9BPX7 CG025_HUMAN
		sp B5ME19 EIFCL_HUMAN
		sp Q99613 EIF3C_HUMAN
		sp Q9BRR8 GPTC1_HUMAN

1007	VGNESPVQELK20000000	sp Q5JSH3 WDR44_HUMAN
1008	AEENTDQASPQEDYAGFER200000000000	sp Q9NU22 MDN1_HUMAN
1009	ATSSSSGSLSATGR200000000000	sp Q03252 LMNB2_HUMAN
1010	TVSSPPTSPRPGSAATVSASTSNIIPPR2002000000000000	sp 076094 SRP72_HUMAN
1011	LPSTSGSEGVFPR200000000000	sp P29317 EPHA2_HUMAN
1012	SPGSTPTTPTSSQAPQK2000000000000000	sp P35658 NU214_HUMAN
1013	DNQESSDAELSSSEYIK20000200000000	sp Q96JM3 CHAP1_HUMAN
1014	KSSEGGVGVGPGGGDEPPTSPR2E+20	sp Q9Y4H2 IRS2_HUMAN
1015	GLRDSHSSEEDEASSQTDLSTISK2. 2E+19	sp Q5JTV8 TOIP1_HUMAN
1016	VAASPKSPTAALNESLVECPK2020000000000000	sp Q53EZ4 CEP55_HUMAN
1017	SQSDLDDQHDYDSVASDEDTDQEPLR200000000000	sp Q9Y2X7 GIT1_HUMAN
1018	SGDETPGSEVPGDK2000200000000000	sp Q1KMD3 HNRL2_HUMAN
1019	FQEQECPPSPEPTR2000000	sp P62070 RRAS2_HUMAN
1020	VMTIPYQPMPASSPVICAGGQDR200000000000	sp Q15365 PCBP1_HUMAN
1021	DLLSDLQDISDSER200000200000	sp Q9UQ88 CD11A_HUMAN
1022	LQQGAGLESPPQQPEPGAASPQR2000000000020000	sp P21127 CD11B_HUMAN
1023	SSSEDAESLAPR200000000000	sp Q9H6F5 CCD86_HUMAN
1024	DKAPVPPQSPAAAPGGTDEKPSGK2000000000000000	sp Q4G0J3 LARP7_HUMAN
1025	LKLEDDSEESDAEFDEGFK20020000000000	sp Q13200 PSMD2_HUMAN
1026	DDAEESSEPEEENVGNDTK2020000000000000	sp Q03468 ERCC6_HUMAN
1027	STAQQELDGKPASPTPVIVASHTANK2000000000000000	sp Q13823 NOG2_HUMAN
1028	HSVTAATPPPSPTSGESGDLLSNLLQSPSSAK2. 0002E+26	sp P35606 COPB2_HUMAN
1029	FTMDLSDSEDFDFDEK200002000000	sp Q96QU8 XPO6_HUMAN
1030	QPPGTQQSHSSPGEITSSPQGLDNPALLR2E+19	sp P11388 TOP2A_HUMAN
1031	LNHVAAGLVSPSLK200000	sp Q5TDH0 DDI2_HUMAN
1032	GGSYSQAASSDSAQGSVSLTACK2000000000000000	sp Q05519 SRS11_HUMAN
1033	EEGPPPPSPDGASSDAEPEPPSGR20000000000000	sp P01891 1A68_HUMAN
1034	VQGEAVSNIQENTQTPTVQEESEEEEEVDETGVEVK2E+19	sp P01892 1A02_HUMAN
1035	GLNSQSSDDHLNKR200000000000	sp P05534 1A24_HUMAN
1036	GLVAAYSGESDSEEEQER30002020000000	sp Q09160 1A80_HUMAN
1037	GGSYSQAASSDSAQGSVSLTACK20002000000000	sp P10316 1A69_HUMAN
1038	SAPASPTHPLMSPR2000200000000000	sp P30447 1A23_HUMAN
1039	FFTTGSDSESESSLSGEELVTKPVGGNYGK2. 02E+25	sp Q9NTJ3 SMC4_HUMAN
1040	TGSYGALAEITASK200000000000	sp E9PAV3 NACAM_HUMAN
1041	KDSNELSDSAGEEDSADLKR2020000000000000	sp Q13765 NACA_HUMAN
1042	KVELSESEEDKGGK200000000000	sp A8CG34 P121C_HUMAN
1043	ASASGGAQVGGPISSGSSASSVTVTR200000000000	sp Q96HA1 P121A_HUMAN
1044	LASPSGSTSSGLEVVPEGTSAPGGGPGTLDDSATICR2E+36	sp P98175 RBM10_HUMAN
1045	GARPPAAGPGGDEDEDEEDTAPESALDTSLDK20002000020000	sp P05534 1A24_HUMAN
1046	IYHLPDAESDEDEDFKEQTR200000000000	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
		sp P85037 FOXK1_HUMAN
		sp B5ME19 EIFCL_HUMAN
		sp Q99613 EIF3C_HUMAN
		sp 014974 MYPT1_HUMAN
		sp Q9Y6X9 MORC2_HUMAN
		sp Q8IYB3 SRRM1_HUMAN
		sp P02545 LMNA_HUMAN
		sp Q13263 TIF1B_HUMAN
		sp Q27J81 INF2_HUMAN
		sp Q15019 SEPT2_HUMAN

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1048	DHAPSIPTSPFGDSLESSTELR200200000000000000	sp	Q9Y4B5	MTCL1_HUMAN
1049	AASLNLYLNQPSAAPLQVSR200000000000000000	sp	Q9NSK0	KLC4_HUMAN
1050	TSSTSSSVNSQTLNR2000000000000000	sp	Q9NYJ8	TAB2_HUMAN
		sp	014950	ML12B_HUMAN
1051	ATSNVFMFDQSQIQEFK200000000000000000	sp	P19105	ML12A_HUMAN
		sp	P24844	MYL9_HUMAN
1052	QVVSVVQDEEVGLPFEASPESPPPASPDPGVTEIR200000002000000000	sp	Q96TA1	NIBL1_HUMAN
1053	KGGEFDEFVNDTDDDLPIISK20000000000	sp	Q6PD62	CTR9_HUMAN
1054	EPGSGGPQNLSPGGR2000000	sp	Q9HOW8	SMG9_HUMAN
1055	EKGSTLDLSDLEAEK20000000000000	sp	000767	ACOD_HUMAN
1056	APSDTGSLSPWDQDR20200000000	sp	015320	CTGE5_HUMAN
1057	IADPEHDHTGFLTEYVATR20300000	sp	P27361	MK03_HUMAN
1058	MQSLQLDCVAVPSSR200000000000200	sp	Q13546	RIPK1_HUMAN
1059	VAAAAGSGSPPGSPGHDR20002000000	sp	Q9BTU6	P4K2A_HUMAN
1060	SPSVSPSKQPVSTSSK202000000000000000	sp	Q96JG6	CC132_HUMAN
1061	KKPSTDEQTSSAEEDVPTCGYLNLSNSR2. 22E+21	sp	Q8N556	AFAP1_HUMAN
1062	SRLTPVSPESSSTEEK200200000000000	sp	Q13501	SQSTM_HUMAN
1063	GSSGSPAHAESYSSGGGGQK2. 02E+19	sp	P15408	FOSL2_HUMAN
1064	YQPLASTASDNDFVTPEPR202000000000000	sp	Q15021	CND1_HUMAN
1065	SPPPTHSTQLGAPSR200000000000000000	sp	Q14814	MEF2D_HUMAN
1066	LPISDSPDTQEIHVIEQEK200000000000000000	sp	Q8WYP5	ELYS_HUMAN
1067	LASPSGTSSGLEVVAPGTSAPGGGPGTLDDSATICR2. 22E+31	sp	Q13263	TIF1B_HUMAN
1068	SLSPGGAALGYR200000000000	sp	Q96T37	RBM15_HUMAN
1069	GDDDEESDEEAVKK2000000000	sp	043395	PRPF3_HUMAN
1070	SSTPLPTISSAENTR220000000000000000	sp	P42166	LAP2A_HUMAN
		sp	P42167	LAP2B_HUMAN
1071	ATNESEDEIPQLVPIGK2000000000000000	sp	076021	RL1D1_HUMAN
1072	AVPIA VADEGESESEDDLKPR202000000000	sp	Q9Y2K6	UBP20_HUMAN
1073	VLDTSSLTQSAPASPTNK2000200000	sp	Q8N122	RPTOR_HUMAN
1074	TPSDDEEDNLFAPPK2000000000000000	sp	Q9Y4E1	FA21C_HUMAN
		sp	Q641Q2	FA21A_HUMAN
1075	ANSFVGTAQYVSPELLTEK200000000000000000	sp	015530	PDPK1_HUMAN
		sp	Q6A1A2	PDPK2_HUMAN
1076	ASVSQTSPQSASSPR20000000000	sp	Q8TDY2	RBCC1_HUMAN
1077	SLESCLDLR200000000000	sp	Q6ZRV2	FA83H_HUMAN
1078	ALAEAESEELPSDVLNDPYFAEEVK2. 000002E+21	sp	Q9H501	ESF1_HUMAN
1079	GYTSDSEVYTDHGRPGK2000000000000000	sp	Q92538	GBF1_HUMAN
1080	EEGPPPPSPDGASSDAEPEPPSGR200000000000	sp	Q9NTJ3	SMC4_HUMAN
1081	RGSQKSTDSPGADAELPESAAAR2. 0002E+20	sp	Q9Y4K1	AIM1_HUMAN
1082	SPLTILQDDNSPGTLTLR2000000200000000	sp	Q99618	CDCA3_HUMAN
1083	QSFDDNDSELEDKDSK200000000000	sp	060841	IF2P_HUMAN
1084	KLSVPTSDEEDEVPAKPR200020000000000000	sp	Q8NE71	ABCF1_HUMAN
1085	SSGTASSVAFTPLQGLEIVNPQAAEK2. 0002E+20	sp	Q8WWY3	PRP31_HUMAN
1086	RNPWSDDESKSESLEETEPVVIPR2. 00020202E+21	sp	Q02880	TOP2B_HUMAN
1087	GTLDEEEDSADTDDIDHR20000000	sp	Q9HCN4	GPN1_HUMAN
1088	SPVGSQAPQAAAPAPAAHVAGNPGGDAAPAATGTAASLATAAGSEDAEKK2E+52	sp	P16989	YBOX3_HUMAN
1089	TPLSSLSATSNNSTVACIDR22002000000000	sp	P00533	EGFR_HUMAN
1090	SGDETPGSEVPGDK2000000000000000	sp	Q1KMD3	HNRL2_HUMAN
1091	SPPLQLSEAELLNQLQR200000000000000000	sp	015015	ZN646_HUMAN



		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
1092	EDALDDSVSSSSVHASPLASSPVR20000	sp A6NKT7 RGPD3_HUMAN
		sp O14715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
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1094	ALVVPEPEPDSNSQER200000	sp Q5VTR2 BRE1A_HUMAN
1095	TSPSSPAPLPHQEATPR200000000000000	sp P04920 B3A2_HUMAN
1096	ESESEDETTPAAPQLIK202000000000000000	sp O60832 DKC1_HUMAN
1097	KGDVEGSQSQDEGEGSGESER200000000000000	sp Q9UQE7 SMC3_HUMAN
1098	EAEQAASEAAGDTPGSSPSSLYEELGQPPR202000000000000000	sp Q86TC9 MYPN_HUMAN
1099	TSSTCSNESLSVGGTSVTPR2020020000000000000	sp O60343 TBCD4_HUMAN
1100	ALSSDSILSPAPDAR20000020000000	sp Q8IVT2 MISP_HUMAN
1101	DGSDYDPYDFSDTEEEMPQVHTPK200000000000000	sp P25205 MCM3_HUMAN
1102	SRSSSVGSSSYPISPAUSR2220020000000000000	sp Q15149 PLEC_HUMAN
1103	WLDESDAEMELR200000000	sp Q9P035 HACD3_HUMAN
1104	RNSCNVGGGGGFK2000000000000	sp Q7L2J0 MEPCE_HUMAN
1105	LFEESDDKEDEDADGK2000000000000	sp O43719 HTSF1_HUMAN
1106	VPVQEIDDDFFPSSGEEAASVGEVGGGGGR2. 2E+19	sp Q03468 ERCC6_HUMAN
1107	FIIGSVSEDNSEDEISNLVK200020000000000	sp Q13085 ACACA_HUMAN
1108	SSSVGSSSYPISPAUSR2000000	sp Q15149 PLEC_HUMAN
1109	TSQVGAASAPAKESPR2000	sp Q13428 TCOF_HUMAN
1110	SSGSETEQVDFSDR2202000000002000	sp O95251 KAT7_HUMAN
1111	QSQQPMKPISPVKDPVSPASQK20000002000000	sp Q12888 TP53B_HUMAN
1112	SVVSDLEADV2002000000000	sp P11388 TOP2A_HUMAN
1113	ASSLGEIDESSEL2000000000000	sp Q16513 PKN2_HUMAN
1114	IASDEEIQGTK2000000000	sp Q8WX93 PALLD_HUMAN
1115	GVMAVTAVTATAASDR2000	sp Q7Z4V5 HDGR2_HUMAN
1116	IVEPEVVGESDSEVEGDWR200000000000	sp P55081 MFAP1_HUMAN
1117	EDKSPSEESAPTTSPESVSGSVSSGSSGR2E+19	sp P54725 RD23A_HUMAN
1118	VSPASPAGSPADFVHGESLGR2. 0000002E+23	sp Q96C92 SDCG3_HUMAN
1119	VSALEEDMDDVESSEEEEEDEKLER2. 00000000002E+25	sp Q8NAV1 PR38A_HUMAN
1120	DFLSGPDYDHTVGSDEDEEVEGEP202000000000000	sp Q6ZW31 SYDE1_HUMAN
1121	TSPCQSPTISEGPVPSLPIRPSR2. 0002E+22	sp O43426 SYNJ1_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
1122	KGGSYSQAASSDSAQGSVSLTACK2E+20	sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
		sp P01892 1A02_HUMAN
1123	GGAPDPSPGATATPGAPAQSSPDAR200000	sp O95365 ZBT7A_HUMAN
1124	LQSWEETWSLIPEK2000000000000	sp Q7Z736 PKHH3_HUMAN
1125	SIGSAVDQGNESIVAK2000000000000	sp Q9H0H5 RGAP1_HUMAN
1126	LSVPTSDEEDEVPAKPR200020000000000000	sp Q8NE71 ABCF1_HUMAN
1127	TCEERPAEDGSDEEDPDSMEAPTR200000001000000	sp Q9H6Y2 WDR55_HUMAN
1128	GAEAFGDSEEDGEDVFEVEK200000000000000	sp Q99549 MPP8_HUMAN
1129	ASAPSPNAQVACDHCLK20000000000000000	sp Q14258 TRI25_HUMAN
1130	YGPADVEDTTGSGATDSK200	sp P24534 EF1B_HUMAN
1131	TTSSSDLTSSSSGPR20000000000000000	sp Q5QP82 DCA10_HUMAN
1132	MDSDEDEKEGEEK2000000000000	sp Q96ST2 IWS1_HUMAN
1133	NAPAAVDEGSI SPR2000	sp P28715 ERCC5_HUMAN
1134	ASDLEDEESAAR2000000000000	sp Q86VM9 ZCH18_HUMAN
1135	NGFPHPEDCNPSEAASEESNSEIEQEIPEVK200200000000000000	sp Q9NR30 DDX21_HUMAN
1136	SVEDDEEGHLICQSGDVLSAR2E+21	sp P49759 CLK1_HUMAN

1137	SAFTPATATGSSPSPVLGQGEK2000000000	sp Q15811 ITSN1_HUMAN
1138	RPPSPDIVLSDNEQPSSPR200000020000000000	sp Q86YP4 P66A_HUMAN
1139	LTPLPEDNSMNVDDQGDPSDR20000000000000	sp P78527 PRKDC_HUMAN
1140	AAVQELSSSILAGEDPEER200000000000	sp P49768 PSN1_HUMAN
1141	ATLLNVPDLSDSIHSANASER20020002000	sp Q07157 ZO1_HUMAN
1142	VAAAAGSGSPPGSPGHDR200000000000	sp Q9BTU6 P4K2A_HUMAN
1143	SSQSSSQFSGIGR2000000000000000	sp Q92841 DDX17_HUMAN
1144	QVVSVVQDEEVGLPFEASPESPPPASPDPGVTEIR2000020000000000	sp Q96TA1 NIBL1_HUMAN
1145	SSSLGSYDDEQEDLTPAQLTR3000000000000000	sp 094988 FA13A_HUMAN
1146	ASSHSSQTQGGGSVTK22000000000000	sp P02545 LMNA_HUMAN
1147	NYQQNYQNSSEGEKNEGSESAPEGQAQQR2E+19	sp P67809 YBOX1_HUMAN
1148	APVQPQQSPAAAPGGTDEKPSGK2000000000000000	sp Q13200 PSMD2_HUMAN
1149	AAFTTPDHAPLSPQSSVASSGSEQTEEQGSSR200000000000000000	sp Q5PRF9 SMAG2_HUMAN
1150	RPMEEDGEKSPSK20000	sp Q12906 ILF3_HUMAN
1151	TDCEPVGLEPAVEQSSAASELAEASSEELAEAPTEAPSPEAR200000	sp Q15424 SAFB1_HUMAN
1152	DPSPEADAPVLGSPEK200000000020000	sp Q14151 SAFB2_HUMAN
1153	KLSSSEPYEEDEFNDDQSIK2200300000000000000	sp 015234 CASC3_HUMAN
1154	SPHSPSPATPPSQSSSCSDAPMLSTVHSAK2.002020002E+31	sp Q9H4L7 SMRCD_HUMAN
1155	VNSNGKESPGSSEFFQEAVSHGK2.00002E+21	sp Q9NYF8 BCLF1_HUMAN
1156	QENCGAQQVPAGGTSTPPSSPVR2000020000	sp Q8N108 MIER1_HUMAN
1157	ILATPPQEDAPSVDIANIR2000000000	sp Q96G46 DUS3L_HUMAN
1158	EVSDDEAEKEDKEEEK2000000000000000	sp P29401 TKT_HUMAN
1159	ELEKPIQSKPQSPVIQAAAVSPK2000000002000	sp P07900 HS90A_HUMAN
1160	TRPGSFQSLSDALSDTPAK2000000000000000	sp Q9Y6D5 BIG2_HUMAN
1161	ISFPGSSESPLSSK2000000	sp Q9H9C1 SPE39_HUMAN
1162	SDSGGSSEPFDR200000000000	sp Q15911 ZFH3_HUMAN
1163	IKNENTEGSPQEDGVELEGLK20000000000000	sp P48634 PRC2A_HUMAN
1164	HTGPNSPDTANDGFVR200000000000	sp P11388 TOP2A_HUMAN
1165	SPSPAPAPAAAAAGPPTR2.02E+19	sp P31943 HNRH1_HUMAN
1166	VISDSESDIGGSDVEFKPDK2.020200002E+19	sp P55795 HNRH2_HUMAN
1167	LPQSSSESSPPSPQPTK2000000	sp Q9H7N4 SFR19_HUMAN
1168	RPDYAPMESSDEEDEFQFIK2200000000000000	sp P52701 MSH6_HUMAN
1169	AKSPQPPVEEEDHFDDTVVCLDTYNCDLHFK2E+30	sp Q9UQ35 SRRM2_HUMAN
1170	TPPPPTLSPAAITVGR20000002000000000	sp P55081 MFAP1_HUMAN
1171	REESPMDVDQSPSAQDTQSIASDGTQGEK2E+20	sp Q00839 HNRPU_HUMAN
1172	VDEDSAEDTQSNQDGK200000000000	sp Q8NDX5 PHC3_HUMAN
1173	LDDAHSLSGAGEGYEPISDDELDEILAGDAEK2.0000030002E+25	sp Q7Z6Z7 HUWE1_HUMAN
1174	SASSGAEGDVSSER200000000000	sp Q9H2G2 SLK_HUMAN
1175	TIGGGDDSFNTFFSETGAGK20000000000000	sp Q5T200 ZC3HD_HUMAN
1176	SQSSEGVSSLSPPSNSLETQSQSLSR2E+24	sp Q8TEA8 DTD1_HUMAN
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1178	TNSPAYSDISDAGEDGEGK200020020000000000	sp Q9BQE3 TBA1C_HUMAN
1179	SDISDQEEDEESEGCVPVINLSK2.002E+23	sp Q71U36 TBA1A_HUMAN
1180	APAGQEEPPTPPSSPLSAEQQLDR2000200000000000	sp Q13748 TBA3C_HUMAN
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1182	GSYDGLGGPIITTVQVITPK20000000000000000000	sp 095155 UBE4B_HUMAN
1183	DQTASAPATPLVNK20002000000	sp 015541 R113A_HUMAN
1184	LTSVSSSVDFDQR200220000000	sp 015014 ZN609_HUMAN
		sp Q15154 PCM1_HUMAN
		sp P13051 UNG_HUMAN
		sp P04792 HSPB1_HUMAN
		sp P61978 HNRPK_HUMAN
		sp Q53SF7 COBL1_HUMAN
		sp Q5T3F8 CSCL2_HUMAN

1185	SAPASPTHPLMSPR2000200000010000	sp P85037 FOXK1_HUMAN
1186	KPPAPSPVQSQSPSTNWSPAVPVK2.00000002E+19	sp Q8IYB3 SRRM1_HUMAN
1187	AQPFGFIDSDDTDAEEER2000000000	sp Q14676 MDC1_HUMAN
1188	QASTDAGTAGALTPQHVR200000000000	sp P46937 YAP1_HUMAN
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1191	KISENSYSLDDLEIGPGLSSSTFDSEK2E+26	sp Q9UHB6 LIMA1_HUMAN
1192	EMLMEDVGSEEEQEEDEAPFQEK1.0100002E+23	sp Q9H1E3 NUCKS_HUMAN
1193	NGFPHPPEPCNPSEAASEESNSEIEQEIPVEQK20000200000000000000	sp Q9NR30 DDX21_HUMAN
1194	GSLDESSLGFGYPK2000000000000000	sp Q9BUZ4 TRAF4_HUMAN
1195	NLTANDPSQEYVASEGEEDPEVEFLK2000000000000000	sp Q86X95 CIR1_HUMAN
1196	GSISSTSEVHSPNVGLR202000000000000000	sp Q8WWQ0 PHIP_HUMAN
1197	TVSASSTGDLPK200200000000	sp P78362 SRPK2_HUMAN
1198	SSSVGSSSYPISPAISR22000000000000000000	sp Q15149 PLEC_HUMAN
1199	EKQSSEEEKETR220000000000	sp Q92504 S39A7_HUMAN
1200	SQDQDSEVNELSR2000000000	sp Q86VM9 ZCH18_HUMAN
1201	VIHFVSGETMEEYSTDEDEVDGLEK100022000000000000	sp Q8N128 F177A_HUMAN
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1203	TADSSSEDEEEYVVEK220200000000000000	sp P45973 CBX5_HUMAN
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1207	VNFSEGETEEDDQDSSHSSVTTVK2.00002E+22	sp Q5JTV8 TOIP1_HUMAN
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1209	SASSDTSEELNSQDSPPK2200000000000000	sp 014745 NHRF1_HUMAN
1210	TPENLITEIR200000000000	sp Q7Z4H7 HAUS6_HUMAN
1211	YSPSQNSPIHHIPSR20020000000000	sp Q9NYF8 BCLF1_HUMAN
1212	HNDIVDSDSAEDR2020000000	sp P78316 NOP14_HUMAN
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1215	EISDDEAEEKGEK20000000000000	sp P08238 HS90B_HUMAN
1216	SRSRSTPPSSAGLGSNSAPPIDSR2.020202E+26	sp Q58FF7 H90B3_HUMAN
1217	TDDTAQVTSDYETNNNSDSSDIVQNEDETECLR2.000202E+21	sp Q8IWX8 CHERP_HUMAN
1218	AGEPNPDAEEANSPDVTAGCDPAGVHPPR2E+25	sp Q14432 PDE3A_HUMAN
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1220	SCSSPAVSAVSQLPLSPK20000000000002000	sp Q92945 FUBP2_HUMAN
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1222	TPSSSSTLAYSPR220000000000	sp Q6IQ22 RAB12_HUMAN
1223	QESDPEDDDVKKPALQSSVATSK2E+22	sp Q6Q0C0 TRAF7_HUMAN
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1225	ESVPEFPLSPPK20000	sp Q9Y4H2 IRS2_HUMAN
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1228	SSGFISELPSEEGK2000000000000000	sp P18615 NELFE_HUMAN
1229	KGSSSVCSVASSSDISLGSTK2E+20	sp Q5VZK9 LR16A_HUMAN
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1231	SGSSSPDSEITELKFPSINH2.22002E+19	sp P0DJ93 SIM13_HUMAN
1232	GYSPYSVSGSGSTAGSR2000000000000200	sp P17812 PYRG1_HUMAN
1233	VSEEQTPPSPAGAMSTAMGR2000000000000000	sp Q15149 PLEC_HUMAN
1234	ADEPSSEESDLEIDK200000000000	sp Q16666 IF16_HUMAN
1235	HQASDSENEEPPKPR20200000000000	sp P50502 F10A1_HUMAN
1236	EAAAGIQWSEEEETEDEEEKEVTPESGPPK2.0002E+22	sp Q8IZP2 ST134_HUMAN
		sp Q96ST2 IWS1_HUMAN
		sp P46087 NOP2_HUMAN

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1238	HGSGADSDYENTQSGDPLLGLEGK203000000000000000	sp Q9Y2X7 GIT1_HUMAN
1239	GSSGSPAHAESYSSGGGGQQK200000000000000000	sp P15408 FOSL2_HUMAN
1240	GQESSDQEVDVESIDFSK202000000000000000	sp Q96JM3 CHAP1_HUMAN
1241	SPLGGGGGSGASSQAACLK2E+19	sp Q68DK7 MSL1_HUMAN
1242	QEPLEEDSPSSSAGLDK200000000000	sp P15408 FOSL2_HUMAN
1243	VCSSDSDSSLQVVK220200000000	sp Q9NSI6 BRWD1_HUMAN
1244	FSGEEGIEDESGTENR200000000000000000	sp Q9Y2W1 TR150_HUMAN
1245	HGSGADSDYENTQSGDPLLGLEGK2.000003E+22	sp Q9Y2X7 GIT1_HUMAN
1246	SPQPVCSPAGSEGTPK20002000000	sp Q6KC79 NIPBL_HUMAN
1247	SSTPPGESYFGVSSLQK200000000000000000	sp Q9UQ35 SRRM2_HUMAN
1248	SQSSEGVSSLSSSPNSLETQSQSLSR2.02E+19	sp O95155 UBE4B_HUMAN
1249	EVSSRPSTPGLSVVSGISATSEIPNKIEDLR2.2E+26	sp Q14C86 GAPD1_HUMAN
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1252	SSGHSSSELSPDAVEK2000020000000	sp Q9UQ35 SRRM2_HUMAN
1253	ASQEANLLTLAQK20000000000000	sp Q6PJG2 EMSA1_HUMAN
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1255	SPPPESVDTPSTK2000000000000000	sp P46013 KI67_HUMAN
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1258	DTVIVSEPSEDEESQGLPTMAR200200000000000000	sp Q9H4L7 SMRCD_HUMAN
1259	RQSSGSATNVASTPDNR2002000000000000	sp Q7Z460 CLAP1_HUMAN
1260	GLLYDSDEEDEERPAR200000000000	sp P49736 MCM2_HUMAN
1261	ELVEPLTPSGEAPNQALLR20000000000000	sp P00533 EGFR_HUMAN
1262	SEGSEYEEIPK200200000000	sp P48634 PRC2A_HUMAN
1263	TIASDSEEEAGKELSDK2020000000000000	sp Q96ST2 IWS1_HUMAN
1264	TQPDGTSVPGEPASPISR2000000	sp Q14980 NUMA1_HUMAN
1265	SVGGSGGGSGFDNLVTR20000000000000	sp P02545 LMNA_HUMAN
1266	GEDVPSEEEEEENGFEEDR2000000000000000	sp Q9ULX3 NOB1_HUMAN
1267	ADRSSLPLDFNDEFSDLGVVQQR200000000000	sp Q96HH9 GRAM3_HUMAN
1268	LSFSDSESDNSADSSLPSR202020000000000000	sp Q03111 ENL_HUMAN
1269	NKSTESLQANVQR20000000000000	sp P26373 RL13_HUMAN
1270	SSGSSSSGLGTVSNPASQR2000000	sp Q9UPP1 PHF8_HUMAN
1271	QEEESPVERPLK20200000000	sp Q96SB4 SRPK1_HUMAN
1272	FIIGSVSEDNSEDEISNLVK202000200000000000	sp Q13085 ACACA_HUMAN
1273	SGSSSPDSEITELK22200000000000	sp P17812 PYRG1_HUMAN
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1275	KPPAPSPVQSQSPSTNWSPAVPVK2020000000000000	sp Q8IYB3 SRRM1_HUMAN
1276	KASQQSNQIQTR20000000000000	sp Q05D32 CTSL2_HUMAN
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1278	TADSSSEDEEEYVVEK2220000000000000	sp P45973 CBX5_HUMAN
1279	NLEHLSSFSSDEDDPGYSQDAYK2200000000000000	sp Q2KHR3 QSER1_HUMAN
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1282	KTSSVSSISQVSPER2200000000000000	sp O15530 PDPK1_HUMAN
1283	GGSTGGGGGFDPPPAYHEVDAEK2E+22	sp P42356 PI4KA_HUMAN
1284	RPASPSSPEHLPATPAESPAQR2.002E+19	sp Q9UPU5 UBP24_HUMAN
1285	KLSSSSEPYEDEFNDDQSIK200300000000000000	sp Q9H7L9 SDS3_HUMAN
1286	HDYDDSSSEEQAEDSYEASPGSETQR2.20002E+21	sp Q9H4L7 SMRCD_HUMAN
1287	QSNASSDVEVEEK2000000000	sp P98175 RBM10_HUMAN
1288	NVLSSLAVYAEDSEPESDGEAGIEAVGSAAEEK2.0003E+29	sp O75475 PSIP1_HUMAN
1289	LQALVNSLCAGQSP200	sp Q9UHR5 S30BP_HUMAN
		sp Q6XQ6 PNCB_HUMAN

1290	VGPAGDEEEEEEEELSSSCEEEGEEDR222000000000	sp P36915 GNL1_HUMAN
1291	ALEQQEAESDSSDTEEKDDDDDEEDVGKR2. 2E+20	sp Q9BXY0 MAK16_HUMAN
1292	SPAPGSPDEEGGAEAPAAGIR2. 00002E+21	sp O15061 SYNEM_HUMAN
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1296	TLSSSAQEDIIR20000000000	sp Q9H788 SH24A_HUMAN
1297	ATAPQTQHVSPMR20000	sp P29692 EF1D_HUMAN
1298	DAVFDGSSCISPTIVQQFGYQR2000000000000	sp P04049 RAF1_HUMAN
1299	SASPPPATSSSSSSR2020000000000000	sp Q9H7N4 SFR19_HUMAN
1300	EPHSPADQPEQQAESTL TSAETR2E+20	sp Q9H2Y7 ZN106_HUMAN
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1302	EVSSRPSTPGLSVVSGISATSEIPNK2. 2E+21	sp Q14C86 GAPD1_HUMAN
1303	NGFPHPEDCNPSEAASEESNSEIEQEIPVEQK20200000000000	sp Q9NR30 DDX21_HUMAN
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1305	TVSASSTGDLPK20002000000	sp P78362 SRPK2_HUMAN
1306	VSDPISTSESSEEEEEAEATAK2. 0002220E+22	sp Q13428 TCOF_HUMAN
1307	KEESEESDDDMGFLFD200000000000	sp P05386 RLA1_HUMAN sp P05387 RLA2_HUMAN
1308	EDKSPSEESAPTTSPESVSGVPSGSSGR2000000000000000000	sp P54725 RD23A_HUMAN
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1310	SAPASPTHPLMSPR2000000000000000	sp P85037 FOXK1_HUMAN
1311	QLSLEGLGVEDLK2000000000000000	sp Q9UMZ2 SYNRG_HUMAN
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1313	VNFSEEGETEEDDQDSSHSSVTTVK2E+22	sp Q5JTV8 TOIP1_HUMAN
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1316	HEVSASTQSTPASSR2000000	sp Q01082 SPTB2_HUMAN
1317	KDSSGSVSPNTLSQEEGDQICLYHIR2. 002E+24	sp Q9H0J9 PAR12_HUMAN
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1320	SSSEYTSQSFQSR200000000000	sp Q9BTU6 P4K2A_HUMAN
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1388	SEGSEYEEIPKR20020000000000	sp Q9H0W8 SMG9_HUMAN
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1411	LSLNNDIFEANSDDQSETK20200000000	sp Q99549 MPP8_HUMAN
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1414	KTSDANETEDHLESLICK20000000000000000	sp Q09161 NCBP1_HUMAN
1415	YLATSQPRPDSSGSH200	sp Q9C0H2 TTYH3_HUMAN
1416	SGSPSDNSGAEEMEVSLAKPK2.0200002E+21	sp P31749 AKT1_HUMAN
1417	SNCLGTDEDSQDSSDGIPSAAPR2000200000000000000	sp Q92993 KAT5_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
1418	KGGSYSQAASSDSAQGS DVSLTACK2002000000	sp P01892 1A02_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
1419	LSHEDDHILEDAGSSDISSDAACTNPNK200200000000000	sp Q92576 PHF3_HUMAN
1420	NAIASDSEADSDTEVPK202000200000000	sp Q8WVC0 LEO1_HUMAN
1421	SSLSSHSHQSQIYR202000000000000	sp O15027 SC16A_HUMAN
1422	GIPLATGDTSPPELLPGAPLPPPK20000000000000000	sp O75821 EIF3G_HUMAN
1423	LFPEFDSDYDEVPEEGPGAPAR20000000000000000	sp Q96P48 ARAP1_HUMAN
1424	SPSPPDGSPAATPEIR200000000000000	sp O00499 BIN1_HUMAN
1425	GSISSTSEVHSPNVGLR220000000000000	sp Q8WWQ0 PHIP_HUMAN
1426	SGSMDPSGAHPSVR2100000000000	sp Q07666 KHDR1_HUMAN
1427	TSPSGGTWSSVVGVPVR20000000000000000	sp Q99700 ATX2_HUMAN
1428	MNSYPYLADR200000000	sp Q9Y580 RBM7_HUMAN
1429	RPGAQPLPPPPPSQSPEPTPHPR20000000000	sp Q9UIS9 MBD1_HUMAN
1430	LQDISSELLATGAGLSESEAE PDGHNITELPQAVAGR2.02E+23	sp Q9NQ55 SSF1_HUMAN
1431	SKSEEHAEDSVMDDHFR20000000000000000	sp Q8NC51 PAIRB_HUMAN
1432	SQSSDTEQQSPTSGGGK200000000000000	sp P23588 IF4B_HUMAN
1433	TPLLSLSATSNNSTVACIDR202200000000000	sp P00533 EGFR_HUMAN
1434	SPLTILQDDNSPGTLTLR2000000000200000000	sp Q99618 CDCA3_HUMAN
1435	QEKPAEKPAETPVATSPTATDSTSGDSSR200000000000000	sp P54727 RD23B_HUMAN
1436	DSQDASAEQSDHDDEVASLASASGGFGTK2E+20	sp Q6P2E9 EDC4_HUMAN
1437	ALDISLSSGEEDEGDEEDSTAGTTK2.022E+21	sp Q8IY81 SPB1_HUMAN
1438	SSGSETEQVVD FSDR2000000000000	sp O95251 KAT7_HUMAN
1439	RPNEDSDEDEEK20000000	sp Q12996 CSTF3_HUMAN
1440	SPDLLDASAVSATS R200000000000000	sp O60499 STX10_HUMAN
1441	GSISSTSEVHSPNVGLR220000000000000	sp Q8WWQ0 PHIP_HUMAN

1442	VTTEIQLPSQSPVEEQSPASLSSLR200000000000000000	sp Q8WWI1 LMO7_HUMAN
1443	LFDEEEDSSEKLFDDSDER2200000020000	sp O43719 HTSF1_HUMAN
1444	GAPSSPATGVLSPQGK2000000000000000	sp Q5SRE5 NU188_HUMAN
1445	SSSMSYVDGFIGTWPK2000000000000000	sp Q08AD1 CAMP2_HUMAN
1446	SDKSPDLAPTPAQSTPR2000000000000000	sp Q9BY44 EIF2A_HUMAN
1447	LHKPPADSGVDLR2000000	sp O15427 MOT4_HUMAN
1448	SSSSSSGGGLLPYPR2002000000000000	sp O60293 ZC3H1_HUMAN
1449	SPGSTPTTSSQAPQK200200000000000000	sp P35658 NU214_HUMAN
1450	DQQPSGSEGEDDDAEAAALKK2020000000000000	sp Q9NXG2 THUM1_HUMAN
1451	ERESEDELEEANGNNPIDIEVDQNK2E+22	sp Q9UKL0 RCOR1_HUMAN
1452	EESSEAEAPVQR2000000000000000	sp Q9BW71 HIRP3_HUMAN
1453	VTNDISPESSPGVGR20002000000	sp Q15154 PCM1_HUMAN
1454	IALESEGRPEEQMESDNCSGGDDWTHLSSK200020000000000000	sp Q13501 SQSTM_HUMAN
1455	GIPLATGDTSPPELLPGAPLPPPK2E+20	sp O75821 EIF3G_HUMAN
1456	DDAEESSEPEEENVGNDTK222000000000000000	sp Q13823 NOG2_HUMAN
1457	AGMSSNQSISPVLDVPR220000020000000000	sp Q9UQ35 SRRM2_HUMAN
1458	SIQDLTVGTGEPGVSSR20000000000000000000	sp O43318 M3K7_HUMAN
1459	SPTGPSNSFLANMGTVAHK2E+20	sp Q96I25 SPF45_HUMAN
1460	KNSTDLDAPEDPTSPK2000000000002000	sp Q96RK0 CIC_HUMAN
1461	VMTIPYQMPASSPVICAGGQDR20000000000000	sp Q15365 PCBP1_HUMAN
1462	VEEDLKADEPSSEESDLEIDK220020000000	sp P50502 F10A1_HUMAN
1463	TGSVQTPPDLESSEEVDAEAGEPEVVGPK2.0020000022E+27	sp Q8IZP2 ST134_HUMAN
1464	QSAGPNSPTGGGGGGGGGTR2E+20	sp P28290 SSFA2_HUMAN
1465	VISDESSEDIGGSDVEFKPDTK2.0202E+19	sp A7E2V4 ZSWM8_HUMAN
1466	SSTPLPTISSAENTR200000000000000000	sp P52701 MSH6_HUMAN
1467	NGFPHPEDCNPSEAASEESNSEIEQEIPVEQK200202000000000000	sp P42167 LAP2B_HUMAN
1468	HGSGADSDYENTQSGDPLLGLEGK2E+22	sp P42166 LAP2A_HUMAN
1469	QSSGSATNVASTPDNR2000000000000000	sp Q9NR30 DDX21_HUMAN
1470	SNPSIQATLNK2000000000000000	sp Q9Y2X7 GIT1_HUMAN
1471	NSSDSAIDNPKPNK2000000000000000	sp Q7Z460 CLAP1_HUMAN
1472	GPAFNMA SPESDFGINLK20000000000000	sp Q6UUV7 CRT3_HUMAN
1473	AESPESAIESTQSTPQK20000000000020000	sp P46100 ATRX_HUMAN
1474	SSTPLPTISSAENTR220000000000000000	sp Q09666 AHNK_HUMAN
1475	GNAEGSSDEEGKLVIDEPAK2000000000000000	sp Q9NTI5 PDS5B_HUMAN
1476	GPKPEPPGSGSPAPPR202000000	sp P42167 LAP2B_HUMAN
1477	SVASNQSEMEFSSLQDMPK2.00200001E+19	sp P42166 LAP2A_HUMAN
1478	ALFKPPEDSQDDESDDAEQQTTK20200000000000	sp P51858 HDGF_HUMAN
1479	TDDTAQVTSYETNNNSDSSDIVQNEDETECLR3.000000022E+23	sp Q9UKJ3 GPTC8_HUMAN
1480	APYQENDGYCPDLELSDSEAESDGNKEK20200000000000	sp Q8N163 CCAR2_HUMAN
1481	HSSISPSTLTLK20200000000	sp Q13769 THOC5_HUMAN
1482	LQPSSPENSLDPFPPR2000000000000000	sp Q14432 PDE3A_HUMAN
1483	TGSCSELDACPSK2000000000000000	sp Q92613 JADE3_HUMAN
1484	ATLLNVPDLSDSIHSANASER20020000000	sp Q14004 CDK13_HUMAN
1485	SKSDGEAKPEPSPR2000000000000000	sp Q8WWM7 ATX2L_HUMAN
1486	TPSPPPPPIEDIALGK202000000000000000	sp Q9H1H9 KI13A_HUMAN
1487	NASASFQLEDKK202000000000	sp Q07157 ZO1_HUMAN
1488	VNSGDTEVGSSLLR2000000000000000	sp P26358 DNMT1_HUMAN
1489	KPATPAEDDEDDIDLFGSDNEEDKEAAQLR2E+29	sp Q5T200 ZC3HD_HUMAN
1490	APSPPLSQVFQTR2000000000000000	sp Q99543 DNJC2_HUMAN
1491	TPLALAGSPTPK200000	sp O75592 MYCB2_HUMAN
1492	SCDELSPVSPTQGGYPSEPTR2000000000000000	sp P29692 EF1D_HUMAN
		sp Q9NRA8 4ET_HUMAN
		sp O95785 WIZ_HUMAN
		sp Q9HCD6 TANC2_HUMAN



1493	SLSSPTVTLAPLEGAK2000000000000000	sp Q96PU5 NED4L_HUMAN
1494	EVSSRPSTPGLSVVSGISATSEIPNK2.002E+24	sp Q14C86 GAPD1_HUMAN
1495	ELQGDGPPSSPTNDPTVK2000000000	sp Q12873 CHD3_HUMAN
1496	SSSMSLTGAYASGIPSSSR2200000000000000	sp Q96HA1 P121A_HUMAN
1497	ENSTVDFSK20000000	sp Q9Y6M7 S4A7_HUMAN
1498	RQSSGSATNVASTPDNR2000000000000000	sp Q6U841 S4A10_HUMAN
1499	DSGNWDTSGSELSEGELEK20200200000000	sp Q7Z460 CLAP1_HUMAN
1500	SKFDSDEEEEDTENVEAASSGK200000000000000000	sp 075400 PR40A_HUMAN
1501	ASSLGEIDESSEL2000000000000000	sp Q8TF01 PNISR_HUMAN
1502	EGEETEGSEEDDKENDK20020000000000	sp Q16513 PKN2_HUMAN
1503	GPSPSSPTPPAAAAPAEQAPR2.00002E+19	sp P49321 NASP_HUMAN
1504	VEEDLKADEPSSEESDLEIDK220000000000	sp Q86VQ1 GLC11_HUMAN
1505	SPSSDLDTAEGDDFELLDQSELSQLDPASSR2.0002E+29	sp P50502 F10A1_HUMAN
1506	TASPLPLRPDAPVEKSPEETATQVPSLESITLK2.0000000000002E+31	sp Q8IZP2 ST134_HUMAN
1507	VSPSPQESLSSSK2000000000000000	sp Q86VR2 F134C_HUMAN
1508	ALSSGGSITSPPLSPALPK200000200000000000	sp Q9HCK8 CHD8_HUMAN
1509	SSDQPLTVPVSPK2000	sp Q5JSH3 WDR44_HUMAN
1510	SSSPVTELASR220000000000	sp Q6Y7W6 PERQ2_HUMAN
1511	QPPPLAPQSPQGGVMGGSNSNQQQMR2E+19	sp Q9ULW0 TPX2_HUMAN
1512	VVPGQFDDADSSDSEN2000000	sp Q9UQ35 SRRM2_HUMAN
1513	DFQDYMEPEEGCQSPQR10000000020000	sp P46937 YAP1_HUMAN
1514	WVEENVSSVTDVALPALLDSDEER200000	sp Q9BRS2 RIOK1_HUMAN
1515	RGPNYTSGYGTNSELSNPSETESER200000000000	sp 043237 DC1L2_HUMAN
1516	LGSSGLSASSIQA AVR200000000000	sp Q8NBN3 TM87A_HUMAN
1517	AFGESSTESDEEEEGCGHHCVR2.2202E+20	sp P51114 FXR1_HUMAN
1518	YGLLANTEDPTEMASLDSDEETVFESR1020020000000000	sp Q7Z6Z7 HUWE1_HUMAN
1519	EKPSFIDLETFNAGSDVGSSEYSEPFMGGAR20002000000000000000	sp 060927 PP1RB_HUMAN
1520	SASPDDLGSNNWEAADLGNEER20000000000000	sp Q9BVV8 CS024_HUMAN
1521	SQLDDHPESDDEENFIDANDEDMEK200000000000000000	sp Q9BRY0 S39A3_HUMAN
1522	EGSEFSFSDGEVAEK20020000000000	sp 000193 SMAP_HUMAN
1523	LGSTSGEESDLER222000200000	sp Q03701 CEBPZ_HUMAN
1524	NSPSNLSSSDTG SVGGTYR2E+19	sp Q9BVS4 RIOK2_HUMAN
1525	MDRTPPPPTLSPA AITVGR1.0020000002E+19	sp Q9BW71 HIRP3_HUMAN
1526	KEESEESDDMGFLFD200200010000000	sp 043166 SI1L1_HUMAN
1527	ASTASPCNNNINAATAVALQEPR2.02E+21	sp Q8NDX5 PHC3_HUMAN
1528	SYLEGSSDNQLKDESETPVDDR2.300002E+22	sp P05386 RLA1_HUMAN
1529	TSPLVSQNEQGSTLR2000000000000000	sp P05387 RLA2_HUMAN
1530	TNSPAYSDISDAGEDGEGK2.000002002E+19	sp Q71RC2 LARP4_HUMAN
1531	SSVSDAPVHITASGEPVPISEESELDQK2E+26	sp Q9UHI6 DDX20_HUMAN
1532	DELHIVEAEAMNYEGSPIK20000	sp Q15652 JHD2C_HUMAN
1533	EDKSPSEESAPTTSPESVSGSVSSGSSGR200000000000000000	sp 015014 ZN609_HUMAN
1534	VISDESIGGSDVEFKPDTK2.020000002E+19	sp 014646 CHD1_HUMAN
1535	LLSTDAEAVSTR200000000000	sp P06748 NPM_HUMAN
1536	GDSETDLEALFNAVMNPK20000000000010000	sp P54725 RD23A_HUMAN
1537	SAGALEEGTSEGQLCGR200000000000000000	sp P52701 MSH6_HUMAN
1538	SSTDSLPGISR2002000000000	sp Q15545 TAF7_HUMAN
1539	AQLSPGIYDDTSAR200000000000	sp P46937 YAP1_HUMAN
		sp Q8N2F6 ARM10_HUMAN
		sp Q9HCD5 NCOA5_HUMAN
		sp 075376 NCOR1_HUMAN

1540	GGSYSQAASSDSAQGSVDVSLTACK2000000000	sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
		sp P01892 1A02_HUMAN
1541	MALPPQEDATASPPR20000	sp Q9UQ35 SRRM2_HUMAN
1542	HQSESQGVGLSDK200000002000	sp P38398 BRCA1_HUMAN
1543	DEEDGKDSDEAEDAELYDDLCPACDK2E+20	sp Q5F1R6 DJC21_HUMAN
1544	SPQPPVEEEDHFDDTVVCLDTYNCDLHFK2E+30	sp Q00839 HNRPU_HUMAN
1545	LPSTSGSEGVVFR200020000000	sp P29317 EPHA2_HUMAN
1546	ATLSSTSGLDLMSSEGEIISPQR20000	sp Q8WWI1 LMO7_HUMAN
1547	THSTSSSLGSGESPFSSR200000000	sp Q9UGV2 NDRG3_HUMAN
1548	SLSDSESDSK200000000000	sp Q13185 CBX3_HUMAN
1549	SPPLSPVGTTPVK2000200000000	sp Q9BVC5 ASHWN_HUMAN
1550	SGSALLQSQSSTEDPKDEPAELKPDSEDLSSQSSASK2E+27	sp Q5VTR2 BRE1A_HUMAN
1551	RTSTPVIIMEGVQEETDTR2000000000000000	sp Q8NFC6 BD1L1_HUMAN
1552	NVLSSLVAEADSEPESDGEAGIEAVGSAAEK3.0002E+25	sp Q9UHR5 S30BP_HUMAN
1553	LIHGEDSDSEGEER20200000000	sp Q9C0C9 UBE20_HUMAN
1554	STSTPNVHMVSTTLPVDSR2000000000000000	sp P04049 RAF1_HUMAN
1555	KPPAPSPVQSQSPSTNWSAPVVKK202000000000000	sp Q8IYB3 SRRM1_HUMAN
1556	SQDATFSPGSEQAESPGPIVSR200000000	sp Q86WB0 NIPA_HUMAN
1557	SDVQEESESDTDDNKDSAAFEDNEVQDEFLEK2.00002E+27	sp Q9UIG0 BAZ1B_HUMAN
1558	DSGNWDTSGSELSEGELEKR2202002000000000	sp O75400 PR40A_HUMAN
1559	SPGVAAAEDGGLK2000000000000000	sp Q9BTE7 DCNL5_HUMAN
1560	SNTPESIAETPPAR2000000000000	sp Q9Y4H2 IRS2_HUMAN
1561	SRSPLDKDTPPSASVVGASVGGHR23000000000000000	sp Q96T37 RBM15_HUMAN
1562	ERSPALKSPLQSVVVR2000020000000000	sp Q9Y2W1 TR150_HUMAN
1563	THSTSSSLGSGESPFSSR2220020000000000	sp Q9UGV2 NDRG3_HUMAN
1564	GEEGSDDETENGPVK2000000000000000	sp Q6PD62 CTR9_HUMAN
1565	SGDEEFKGEDELCDSGR200000000000000000	sp O43823 AKAP8_HUMAN
1566	VSMPDVELNLKSPK1000000002000	sp Q09666 AHNK_HUMAN
1567	DPSASPGDAGEQAIR2000000000000	sp P18206 VINC_HUMAN
1568	QASLDGLQQLR20000000000	sp Q3MI16 TBC25_HUMAN
1569	EAEQAASEAAGDTPPGSSPSSLYEEPLGQPPR2.0000002E+28	sp Q86TC9 MYPN_HUMAN
1570	ALSSGGSITSPPLSPALPK200000020000000000	sp Q6Y7W6 PERQ2_HUMAN
1571	ASVLDTSMSAGSGSPSK10000000200	sp Q9UPQ0 LIMC1_HUMAN
1572	SYLEGSSDNQLKDESTPVDDR3.00022E+21	sp Q9UHI6 DDX20_HUMAN
1573	GPLSPPTAGLEESSGQGPSSPVALLGQVQDFQQAECQPK2.00002E+27	sp A1L390 PKHG3_HUMAN
1574	AAVLSDESEDEEKASAK2020000000000	sp Q96ST2 IWS1_HUMAN
1575	KYSDSSLPPSNSGK2000000000000	sp Q68CP9 ARID2_HUMAN
1576	TAFDEAIAELDTLSEESYK2000000	sp P63104 1433Z_HUMAN
1577	KKASSDSEDSSEEEEEVQGPPAK2.22000022E+21	sp Q14978 NOLC1_HUMAN
1578	GSVSPSTPSPPKPTGIFQTSANSSFEVVK2.0022E+27	sp O15027 SC16A_HUMAN
1579	NASASFQLEDK200000000000	sp Q99543 DNJC2_HUMAN
1580	NTSSDNSDVEVMPAQSPR2002000010000000	sp Q9HCK8 CHD8_HUMAN
1581	SFNGTDNHIR200000000000	sp Q5T0W9 FA83B_HUMAN
1582	IPNYQLSPTK20000	sp O60934 NBN_HUMAN
1583	SPSPEPIYNSEGK202000000000000	sp Q15637 SF01_HUMAN
1584	AGMSSNQSISSPVLDVAVPR100000002000000000	sp Q9UQ35 SRRM2_HUMAN
1585	LASTSDIEEK20000000	sp O14974 MYPT1_HUMAN
1586	MHNLHGTPPPSEGSDEEEEEDEEDEEER2.002E+19	sp Q9BQ67 GRWD1_HUMAN
1587	SSSMSSLTGAYTSGIPSSSR202000000000000000	sp A8CG34 P121C_HUMAN

1588	LPPFPGLEPESEGAAGGSEPEAGDSDTEGEDIFTGAAVVK2. 00000002E+24	sp Q13596	SNX1_HUMAN
1589	SSVAAPEKSSSNDVDEETAESDTSPVLEK2. 22E+22	sp Q99590	SCAFB_HUMAN
1590	KPVNYSQFDHSDSDDDFVSATVPLNK2020000000000000	sp Q96B01	R51A1_HUMAN
1591	KRDSEEEFGSER2000002000	sp Q12873	CHD3_HUMAN
1592	RPMEEDGEKSPSK200	sp Q12906	ILF3_HUMAN
1593	ASSHSSQTQGGGSVTK2020000000000000	sp P02545	LMNA_HUMAN
1594	EMLMEDVGSEEEQEEDEAPFQEK2000000000000000	sp Q9H1E3	NUCKS_HUMAN
1595	HTDDEMTGYVATR300000	sp Q16539	MK14_HUMAN
1596	EHYPVSSPSSPPAQPGGVSR2002000000000000	sp 075179	ANR17_HUMAN
1597	TLSADASVNTLPVVVSR2000000000000000	sp Q92615	LAR4B_HUMAN
1598	NLATSADTPPSTVPGTGK2002000000000000	sp Q96Q15	SMG1_HUMAN
1599	SPEASASAPHTPR2000000000000000	sp Q8TF72	SHRM3_HUMAN
1600	VSPSDTTPLVSR2000000000000000	sp Q9HCK8	CHD8_HUMAN
1601	HSSNPPLSHVGVMDSR2000000000000000	sp Q6PKG0	LARP1_HUMAN
1602	SPGDFTSAAQLASTPFHK2000000000000000	sp 043683	BUB1_HUMAN
1603	ESESEDETTPAAPQLIK2020200000000000	sp 060832	DKC1_HUMAN
1604	VTFGEDLSPEVFDLPLANTPLR2000000000000000	sp Q69YH5	CDCA2_HUMAN
1605	TSPADHGGVSGSESGSAVDSVAGEHSVSGR2000000000000000	sp Q5T4S7	UBR4_HUMAN
1606	EGSEFSFDGEVAEK2020000000	sp Q9BVS4	RIOK2_HUMAN
1607	QITQEEDDSDEEVAPENFFSLPEK2000000000000000	sp Q92733	PRCC_HUMAN
1608	HGLQLGAQSPGR20000	sp Q8N1G0	ZN687_HUMAN
1609	GTGSGGQLQDLDCSSDDEGAAQNSTKPSATK2. 2E+19	sp P08240	SRPR_HUMAN
1610	SSPVNNHSPGQTPTLGGPEAR2000000000000000	sp Q5SW79	CE170_HUMAN
1611	GLLSNHTGSPR2000	sp Q96L14	C170L_HUMAN
1612	SASQSSLDKLDQELK220000000000	sp Q13111	CAF1A_HUMAN
1613	SASQSSLDKLDQELK220000000000	sp 060271	JIP4_HUMAN
1614	DTSENADGQSDENKDDYTIPDEYR2E+22	sp P43243	MATR3_HUMAN
1615	SPATSPISSNSHR20002000000000	sp Q9NR09	BIRC6_HUMAN
1616	DNLLDTYSADQGDSSSEGGTLAR2000000000	sp Q6ZRP7	QSOX2_HUMAN
1617	SPVGKSPSTGTYGSSQK2000000000000000	sp Q9Y2W1	TR150_HUMAN
1618	GYSPYSVSGSGSTAGSR300000020000000000	sp Q15149	PLEC_HUMAN
1619	LASPSGSTSSGLEVVAPEGTSAPGGGPGTLDDSATICR2. 2E+30	sp Q13263	TIF1B_HUMAN
1620	QSPASPPPLGGAPVR2000000000000000	sp Q14160	SCRIB_HUMAN
1621	DLFSLDSEDSPASPPLR200000	sp Q9BTK6	PAGR1_HUMAN
1622	GNDSGEGESDDPEK2000000000000000	sp 075351	VPS4B_HUMAN
1623	EDVTPPLSPDDLK200020000000	sp Q9Y4B5	MTCL1_HUMAN
1624	QVQSLTCEVDALK200000000000	sp P08670	VIME_HUMAN
1625	YLVDGTKPNAGSEEISSEDELVEEK2000220000000000	sp Q9NY61	AATF_HUMAN
1626	DIEISTEEKDTGDLK2200000000000000	sp Q4G0J3	LARP7_HUMAN
1627	GDAASSPAPAASVGSQGGAR2000000000000000	sp Q8WVB6	CTF18_HUMAN
1628	DELHIVEAEAMNYEGSPIK1000020000	sp P06748	NPM_HUMAN
1629	LPLVPESPR2000	sp Q86WB0	NIPA_HUMAN
1630	SRSDIDVNAASAK2000000000000000	sp Q7Z460	CLAP1_HUMAN
1631	AESISEEADSEPGR200002000000	sp Q9NSI6	BRWD1_HUMAN
1632	LEPVSPSPPHTDPELELVPPR20020000000000000000	sp Q9Y618	NCOR2_HUMAN
1633	STSAPQMSPGSSDNQSSSPQPAQQK1. 2E+19	sp Q14157	UBP2L_HUMAN
1634	SSPKEELHPAAPSQ LAPSFSSSSSSSSGPR2E+29	sp 043166	SIIL1_HUMAN
1635	SGSAAQAEGLCK200000000000	sp Q92685	ALG3_HUMAN
1636	SESELIDELSEDFDR2000000000000000	sp P20810	ICAL_HUMAN
1637	YLVDGTKPNAGSEEISSEDELVEEK220000000000	sp Q9NY61	AATF_HUMAN

		sp	P05534	1A24_HUMAN
		sp	P30447	1A23_HUMAN
1637	KGGSYSQAASSDSAQGSVSLTACKV200020000000000	sp	P01891	1A68_HUMAN
		sp	P01892	1A02_HUMAN
		sp	Q09160	1A80_HUMAN
		sp	P10316	1A69_HUMAN
1638	KEESEESDDDMGFGLFD200000000000	sp	P05387	RLA2_HUMAN
		sp	P05386	RLA1_HUMAN
1639	SPHSPPIATPPSQSSSCSDAPMLSTVHSAK2.0222E+19	sp	Q9NYF8	BCLF1_HUMAN
1640	SSSVGSSSSYPISPAISR200	sp	Q15149	PLEC_HUMAN
1641	HGSGADSDYENTQSGDPLLGLEGK2.0002E+22	sp	Q9Y2X7	GIT1_HUMAN
1642	ASQGLLSSIENSESDSSEAK2020000000	sp	Q5UIP0	RIF1_HUMAN
1643	KQSGSPTLDTAPNGR20200000000000	sp	Q9H2J7	S6A15_HUMAN
1644	DYDEEEQGYDSEKEK200000	sp	Q05519	SRS11_HUMAN
1645	SPVSGGAPQAAAPAPAAHVAGNPGGDAAPAATGTAAAASLATAAGSEDAEK200000000000000000	sp	P16989	YBOX3_HUMAN
1646	GIPLATGDTSPPELLPGAPLPPPK200000000000000000	sp	Q075821	EIF3G_HUMAN
1647	SLSPGKENVSALDMEK200000000000000000	sp	Q9H307	PININ_HUMAN
1648	SIHEVDISNLEAALR200000000000000000	sp	Q9Y2H5	PKHA6_HUMAN
1649	SELGNQSPSTSSR20000000	sp	Q5JTV8	TOIP1_HUMAN
1650	EAEGEDDRDSANGEDDS200000000	sp	P07910	HNRPC_HUMAN
1651	LLDESESAASSR2002000	sp	Q8N108	MIER1_HUMAN
1652	ASPAGTAGGPGAGAAAGGTGPLAAR2E+20	sp	Q12962	TAF10_HUMAN
1653	KEESEESDDDMGFGLFD2000000000000000	sp	P05386	RLA1_HUMAN
		sp	P05387	RLA2_HUMAN
1654	EVSDSEAGGPGQGER2000000000000000	sp	Q9BW71	HIRP3_HUMAN
1655	TASGSSVTSLDGTR20220000000000	sp	Q92597	NDRG1_HUMAN
1656	DKSDSDEGLLFSR20200000000000	sp	Q9H4G0	E41L1_HUMAN
1657	NTSSDNSDEVMPAQSPR2002000010002000	sp	Q9HCK8	CHD8_HUMAN
1658	QASPTVVVER200000000	sp	Q8TB72	PUM2_HUMAN
1659	SRSPTPPSSAGLGSNSAPPIDSR2.0022E+20	sp	Q8IWX8	CHERP_HUMAN
1660	DSGNWDTSGSELSEGELEK20020020000000	sp	Q075400	PR40A_HUMAN
1661	FTSQDASTPASSELNK2000020000000000	sp	Q9HAW4	CLSPN_HUMAN
1662	TTPNSGDVQVTEDAVR2000000000000000	sp	P35269	T2FA_HUMAN
1663	ENSPSSQSAGLSSINK20000000000000	sp	Q9H2Y7	ZN106_HUMAN
1664	KLPPPPQAPPEEENESEPEEPPSGVEGAQSR200000000000000000	sp	Q060341	KDM1A_HUMAN
1665	YMLTHQELASDGEIETK200000000	sp	Q02241	KIF23_HUMAN
1666	ESEDKPEIEDVGSDEEEKK200000000	sp	P07900	HS90A_HUMAN
1667	LFEDDSNEKLFDEEEDSSEK2000000000002000	sp	Q043719	HTSF1_HUMAN
1668	ELEKPIQSKPQSPVIQAAVSPK200000000000002000	sp	Q9Y6D5	BIG2_HUMAN
1669	STLTSSYNCPSPYADGR20000000	sp	Q9HCU4	CEL2R_HUMAN
1670	ENYDSEEEEDDDVASSR3002000000000000	sp	Q9Y2U8	MAN1_HUMAN
1671	LLPEGEETLESDEK200000	sp	Q8N6N3	CA052_HUMAN
1672	ASLGSLEGEAEAEASSPK200200000000000000	sp	Q09666	AHNK_HUMAN
1673	CAPSAGSPAAVGR200000000	sp	Q7L2J0	MEPCE_HUMAN
		sp	P51812	KS6A3_HUMAN
1674	AYSFCGTVEYMAPEVVNR2000000000000000	sp	Q9UK32	KS6A6_HUMAN
		sp	Q15418	KS6A1_HUMAN
1675	KVPVQEIDDDFFPSSGEEAEASVGEAGGGGR2.2E+19	sp	Q03468	ERCC6_HUMAN
1676	LHQLSGSDQLESTAHSR20200000000000	sp	Q060271	JIP4_HUMAN
		sp	P07355	ANXA2_HUMAN
1677	LSLEGDHSTPPSAYGSVK200000000000000000	sp	Q6NMY6	AXA2L_HUMAN
		sp	P52948	NUP98_HUMAN
1678	DSENLASPSEYPENGER200000000000			
1679	LPSGGAASPTGSAVDIR2000000000000000	sp	Q09666	AHNK_HUMAN

1680	SEPVKEESSELEQPFAQDTSSVGPDR2000000	sp Q15424 SAFB1_HUMAN
		sp Q14151 SAFB2_HUMAN
1681	QICLVMLETLSQSPQGR200000	sp Q15365 PCBP1_HUMAN
1682	YSGAYGASVSDEELK200000000000000	sp Q9NX63 MIC19_HUMAN
1683	GPTSSPCEEEGDEGEEDR220000000000000	sp Q86VM9 ZCH18_HUMAN
1684	GPTSSPCEEEGDEGEEDR200000000000000	sp Q86VM9 ZCH18_HUMAN
1685	AGSSTPGDAPPAVAEVQGR200000000000000	sp Q7Z6Z7 HUWE1_HUMAN
1686	GARPPAAGPGGDEDEDEEDTAPESALDTSLDK20002000220000	sp Q27J81 INF2_HUMAN
1687	VVPGQFDDADSSDSEN2000000	sp Q9BRS2 RIOK1_HUMAN
1688	LKNDSLFLGLGLEEAGPKESSEEGK2200000	sp Q3YEC7 RABL6_HUMAN
1689	TLDAEVVEKPAK200000000000000	sp Q9H3P2 NELFA_HUMAN
1690	KMSQPGSPSPK2000200000	sp Q14839 CHD4_HUMAN
1691	GRESDEDEDASETDLAK2000200020000000	sp Q9H7L9 SDS3_HUMAN
1692	METVSNASSSSNPSSPGR1000000000000020000	sp Q9NTI5 PDS5B_HUMAN
1693	METEADAPSPAPSLGER2000200000	sp Q12873 CHD3_HUMAN
		sp Q71U36 TBA1A_HUMAN
		sp P68363 TBA1B_HUMAN
1694	TIGGGDDSFNTFFSETGAGK200000000000000	sp Q6PEY2 TBA3E_HUMAN
		sp Q13748 TBA3C_HUMAN
		sp Q9BQE3 TBA1C_HUMAN
1695	HGSGADSDYENTQSGDPLLGLGK2000000000000000000	sp Q9Y2X7 GIT1_HUMAN
1696	DNSPEPNDPEEPQEVSSTPSDK2E+20	sp Q9HB58 SP110_HUMAN
1697	SDSSGGYNLSDI IQSPSSTGLLK2E+21	sp Q9P2D0 IBTK_HUMAN
1698	AALGLQSDDEDAAVDIDEQIESMFNSK2E+21	sp Q7L014 DDX46_HUMAN
1699	SLESSVIMDR2000000000	sp Q9H2Y7 ZN106_HUMAN
1700	DTQSPSTCSEGLLGWSQK2000000000000000000	sp Q9C0C2 TB182_HUMAN
1701	SNTPESIAETPPAR2020000000000000	sp Q9Y4H2 IRS2_HUMAN
1702	SQSLPTLLSPVR200000000000000	sp P28290 SSFA2_HUMAN
1703	DSGNWDTSGSELSEGELEK220200200000000	sp O75400 PR40A_HUMAN
1704	YSPSQNSPIHHIPSR30000020000000000	sp Q9NYF8 BCLF1_HUMAN
1705	YMAENPTAGVVQEEEEEDNLEYDSGNPIAPTK1E+31	sp Q86XP3 DDX42_HUMAN
1706	GQSTDSLPQTICR2000000000	sp Q96NE9 FRMD6_HUMAN
1707	DRPGSPESPLLDAPFSR2002000000000000	sp Q8TF72 SHRM3_HUMAN
1708	HGLAHDEMKSPPR2000	sp Q9Y2W1 TR150_HUMAN
1709	VPPAPVPCPPSPGSAVPSSPK2000000002000	sp O95817 BAG3_HUMAN
1710	EMEHNTVCAAGTSPVGEIGEEK20000000000	sp P18583 SON_HUMAN
1711	NHSPSPPTPTGAAPSLASPK2. 020002E+19	sp Q5SYE7 NHSL1_HUMAN
1712	AESSSSISSASLSAANTPTVGVSR2. 202E+21	sp Q0JRZ9 FCHO2_HUMAN
1713	HASSPESPKPAPAPGSHR2200000000000000000	sp Q9UQ35 SRRM2_HUMAN
1714	AGSSTPGDAPPAVAEVQGR2000000000000000000	sp Q7Z6Z7 HUWE1_HUMAN
1715	AASLTEDR2000000	sp Q04637 IF4G1_HUMAN
1716	YTPTSPSYSPSSPEYTPTSPK32000000000000000	sp P24928 RPB1_HUMAN
1717	SATVKPGAVGAGEFVSPCESGDNTGEPALAEQR2. 0002E+19	sp Q12888 TP53B_HUMAN
1718	LHNENAEMSDSSSSGTETDLHGSLR22220000000000000	sp Q9H832 UBE2Z_HUMAN
1719	QGLAETASPVAVSLR2000000000	sp Q15154 PCM1_HUMAN
1720	NASASFQLEDK20200000000	sp Q99543 DNJC2_HUMAN
1721	RKSEDGTPAEDGTPAATGGSQPPSMGR2E+25	sp Q9BQG0 MBB1A_HUMAN
1722	EGMNPYSYDEYADSDQHDAYLER200000000000000	sp Q08945 SSRP1_HUMAN
1723	SGSSSPDSEITELKFPSINH2. 22E+19	sp P17812 PYRG1_HUMAN
1724	ALSGTSPDDVQPGPSVGPSPK2000000000000000000	sp Q9Y314 NOSIP_HUMAN
1725	SVVSDLEADDVK200000000000000	sp P11388 TOP2A_HUMAN
1726	LDDAHSLSGSGAGEGYEPISSDELDEILAGDAEK2. 0020000000002E+28	sp Q5T200 ZC3HD_HUMAN
1727	KPTAGQTSATAVSDDDIQTISSGSEGDDFEDKK2. 0000202202E+20	sp Q15047 SETB1_HUMAN

1728	DLGHPVEEEDELESGDQEDEDDESEDPGKDL DHI2E+21	sp 060763 USO1_HUMAN
1729	SSDEENGPSSPDLDR22000000002000000	sp Q96B36 AKTS1_HUMAN
1730	EVSSRPSTPGLSVVSGISATSEDI PNK2. 2E+25	sp Q14C86 GAPD1_HUMAN
1731	QGSITSPQANEQSVTPQR20000	sp Q9UQ35 SRRM2_HUMAN
1732	DSALQDTDDSDDDPVLIPGAR2002000000000000	sp Q58WW2 DCAF6_HUMAN
1733	HSTPSNSSNPSGPPSPNSPHR20020000	sp Q9Y5S2 MRCKB_HUMAN
1734	GIPLATGDTSPPELLPGAPLPPPK2. 0002E+20	sp 075821 EIF3G_HUMAN
1735	GTSDSSSGNVSEGESPPDSQEDSFQGR2E+23	sp Q8WUB8 PHF10_HUMAN
1736	DKEVSDDEAEK200000000	sp P07900 HS90A_HUMAN
1737	NDSGEENVPLDLTR2000000000000	sp Q6R327 RICTR_HUMAN
1738	LSSTDDGYIDLQFK22000000000000	sp Q96A57 TM230_HUMAN
1739	SIEVENDFLPVEK20000000000000	sp Q96B97 SH3K1_HUMAN
1740	DDDTASVTLNSSATGR22000000000	sp P21675 TAF1_HUMAN
1741	VSLEPHQGGPTPEK200000	sp Q14980 NUMA1_HUMAN
1742	GLGSSPDLELLLR200000000000	sp Q9NQQ7 S35C2_HUMAN
1743	LSPTSMQDGLDLPSETDLR2100000000000000	sp Q9UK58 CCNL1_HUMAN
1744	RGSSGSVDETLFALPAASEPVIR2. 02E+20	sp Q9NQC3 RTN4_HUMAN
1745	HIISATSLSTSPTELGSR200000000	sp Q5JSZ5 PRC2B_HUMAN
1746	LFDEEEDSSEKLFDDSDER200000020000	sp 043719 HTSF1_HUMAN
1747	SQSDLDDQHDYDSVASDETDQEPLR30000200000000000	sp Q9Y2X7 GIT1_HUMAN
1748	TPKDSPGIPPSANAHQLFR2. 0002E+19	sp P51812 KS6A3_HUMAN
1749	SLESSVIMDR2002000000	sp Q9H2Y7 ZN106_HUMAN
1750	VPDEEENEESDNEKETEK2000000000	sp P11388 TOP2A_HUMAN
1751	DITDPLSLNTCTDEGHVVLASPLK20000	sp Q7L2J0 MEPCE_HUMAN
1752	TSSGTSLSAMHSSGSSGK2200000000000000	sp Q12888 TP53B_HUMAN
1753	VIGQDHDFSESSEEEAPAEASSGALR2022000000000000000	sp P23497 SP100_HUMAN
1754	SPSLSLAPLSR2000000002000	sp Q969H4 CNKR1_HUMAN
1755	KGSDDAPYSPTAR200000000000	sp 075151 PHF2_HUMAN
1756	ESYTEEIVSEAESHVSGISR2000200200000	sp Q5QJE6 TDIF2_HUMAN
1757	EEETSIDVAGKPNEVTK20000000000000	sp P53985 MOT1_HUMAN
1758	EYDSGDTDEIIAMK200200000100	sp Q76FK4 NOL8_HUMAN
1759	SEDEDSLEEAGSPAPGPCPR2. 00002000002E+20	sp Q8TBB5 KLDC4_HUMAN
1760	EAANEAGDSSQDEAEDDVK220000000000	sp Q9NVU0 RPC5_HUMAN
1761	TSDANETEDHLESLICK20000000000000000	sp Q09161 NCBP1_HUMAN
1762	MDRTPPPPTLSPAAITVGR20000002000000000	sp Q8NDX5 PHC3_HUMAN
1763	DAPGEDEEEDGVSEAASLEPK20002000000	sp Q9H0E9 BRD8_HUMAN
1764	LDSSEMDSHSENYDTMSSPLPGK2. 010020000001E+20	sp Q9NTI5 PDS5B_HUMAN
1765	KPVTVSPTTPTSPTEGEAS200000000	sp Q9Y6G9 DC1L1_HUMAN
1766	EATNTTSEPSAPSQDLLDLSPSPR2000	sp 075674 TM1L1_HUMAN
1767	NENTEGSPQEDGVELEGLK20000000000000000	sp P11388 TOP2A_HUMAN
1768	GSLAEAVGSPPPAATPTPTPTR20000000000000000	sp Q9Y6I3 EPN1_HUMAN
1769	GADNDGSGSESGYTTPK202000000000	sp Q7Z417 NUFP2_HUMAN
1770	ASLGSLEGEAEAEASSPK2000	sp Q09666 AHNK_HUMAN
1771	SPGMLEPLGSSR20000000000000	sp Q9UQ35 SRRM2_HUMAN
1772	STSQGSINSPVYSR20000000000000	sp 014639 ABLM1_HUMAN
1773	AGSISTLDSLDFAR20000000000000	sp Q9P260 K1468_HUMAN
1774	VTDALPEPEPPGAMAASEDEEEEEEALEAMQSR2000000000000000000	sp Q9Y3E7 CHMP3_HUMAN
1775	SLSPQEDALTGSR20200000000000	sp Q96EN8 MOCOS_HUMAN
1776	KLSSSSEPYEDEFNDDQSIK2000300000000000000	sp Q9H4L7 SMRCD_HUMAN
1777	ADEPSSEESDLEIDK20020000000	sp P50502 F10A1_HUMAN
1778	SPHSPSPATPPSQSSSCSDAPMLSTVHSAK2. 00202E+31	sp Q8IZP2 ST134_HUMAN
1779	ESLKEEDESDDNM2000010	sp Q9NYF8 BCLF1_HUMAN
1780	SIPQLDLGWPDTIAYR20000000000000000	sp P25788 PSA3_HUMAN
		sp A6ND36 FA83G_HUMAN

1781	LTPVSPESSTEEK2000000000	sp Q13501 SQSTM_HUMAN
1782	VSALEEDMDDVESSEEEEEDEKLER1.000022E+19	sp Q8NAV1 PR38A_HUMAN
1783	KPGPPLSPEIR200000	sp Q96JM3 CHAP1_HUMAN
1784	AGLSPERPPK20000000	sp Q9H6F5 CCD86_HUMAN
1785	QICVVMLETLSQSPPK20000	sp Q15366 PCBP2_HUMAN
1786	LTVSDGESGEEKK20002000000	sp P46100 ATRX_HUMAN
1787	VDGETASDESER2000000	sp O95674 CDS2_HUMAN
1788	YGPADVEDTTGSGATDSKDDDDIDLFGSDDEEESEEAK200000000000	sp P24534 EF1B_HUMAN
1789	TASISSPSEGTPTVGSYGCTPQSLPK2.022E+25	sp Q6PKG0 LARP1_HUMAN
1790	SSSPVTELASR2000000000	sp Q9UQ35 SRRM2_HUMAN
1791	SIGSAVDQGNESIVAK200200000000000000	sp Q9H0H5 RGAP1_HUMAN
1792	GLRDSHSSEDEASSQTDLSQTISK20000000000000000000	sp Q5JTV8 TOIP1_HUMAN
1793	SVCGHLENTSVGNPNPSSAENSFR200020000000000000	sp Q96HH9 GRAM3_HUMAN
1794	GSSAAASPGSPPPGR2000000	sp Q86UX6 ST32C_HUMAN
1795	RKPSPEPEGEVGPVK2000000000000000	sp Q7Z5L9 I2BP2_HUMAN
1796	THSTSSSLGSGESPFGR2000000000000000	sp Q9UGV2 NDRG3_HUMAN
1797	ATSSSSGSLSATGR20000000000	sp Q03252 LMNB2_HUMAN
1798	SSVSDAPVHITASGEPVPISEESELDQK2E+28	sp O14646 CHD1_HUMAN
1799	SSTPGSPTHVSSGSNASR2000000000000000	sp Q14CZ0 CP072_HUMAN
1800	TISPSTSNIPLMR20020000000000	sp O94806 KPCD3_HUMAN
1801	SPAGSPDEEGGAEAPAAGIR20000000000000000000	sp Q15139 KPCD1_HUMAN
1802	AVTIANSKPKSEK2000000000	sp O15061 SYNEM_HUMAN
1803	NLEHLSSFSDDEDDPGYSQDAYK20002000000000000000	sp Q8NEF9 SRFB1_HUMAN
1804	KTSSDDESEEDDLLQR20000200000000000000	sp Q2KHR3 QSER1_HUMAN
1805	AGSPLCNSNLQDSEEDIEPSYSLFR20000200000000000000	sp Q9Y5J1 UTP18_HUMAN
1806	TLSDESIYNSQR20200000000000	sp Q9NRY4 RHG35_HUMAN
1807	RKSLDSESDDSK202000000000	sp O43166 SI1L1_HUMAN
1808	SPCETISSPSTLESK200000000000000000	sp Q13185 CBX3_HUMAN
1809	GPPDFSSDEEREPTPVLGSGAAAAGR2000000000000000	sp Q9COD5 TANC1_HUMAN
1810	KDSEEEVSLGSDIEEGNHQVEDGCR2E+25	sp P42166 LAP2A_HUMAN
1811	TDEVPAGGRSEAEDDEDYVPYVPLR2.02E+20	sp P42167 LAP2B_HUMAN
1812	VTETEDSDSDSDDEDDVHVTIGDIK2.020002E+26	sp Q9Y487 VPP2_HUMAN
1813	GGDDHDDTSDSDGLTLK22020000000000	sp Q9UJV9 DDX41_HUMAN
1814	QPLSEASNQQPLSGEEK2000000	sp Q6UN15 FIP1_HUMAN
1815	CEERSPSFGEDYYGPSR2020000000000000	sp Q9BTC0 DIDO1_HUMAN
1816	SPLSDYMNLDSSPK200000000000000000	sp Q9NQW6 ANLN_HUMAN
1817	KGSITEYTAAEEK20000000000000	sp P49761 CLK3_HUMAN
1818	VGSPLDYSLVLPSTNGQSPGK2E+20	sp Q9Y4H2 IRS2_HUMAN
1819	TAVQYIESSDSEIETSELPQK2020000000000000	sp Q12982 BNIP2_HUMAN
1820	HIVSNDSSDSDDESHEPK2000000000	sp Q96T23 RSF1_HUMAN
1821	KGGYSQAASSDSAQGSVSLTACK2000000000	sp Q14527 HLTF_HUMAN
1822	QICLVMLETLSQSPQGR1000000200000	sp P47712 PA24A_HUMAN
1823	GPPDFSSDEEREPTPVLGSGAAAAGR2E+21	sp P05534 1A24_HUMAN
1824	DSTSQHDDDNISTTSGFSSR20000000000000000000	sp Q09160 1A80_HUMAN
1825	RDSSESQLASTESDKPTTGR20000000000000000000	sp P30447 1A23_HUMAN
1826	DQPPFGSDSDSVEADK2002000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
		sp Q15365 PCBP1_HUMAN
		sp P42166 LAP2A_HUMAN
		sp P42167 LAP2B_HUMAN
		sp P49790 NU153_HUMAN
		sp Q96B23 CRO25_HUMAN
		sp Q14676 MDC1_HUMAN

1827	STSAPQMSPGSSDNQSSSPQPAQQK2E+23	sp Q14157 UBP2L_HUMAN
1828	QGLAETASPVAVSLR200002000	sp Q15154 PCM1_HUMAN
1829	AGEPDGESLDEQPSSSSSK200000000000	sp Q9H3Q1 BORG4_HUMAN
1830	INPDNYGMDLNSDDSTDDEAHPR2002200000000	sp Q9NQS7 INCE_HUMAN
1831	TDGFAEAIHSPQVAGVPR2000000000	sp P12270 TPR_HUMAN
1832	TCNSPQNSTDSVSDIVPDSPPFGALGSDTR2.02E+22	sp Q14694 UBP10_HUMAN
1833	TPTAPAVNLAGAR20000000000000	sp Q9UQ35 SRRM2_HUMAN
1834	NSSDSAIDNPKPNK2020000000000	sp P46100 ATRX_HUMAN
1835	LDQPVSAPPSPR2000	sp Q16204 CCDC6_HUMAN
1836	SHSLSQHTATSSK200000000000	sp P49790 NU153_HUMAN
1837	GVMAVTAVTATAASDR10000000002000	sp Q7Z4V5 HDGR2_HUMAN
1838	SSSLGSTPHEELER22000000000000	sp Q9NRA8 4ET_HUMAN
1839	ASDPQSPPQVSR20000000	sp Q7L7X3 TAOK1_HUMAN
1840	DSGNWDTSGSELSEGELEK2000020000000	sp 075400 PR40A_HUMAN
1841	HASSSPESPKPAPAPGSHR2020000000000000	sp Q9UQ35 SRRM2_HUMAN
1842	QSPGPALAR200000000	sp P85037 FOKK1_HUMAN
1843	GDPDSEADSIDSDQEDPLK2000200200000000	sp Q9H9L4 KANL2_HUMAN
1844	DSDQVAQSDGEESPAEEQLLGEHIK2.00002E+19	sp 094979 SC31A_HUMAN
1845	HTGMASIDSSAPETTSOSSPTLSR20000000	sp A7E2V4 ZSWM8_HUMAN
1846	SYLEGSSDNQLKDESETPVDDR2.30002E+22	sp Q9UHI6 DDX20_HUMAN
1847	KSVSSPTSSNTPPTK2000000000000000	sp Q5M775 CYTSB_HUMAN
1848	SPDPFGAVAAQK2000000000000	sp Q9Y3Q8 T22D4_HUMAN
1849	AEEPPSQLDQDTQVQDMDEGSDEEEGQK2000000000	sp Q15459 SF3A1_HUMAN
1850	LPSTSGSEGVPR200000000000	sp P29317 EPA2_HUMAN
1851	SGSSQELDVKPSASPQER2000000000200000	sp Q9UQ35 SRRM2_HUMAN
1852	SPALKSPLQSVVVR2000020000000000	sp Q9Y2W1 TR150_HUMAN
1853	SSTPLPTISSAENTR2000000000000000	sp P42167 LAP2B_HUMAN
1854	KPEKPLFSSASPQDSSPR200000000	sp P42166 LAP2A_HUMAN
1855	SSLSGDEEDELFK20020000000000	sp 000712 NFIB_HUMAN
1856	ASSVSTKESPAR20000	sp Q5T1M5 FKB15_HUMAN
1857	LASPELER2000000	sp Q13428 TCOF_HUMAN
		sp P05412 JUN_HUMAN
		sp P17535 JUND_HUMAN
		sp 015320 CTGE5_HUMAN
		sp P0CG41 CTGE8_HUMAN
		sp Q8IX94 CTGE4_HUMAN
1858	SNSELEDEILCLEK2020000000000000	sp A4FU28 CTGE9_HUMAN
		sp A4D2H0 CTGEF_HUMAN
		sp Q86UF2 CTGE6_HUMAN
1859	DLHQGIEAASDEEDLR20000000	sp Q9UKS6 PACN3_HUMAN
1860	EGITGPPADSSKPIGPDDAIDALSSDFTCGSPTAAGK20000000	sp P20810 ICAL_HUMAN
1861	TTSPSSDTLLDR200200000000	sp Q96QT6 PHF12_HUMAN
1862	SYLEGSSDNQLKDESETPVDDR2.00002E+22	sp Q9UHI6 DDX20_HUMAN
1863	LGGLRPESPELTSVSR20000000000	sp Q9H6F5 CCD86_HUMAN
1864	IDDRSDEEGASDR2000000000	sp Q8TDD1 DDX54_HUMAN
1865	ENPPVEDSDEDDKR20000000	sp Q8TCJ2 STT3B_HUMAN
1866	WDGSEEDENSK2000000000	sp 060841 IF2P_HUMAN
1867	EHNSMWSLSSPDAAEVEPDFSSIERS2000000000000000	sp Q27J81 INF2_HUMAN
1868	SSSMSSLTGAYTSGIPSSSR2200000000000000	sp A8CG34 P121C_HUMAN
1869	YNLDASEEEDSNK200000000	sp 095218 ZRAB2_HUMAN
1870	ESEDKPEIEDVGSDEEEKKGDK200000000000	sp P07900 HS90A_HUMAN
1871	SSPELLPSGVTDENEVTTAVTEK2E+22	sp P42566 EPS15_HUMAN
1872	SPGSTPTTPTSSQAPQK220000000000	sp P35658 NU214_HUMAN
1873	GFEEHKSDDSSDDEQEKKPEAPK22000000000000	sp Q13435 SF3B2_HUMAN



1874	TASISSPSEGTPTVGSYGCTPQSLPK2E+23	sp Q6PKG0 LARP1_HUMAN
1875	TEHSHSPGDSSAPIQNTDCCHSSER2E+20	sp Q13129 RLF_HUMAN
1876	GLSVDSAQEVK2000000000	sp Q9Y4A5 TRRAP_HUMAN
1877	EQGTESRSSTPLPTISSSAENTR2220000000000000	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
1878	VVEAVNSDSDSEFGIPK202020000000	sp Q02880 TOP2B_HUMAN
1879	LSGGLGAGSCR2000	sp Q04695 K1C17_HUMAN
1880	ALIQTCLNSPDSPPR20000	sp P11831 SRF_HUMAN
1881	RSSQSPPTAVPASDSPPTK200200000000000000	sp Q3KQU3 MA7D1_HUMAN
1882	NGSEADIDEGLYSR20000000000000	sp P22314 UBA1_HUMAN
1883	EESEEAEPVQR200200000000	sp Q9BW71 HIRP3_HUMAN
1884	SKSDSYTLDPDTLR20000000000000	sp 075592 MYCB2_HUMAN
1885	LSSGFDDIDLPSAVK2000000000000000	sp Q9Y446 PKP3_HUMAN
1886	EAYNDSSDEESSHHGPGVQCAHQ2. 2E+19	sp 060884 DNJA2_HUMAN
1887	TQEDEEEISTSPGVSEFVSDAFDACNLNQEDLR2E+33	sp Q96A49 SYAP1_HUMAN
1888	SSFASSASDASKPSSPR2000	sp Q96IF1 AJUBA_HUMAN
1889	LQPQEISPPPTANLDR200000000000	sp Q05397 FAK1_HUMAN
1890	SAPTSPTATNISSK22000000000000	sp Q9H792 PEAK1_HUMAN
1891	SEDPPGQEAEGSEEGSSASGLAK20000000000000	sp 075153 CLU_HUMAN
1892	GLRDSHSSEDEASSQTDLSTISK2. 02E+21	sp Q5JTV8 TOIP1_HUMAN
1893	KLEGNPQGSNGVK2000000	sp P61006 RAB8A_HUMAN
1894	SYLEGSSDNQLKDESTPVDDR2. 000022E+22	sp Q9UHI6 DDX20_HUMAN
1895	KNSTDLDLAPEDPTSPK2000020000002000	sp Q96RK0 CIC_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
1896	EDALDDSVSSSSVHASPLASSPVR20000	sp 014715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
		sp A6NKT7 RGPD3_HUMAN
1897	SPTAALNESLVECPK20000020000000	sp Q53EZ4 CEP55_HUMAN
1898	LQSPIKEENTTAVEEIGR200000000000000000	sp Q9NS73 MBIP1_HUMAN
		sp Q00526 CDK3_HUMAN
1899	IGEGTYGVVYK23000000	sp P06493 CDK1_HUMAN
		sp P24941 CDK2_HUMAN
1900	REEGPPPPSPDGASSDAEPEPPSGR200000000000000000	sp Q9NTJ3 SMC4_HUMAN
1901	DDDTASVTLNSSATGR20200000000000	sp P21675 TAF1_HUMAN
1902	GPPQSPVFEGVYNNR20000000000000	sp Q8WWM7 ATX2L_HUMAN
1903	SGSAAQAEGLCK20000000000000	sp Q92685 ALG3_HUMAN
1904	GPPDFSSDEEREPTPVLGSGAAAAGR2E+20	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
1905	SASSDTSEELNSQDSPPK220220000000000000	sp 014745 NHRF1_HUMAN
1906	NMGGPYGGNYGPGSGGGYGR1E+24	sp P22626 ROA2_HUMAN
1907	ATSSSGSLSATGR200000000000	sp Q03252 LMNB2_HUMAN
1908	KPPAPSPVQSQSPSTNWSPAVPVK2. 00000002002E+19	sp Q8IYB3 SRRM1_HUMAN
1909	HQEVQDQDPVFQGS DSSGYQSDHKK22030200000	sp Q3B726 RPA43_HUMAN
1910	LFEESDDKEDEDADGKEVEDADEK2E+20	sp 043719 HTSF1_HUMAN
1911	SNSLPHSAVSNAGSK200000000000000000	sp Q8TBZ3 WDR20_HUMAN
1912	ECINAAPDPSK20000	sp Q7Z569 BRAP_HUMAN
1913	KVVDYSQFQESDDADEDYGR200000000000	sp Q9H1E3 NUCKS_HUMAN
1914	RDSSESQLASTESDKPTTGR200000000000000000	sp Q96B23 CRO25_HUMAN
1915	SSSVSPSSWK2000000	sp Q96JM3 CHAP1_HUMAN
1916	AEAPSPDVAPAGK220000000000	sp Q9P1Y6 PHRF1_HUMAN
1917	HGSFNTVQASER200000000000	sp Q96AV8 E2F7_HUMAN
1918	ATTPADGEEPAPAEALAAAR2E+19	sp Q96S44 PRPK_HUMAN
1919	RSSLPLDHGSPAQENPESEK200000000000000000	sp Q5VZ89 DEN4C_HUMAN

1920	LPGSGAVQAASPER20000	sp Q86U90 YRDC_HUMAN
1921	SLSSGESGPGSPTHSHLSR2000200000000000	sp Q6P0Q8 MAST2_HUMAN
1922	FLETDSEEEQEVEVNEK20200000000000	sp Q76FK4 NOL8_HUMAN
1923	SLESSVIMDR200000000000	sp Q9H2Y7 ZN106_HUMAN
1924	TSSGTSLSAMHSSGSSGK20000001000000000	sp Q12888 TP53B_HUMAN
1925	SDISDQEEDEESEGPCVSNLSK2E+23	sp Q15154 PCM1_HUMAN
1926	GSDALSETSSVSHIEDLEK22200000000000	sp Q6P996 PDXD1_HUMAN
1927	SPHQLLSPSSFSPSATPSQK2E+20	sp Q14181 DPOA2_HUMAN
1928	QPDISCILGTGGKSPR2000	sp P57740 NU107_HUMAN
1929	AGDLLEDSPKRPK2000000	sp P51858 HDGF_HUMAN
1930	LTSVSSVDFDQR2002000000000	sp Q5T3F8 CSCL2_HUMAN
1931	AQSLVISPPAPSPR2000	sp P46013 KI67_HUMAN
1932	TSSAFVGTPEASPEPK2000200000	sp Q99460 PSMD1_HUMAN
1933	TPSSSSTLAYSPR220000000000	sp Q6Q0C0 TRAF7_HUMAN
1934	KTSFDQSDVDIFPSDFPTEPPSLPR2.000002E+25	sp Q02880 TOP2B_HUMAN
1935	SPLNSCKDPYGGSEGTFSSR2.0002E+20	sp Q8N4S9 MALD2_HUMAN
1936	HSTPSNNSNPSGPPSPNSPHR20000000	sp Q9Y5S2 MRCKB_HUMAN
1937	DSAIPVESDTEDEGAPR20200000000	sp Q96D46 NMD3_HUMAN
1938	STQESLTAGGTDLK20000000000000	sp Q15345 LRC41_HUMAN
1939	VSALEEDMDDVESSEEEEEDEK220000000000	sp Q8NAV1 PR38A_HUMAN
1940	RPSASSPNNNTAAK2000000000000	sp O75151 PHF2_HUMAN
1941	KLTSDEEGEPSGK200000000000	sp Q8WVC0 LEO1_HUMAN
1942	DPTSLLGVLQAEADSTSEGLEDAVHSR22000000000000	sp Q27J81 INF2_HUMAN
1943	KLSSSEPYEEDFNDDQSIKK2.202E+20	sp Q9H4L7 SMRCD_HUMAN
1944	SGAQASSTPLSPTR20000	sp P02545 LMNA_HUMAN
1945	NLNSNLFSVNR2000200000	sp P52948 NUP98_HUMAN
1946	NKLEGSDVDSELEDR20002000000	sp Q92614 MY18A_HUMAN
1947	DGTAPPPQSPGSPGTGQDEEWSDEESPR20000000	sp Q9UKS6 PACN3_HUMAN
1948	SPGSISYLPFFTK2000000000000000	sp O75376 NCOR1_HUMAN
1949	ASFGATGGPPAHLTPSPLSGPGSSLLEK2020000000000000	sp O14686 KMT2D_HUMAN
1950	SIDPDSIQSALLASGLGSK2E+19	sp A3KN83 SBN01_HUMAN
1951	SASQSSLDKLDQELK20020000000000	sp O60271 JIP4_HUMAN
1952	QSPECENFLDVGLGR2000000000000000	sp Q12802 AKP13_HUMAN
1953	LSSTDDGYIDLQFK200000000000	sp Q96A57 TM230_HUMAN
1954	FICVTPTCSNTIDLPMSPR20000000000002000	sp P40763 STAT3_HUMAN
1955	RPAAAAAAGSASPR202000	sp Q96S55 WRIP1_HUMAN
1956	MESEGGADDSAEGLDLDLDDDDNEDRGDDQLELIK1.020000002E+34	sp P07910 HNRPC_HUMAN
1957	SGSAAQAEGLCK20200000000000	sp Q92685 ALG3_HUMAN
1958	TGQAGSLSGSPKPFSPQLSAPITTK2020000000000000000	sp Q9ULU4 PKCB1_HUMAN
1959	KRSSSEDDDLVDESDFDDASINSYSVSDGSTSR2.22000000002E+31	sp Q14839 CHD4_HUMAN
1960	AEEEEASTEEEDKEGAVVSAPSVK2200000000000000000	sp Q14160 SCRIB_HUMAN
1961	VDHGAEIITQSPGR20000	sp P11137 MTAP2_HUMAN
1962	TAVSVTQGGHSR200	sp P40425 PBX2_HUMAN
1963	LPSTDDCPAIGTPLR20000000020000	sp Q15648 MED1_HUMAN
1964	DHSPTPSVFNSDEER20200000200000	sp Q6UN15 FIP1_HUMAN
1965	VSKDDRSDVESSEEDVTCTK2000222000000000000	sp Q8N5B7 CERS5_HUMAN
1966	AHSPAEGASVSSSPGPK200000000000000000	sp Q8NFG4 FLCN_HUMAN
1967	ADEASELACPTPK2000	sp P49327 FAS_HUMAN
1968	HCAPSPDRSPELSSSR2000200000000	sp Q96T37 RBM15_HUMAN
1969	GSPATSPHLGR2000000	sp Q13671 RIN1_HUMAN
1970	AQTPPGPSLSGSK2000000000000	sp Q9UQ35 SRRM2_HUMAN
1971	SPSDSSTASTPVAEQIER2000000000000000	sp Q16643 DREB_HUMAN
1972	DSQDASAEQSDHDDEVASLASGGFGTK2E+24	sp Q6P2E9 EDC4_HUMAN
1973	SSPGAVAGLSNAPGTPR20000000000002000	sp Q9BWG4 SSBP4_HUMAN

1974	STPESGSDKESVGSSTSNEGGR2.002E+20	sp Q6GQQ9	OTU7B_HUMAN
1975	SPQTLAPVGEDAMK20000000000000	sp Q92797	SYMPK_HUMAN
1976	SVASNQSEMEFSSLQDMPK2002010000001000	sp Q8N163	CCAR2_HUMAN
1977	SSAEAQTPEDTPNK20000000000000	sp 043493	TGON2_HUMAN
1978	SQSSEGVSSLSSSPSNSLETQSQSLSR200	sp 095155	UBE4B_HUMAN
1979	GVHIHQAGGSPPASSTSSSSLTNDVAK20000000000000	sp Q8N122	RPTOR_HUMAN
1980	GSFKDDPQLYQEIQER2000000000000000	sp Q5VV41	ARHGG_HUMAN
1981	DSELKTPPSVGPSPVTVTLESAPSALEK2.00002E+21	sp Q96T58	MINT_HUMAN
1982	KKEEPSQNDISPK2000	sp Q9NR30	DDX21_HUMAN
1983	ASIGQSPGLPSTTFK200000000000	sp Q5VT52	RPRD2_HUMAN
1984	DYEPSPSPAPGAPPPPPQR2000000000000000	sp P55196	AFAD_HUMAN
1985	GRGPPSSDSEPEAELER22202000000000	sp Q7Z4V5	HDGR2_HUMAN
1986	ELEENDSENSEFEDDGSEK20020000002000	sp 043719	HTSF1_HUMAN
1987	AEAGAGSATEFQFR2000000000	sp Q9NQ39	RS10L_HUMAN
1988	AGDLEDSPK2000	sp P46783	RS10_HUMAN
1989	IALESEGRPEEQMESDNCSGGDDWHLSSK20000000000000	sp P51858	HDGF_HUMAN
1990	TNSPAYSDISDAGEDGEGKVDVSK2.0002002E+22	sp Q13501	SQSTM_HUMAN
1991	EIQNGNLHESDESVPVR2000000000	sp 015014	ZN609_HUMAN
1992	LQEEGGGDEEETGSPSEDGMQSAR200000000000100000	sp Q86UP2	KTN1_HUMAN
1993	ELVGDTSQEGDHEPSGSETEEDTSSSPHR2.00000002E+23	sp Q13610	PWP1_HUMAN
1994	KPSPSEPEPWKFPFAVSPPEPR2.0002E+20	sp PODJ93	SIM13_HUMAN
1995	AVPMAPAPASPGSSNDSSAR20000000000000	sp Q96JM3	CHAP1_HUMAN
1996	GESAPTLSTSPSPSSPSPSPTLGR202000000000	sp Q66K74	MAP1S_HUMAN
1997	QEIESDESSELDGELQDR20202000000000	sp Q9H330	TM245_HUMAN
1998	GYSPYSVSGSGSTAGSR2000000000000000	sp Q9BYW2	SETD2_HUMAN
1999	DTVIIIVSEPEDEESQGLPTMAR200200002000000000	sp Q15149	PLEC_HUMAN
2000	NTSSDNDVEVMPAQSPR2002000000002000	sp Q9H4L7	SMRCD_HUMAN
2001	TGSVQTPPDLESSEEVDAEAGAPEVVGPK2.000000002E+27	sp Q9HCK8	CHD8_HUMAN
2002	SGDETPGSEVPGDK2000200200000000	sp P28290	SSFA2_HUMAN
2003	LQEEGGGDEEETGSPSEDGMQSAR200000020000000000	sp Q1KMD3	HNRL2_HUMAN
2004	KPSVGVPPPASPSPYPR2000000020000000	sp Q13610	PWP1_HUMAN
2005	ALEVEESNSEDEASFK202000000000	sp Q9P206	K1522_HUMAN
2006	SSSQSPSPGNHTSNSSNASNATVVPQNSSAR2.00202E+31	sp Q03111	ENL_HUMAN
2007	HEAPSPISGQPCGDDQNASPSK200000000000000000	sp Q9BTA9	WAC_HUMAN
2008	IVSGSPISTPSPSPLPR2000000000200000	sp 075376	NCOR1_HUMAN
2009	QLAQSSALASLTGLGGLGGYSGDSEDER30000200000	sp Q9ULM3	YETS2_HUMAN
2010	TGSVQTPPDLESSEEVDAEAGAPEVVGPK2.000000022E+27	sp Q8TF01	PNISR_HUMAN
2011	SPHSPSIATPPSQSSSCDAPMLSTVHSAK2.020002E+28	sp P28290	SSFA2_HUMAN
2012	MESEGGADDSAEGLDLLDDDDNEDRGDDQLELIKDDEK1.000000002E+38	sp Q9NYF8	BCLF1_HUMAN
2013	TEELIESPKLESSEGEIIQTVDR200000000000000000	sp P07910	HNRPC_HUMAN
2014	SKSPNNEGDVHFSR2000000000000000	sp Q99590	SCAFB_HUMAN
2015	MVIQGPSSPQGEAMVTDVLEDQK200000100000000000	sp Q8NAP3	ZBT38_HUMAN
2016	SEDDSAVPLAK20000000000000	sp Q12888	TP53B_HUMAN
2017	SPTGNTPPVIDSVSEK2000000000000000	sp Q9Y2W2	WBP11_HUMAN
2018	SGSMDPSGAHPSVR2000000000000000	sp P25054	APC_HUMAN
2019	KPSDDKHFGQAPSK20000000000000	sp Q07666	KHDR1_HUMAN
2020	ATVPVAAATAAEGEGSPPAVAAVAGPPAAAEEVGGVGGSSR2E+40	sp Q5VT52	RPRD2_HUMAN
2021	SASADNLTLPK2000000000	sp 075179	ANR17_HUMAN
2022	NHETDGGSAHGDDDDGPHFEPVVPLPK2E+22	sp Q8ND76	CCNY_HUMAN
2023	KAASPSQSVR202000000	sp P49792	RBP2_HUMAN
2024	WDKDDFESEEDVK20000000	sp Q8IYB3	SRRM1_HUMAN
2025	TDGSIQDRQPVTVADYISR202000000000000000	sp Q7Z6E9	RBBP6_HUMAN
		sp P51116	FXR2_HUMAN

2026	RPDDVPLSLSPSK20000	sp Q9UFC0 LRWD1_HUMAN
2027	STIGVMVTASHNPEEDNGVK200000000000	sp O95394 AGM1_HUMAN
2028	SEALSSVLDQEEAMEIK200022000000000000	sp Q12888 TP53B_HUMAN
2029	QPSDASETTGLVQR2000000000000	sp O15085 ARHGB_HUMAN
2030	NYQQNYQNSSEGEKNEGSESAPEGQAQR2. 02E+21	sp P67809 YBOX1_HUMAN
2031	VYLSDSDNNSLETGEILK2020000000000000	sp Q9NSI6 BRWD1_HUMAN
2032	SPGPGPGGAGAEPGATGGSSHFISSR2E+27	sp Q9ULI0 ATD2B_HUMAN
2033	GSETGSETHESDLAPSDK200020000000000000	sp P48634 PRC2A_HUMAN
2034	TVSLTPSPTTQVETPDLVDHDNTSPLFR2. 02E+24	sp Q05209 PTN12_HUMAN
2035	TLDSGTSEIVKSPR2000	sp Q9H501 ESF1_HUMAN
2036	AVPMAPAPASPGSSNDSSAR100000200000000000	sp Q66K74 MAP1S_HUMAN
2037	SSPNVANQPPSPGGK200000	sp P55196 AFAD_HUMAN
2038	SPEPDGPAPAFLR2000000000000000	sp Q9Y4F5 C170B_HUMAN
2039	VQNSPPVGR2000000	sp O95359 TACC2_HUMAN
2040	GVHIHQAGGSPPASSTSSSSLTNDVAK200000000000000000	sp Q8N122 RPTOR_HUMAN
2041	SPTESVNTVPVGK20000000000000	sp Q8WXG6 MADD_HUMAN
2042	KGSDDDGGDSPVQDIDTPEVDLYQLQVNTLR2. 0000002E+22	sp O75446 SAP30_HUMAN
2043	RVSEVEEKEPVPQPLPSDDTR2E+20	sp O43290 SNUT1_HUMAN
2044	REEDEENDNDNESDHDEADS200000000	sp Q9NRG0 CHRC1_HUMAN
2045	RDSPLQSGGQNSQAGQR200000000000000000	sp O95819 M4K4_HUMAN
2046	RLSPSSSAASSYSFSDLNSTR200000000000000000	sp P50402 EMD_HUMAN
2047	REEQDTSDGESVTHHIR200000000000	sp P36915 GNL1_HUMAN
2048	KESKEETSIDVAGKPNEVTK2E+19	sp P53985 MOT1_HUMAN
2049	TPLSSLSATSNNSTVACIDR2022002000000000	sp P00533 EGFR_HUMAN
2050	SPQTLAPVGEDAMKTPSPAAEDAR2. 0000000000102E+24	sp Q92797 SYMPK_HUMAN
2051	GARPPAAGPGGDEDEDEEDTAPESALDTSLDK20000000220000	sp Q27J81 INF2_HUMAN
2052	RASPPDPSPSAASASER200000000000000000	sp Q9Y2F5 ICE1_HUMAN
2053	SSSSGDDYAYTQALLLHQR200000000000000000	sp Q8N5C8 TAB3_HUMAN
2054	SKSGSEEVLCDSCIGNK2000000000000000	sp Q14134 TRI29_HUMAN
2055	SPPHSGFQQYQQADASK200000000000000000	sp Q92540 SMG7_HUMAN
2056	RDSSESQLASTESDKPTTGR202000000000000000	sp Q96B23 CRO25_HUMAN
2057	EVDATSPAPSTSSTVK200000000000	sp Q16666 IF16_HUMAN
2058	GPPDFSSDEEREPTPVLGSGAAAAGR2E+20	sp P42166 LAP2A_HUMAN
		sp P42167 LAP2B_HUMAN
2059	KGAGDGSDEEVDGKADGAEAKPAE200000000000000000	sp P35579 MYH9_HUMAN
2060	TSGSDDPGICSNTDSTQAQVLLGK2E+21	sp Q8NDI1 EHBP1_HUMAN
2061	SRSSSPVTELASR222000000000	sp Q9UQ35 SRRM2_HUMAN
2062	RLSGSSEDEEDSGKGEPTAK202200000000000000	sp Q9BW71 HIRP3_HUMAN
2063	ASLGSLEGEAAEASSPK20000000002000	sp Q09666 AHNK_HUMAN
2064	ALFKPPEDSQDDESDDAEEEEQTTK200002020000000000	sp Q13769 THOC5_HUMAN
2065	ASMSEFLESEDGEVEQQR20000000000	sp Q15022 SUZ12_HUMAN
2066	SSSISEEKGSDDEKPR20000000020000000	sp Q9UKV3 ACINU_HUMAN
2067	IIEQLSPAATSR200000000	sp Q03468 ERCC6_HUMAN
2068	GTGSGGQLQDLDCSSSDDEGAAQNSTKPSATK220000000000000000	sp P08240 SRPR_HUMAN
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2070	LAAPSVSHVSPR2000	sp Q8WXE1 ATRIP_HUMAN
2071	QNSPVTQDGIQESPWQDR200000000000000000	sp O75815 BCAR3_HUMAN
2072	DSGSDEDFLMEDDDSDYGSSKK2. 0200000000002E+22	sp Q9H1E3 NUCKS_HUMAN
2073	GPPDFSSDEEREPTPVLGSGAAAAGR2. 20000002E+21	sp P42166 LAP2A_HUMAN
		sp P42167 LAP2B_HUMAN
2074	GEPNVSYICSR300000	sp P49840 GSK3A_HUMAN
		sp P49841 GSK3B_HUMAN
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2077	VMTIPYQMPASSPVICAGGQDR1.0000000002E+22	sp Q15365 PCBP1_HUMAN
2078	GSVSSVSPKPPSSFK2000020000000000	sp Q9NZJ0 DTL_HUMAN
2079	SWGQQAQEYQEYK2000000000000000	sp Q5VZK9 LR16A_HUMAN
2080	LSTTPTSSLHEDGVEDFR22000000000000	sp Q94842 TOX4_HUMAN
2081	TPPGPPPDDEDDPVPLPVSGDK2E+24	sp Q69YN4 VIR_HUMAN
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2083	TPLSFTNPLHSDDSDSDER2002020000	sp Q05209 PTN12_HUMAN
2084	ADEPSSEESDLEIDK220000000000	sp P50502 F10A1_HUMAN
		sp Q8IZP2 ST134_HUMAN
2085	AIGSTSKPQESPK2000	sp P18887 XRCC1_HUMAN
2086	GGTTSWTSGQPSYDSSR2000000000	sp Q99081 HTF4_HUMAN
2087	ESEVCPVTNSPSPPLPPPPLPK2.0000000002E+24	sp Q9ULJ3 ZBT21_HUMAN
2088	EHYPVSSPSPPAQPGGVS20200000000000	sp Q75179 ANR17_HUMAN
2089	VSPSPQESLSSKSDTDTGVCSTDEDPDDK2.02002022E+29	sp Q5JSH3 WDR44_HUMAN
2090	RRGDSESEDEQDSEEV2020000000000000	sp Q9NZ63 CI078_HUMAN
2091	IESDTEETQDTSVDHNETGTGESSVEENEK2E+20	sp Q6PL18 ATAD2_HUMAN
2092	DFQEYVEPGEDFPASPQR20000	sp Q9Y6G9 DC1L1_HUMAN
2093	DDSDDGGGWITPSNIK2000000000000000	sp Q9ULX3 NOB1_HUMAN
2094	SLGPSLATDKS200000000000	sp Q9H3N1 TMX1_HUMAN
2095	AAHTEDINACTLTTSR2000	sp P04049 RAF1_HUMAN
2096	AGMSSNQSISSPVLDAVPR100002002000000000	sp Q9UQ35 SRRM2_HUMAN
2097	GAPSSPATGVLPSQPK20000000200000	sp Q5SRE5 NU188_HUMAN
2098	EVHDELEDLPSPPPLSPPPTTSPHK20000020000000000	sp Q9NQX3 GEPH_HUMAN
2099	SGSLDSSFQELPASSK2000000000000000	sp Q9UMZ2 SYNRG_HUMAN
2100	SEPVKEESSELEQPFQDTSSVGPDR2.2E+19	sp Q15424 SAFB1_HUMAN
		sp Q14151 SAFB2_HUMAN
2101	EDKSPSEESAPTTSPESVSGVSSGSSGR2E+22	sp P54725 RD23A_HUMAN
2102	AYACHAEHENDSDDDDEAEDDDETEELGSEDDIDEDGQEYLEILAK2.00002E+24	sp Q95373 IPO7_HUMAN
2103	KPSDSLVAASSR20020000	sp Q5T0W9 FA83B_HUMAN
2104	EISSPESCPPEIK220000000000	sp Q92889 XPF_HUMAN
2105	VPLAPITDPQQLQLSPLK20000	sp P31350 RIR2_HUMAN
2106	DIKEESDEEEEDDEESGR20000000000000	sp Q96EV2 RBM33_HUMAN
2107	SPGSTPTTSSQAPQK200000000000	sp P35658 NU214_HUMAN
2108	AMDNHSDEEELAAFCPLDDSTVAR2.02E+21	sp Q86VR2 F134C_HUMAN
2109	SFSLASSNSPISQR2000000	sp Q9BXB4 OSB11_HUMAN
2110	HQEVQDQDPVFQGSQSDSSGYQSDHKKK20220002000000	sp Q3B726 RPA43_HUMAN
2111	SDAEEDGGTVSQEEDRKP2.000000002E+20	sp P27824 CALX_HUMAN
2112	VISDESIDIGSDVEFKPDK2.02E+19	sp P52701 MSH6_HUMAN
2113	ASEDTTSGSPPK200000	sp Q9BZZ5 API5_HUMAN
2114	LEGSDDLLSDSEEHSR2000000000	sp Q9NR09 BIRC6_HUMAN
2115	TGSYGALAEITASK300000000000	sp Q14974 MYPT1_HUMAN
2116	SPSGPVKSPPLSPVGTTPVK20002000000000	sp Q9BVC5 ASHWN_HUMAN
2117	SSPELEDTATSSK20000000000000	sp P46013 KI67_HUMAN
2118	EEPLSEEPCTSTAIASPEK2000000000020000	sp Q9Y2X3 NOP58_HUMAN
2119	GPPDFSSDEEREPTPVLGSGAAAAGR2.20000002E+21	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
2120	HASSPESPKPAPAGSHR222002000000000000	sp Q9UQ35 SRRM2_HUMAN
2121	SPSPDGSPAATPEIR2000020000000000	sp Q00499 BIN1_HUMAN
2122	TVSTSTSGNDFEETK22000000000000	sp Q8TF72 SHRM3_HUMAN
2123	NLETLPFSFSDSEEDSVAK220000000000	sp Q9ULL5 PRR12_HUMAN
2124	SDSLSPPR2020000	sp Q9UPQ0 LIMC1_HUMAN
2125	DHSPTPSVFNSDEER20000000200000	sp Q6UN15 FIP1_HUMAN
2126	LVSPGSANETSSILVESVTR200000000000000000	sp Q75369 FLNB_HUMAN

2127	KPSPAQAAETPALELPLPSVPAPAPL2E+24	sp Q8WX92 NELFB_HUMAN
2128	TASGSSVTSLDGTR202020000000000	sp Q92597 NDRG1_HUMAN
2129	GSSGSVDETLFALPAASEPVIR2. 2E+21	sp Q9NQC3 RTN4_HUMAN
2130	HGGSPQPLATTPLSQEPVNPPEASPTR20000	sp Q9UQ35 SRRM2_HUMAN
2131	TSIDSIDSGVELTTSFK20020020000000000	sp O15403 MOT7_HUMAN
2132	HLDGEEDGSSDQSQASGTTGGR220000000000000	sp Q9UNF1 MAGD2_HUMAN
2133	SRSSDIVSSVR20000000000	sp Q14C86 GAPD1_HUMAN
2134	LATNTSAPDLK2000000	sp P27816 MAP4_HUMAN
2135	SEVQQPVHPKPLSPDSR200200	sp Q9UHB6 LIMA1_HUMAN
2136	DGDKSPMSSLQISNEK2000000000000	sp Q9UER7 DAXX_HUMAN
2137	SVSASSGPCSPAAPHPPPPR20002000000000000	sp Q86UY5 FA83A_HUMAN
2138	QSETVDQNSDSEMLAILK202000000000	sp P46100 ATRX_HUMAN
2139	SSSEDDLDVESDFDDASINSYSVSDGSTSR2. 22E+31	sp Q14839 CHD4_HUMAN
2140	KLSSSEPEYEDEFNDDQSIKK2200300000000000000	sp Q9H4L7 SMRCD_HUMAN
2141	DEVFHTDQDDPSSDDEGMPYTRPVK222000000000000	sp Q5JSH3 WDR44_HUMAN
2142	KLSDPCPVESK20000000000	sp O60244 MED14_HUMAN
2143	SHVEDGDIAGAPASSPEAPAEQDPVQLK200000000000000	sp Q9P2E9 RRBP1_HUMAN
2144	SKFDSDEEEDTENVEAASSGK20000000000000000	sp Q8TF01 PNISR_HUMAN
2145	RNSSEASSGDFLDL200000000000000	sp Q9UK76 HN1_HUMAN
2146	LGSVDSFER20000000	sp O60343 TBCD4_HUMAN
2147	NMTVEQLLTGSPSPTVEPEKPTR1. 000000000002E+23	sp Q9C0C9 UBE20_HUMAN
2148	NSLDDLVGPPSNLEGQSDER2E+19	sp Q9Y6Q9 NCOA3_HUMAN
2149	GSSDGRGSDSEDLPHR20200000000	sp Q9UPQ0 LIMC1_HUMAN
2150	SSSNDVDEETAESDTPVLEK2000020000000000000	sp Q99590 SCAFB_HUMAN
2151	AGSSGNSCITYQPSVSGEHK2000000000000000000	sp Q99755 PI51A_HUMAN
2152	THSTSSSLGSGESPF20000000000000000	sp Q9UGV2 NDRG3_HUMAN
2153	GQKPALDCSQVSPRPATSPENNASLSDTSPMDSSPSGIPK2. 2E+24	sp P18858 DNLI1_HUMAN
2154	YYDDIYFDSSEDEDR202000000	sp Q56P03 EAPP_HUMAN
2155	RNSSEASSGDFLDL200000000000000	sp Q9UK76 HN1_HUMAN
2156	SPISDNSGCDAPGNSNPSLSPSSAESEK200000000000000	sp Q99590 SCAFB_HUMAN
2157	NSSPAPPQAPGK2000000000000	sp Q9NYV4 CDK12_HUMAN
2158	TPMYGSQTPLQDGSR200000000	sp O00267 SPT5H_HUMAN
2159	QRPVPQSSASLDEYTLMR20000000000	sp Q9Y4H2 IRS2_HUMAN
2160	SPQQTVPYVVPLSPK2000000000002000	sp Q9UQR0 SCML2_HUMAN
2161	HLDGEEDGSSDQSQASGTTGGR200000000000000	sp Q9UNF1 MAGD2_HUMAN
2162	ESCSSPSTVGSLLTTR20000000000000000	sp Q7Z589 EMSY_HUMAN
2163	SSTPPGESYFGVSSLQLK20000000000000000	sp Q9UQ35 SRRM2_HUMAN
2164	DLDDALDKLSDSLGQR20000000	sp P20810 ICAL_HUMAN
2165	SAAQAAAQTNNAAGK2000000	sp Q8NC51 PAIRB_HUMAN
2166	LGVSVSPSR20000	sp Q86VM9 ZCH18_HUMAN
2167	AYSPTCSPTLGFK2000200000000	sp Q5TGY3 AHDC1_HUMAN
2168	SQEMVHLVNK200000000000	sp P16070 CD44_HUMAN
2169	KISVVSATK20000000	sp Q9UKV3 ACINU_HUMAN
2170	GAPPGSPEPPALLAAPLAAGACPGGR2E+21	sp Q9Y4B5 MTCL1_HUMAN
2171	QASPETSASPDGSQNLVYETELLR2E+22	sp Q14789 GOGB1_HUMAN
2172	TSPLKDNPSPEPQLDDIK2000000200000000000	sp Q96JC9 EAF1_HUMAN
2173	ASGPPVSELITK2000000000000	sp P10412 H14_HUMAN
		sp P16403 H12_HUMAN
		sp P16402 H13_HUMAN
2174	STSAPQMSPGSSDNQSSSPQAQK2000000000000000000	sp Q14157 UBP2L_HUMAN
2175	GPSPSSPTPPAAAAPAEQAPR2. 02E+19	sp Q86VQ1 GLC11_HUMAN
2176	QDPVTYISETDEEDDFMCK2000020000001000	sp P35251 RFC1_HUMAN
2177	FTSQDASTPASSELNK2000200000000000	sp Q9HAW4 CLSPN_HUMAN
2178	RPASVSSAAVEHEQR200000000000000	sp Q7Z5L9 I2BP2_HUMAN

2179	ASESSSEEKDDYEIFVK2220000000000000	sp P18583 SON_HUMAN
2180	QSFDDNDSELEDKDSK200000000000000000	sp O60841 IF2P_HUMAN
2181	SVQEGENPDDGVR2000000000000000	sp P42696 RBM34_HUMAN
2182	TPHVQAVQGPLGSPPK20000	sp Q15149 PLEC_HUMAN
2183	KADTTTPTTAILAPGSPASPPGSLEPK2002000000000	sp P25440 BRD2_HUMAN
2184	KGSSSSVCSVASSDISLGSTK200000000000000000	sp Q9ULT8 HECD1_HUMAN
2185	TASPPPPPK2020000000	sp Q8IYB3 SRRM1_HUMAN
2186	APAGQEEPGTPSSPLSAEQLDR2000000000000000	sp P13051 UNG_HUMAN
2187	IGFPETEEELIEIASENSDFIFPSAPDVK2002000000000000	sp Q9Y3F4 STRAP_HUMAN
2188	DELNEELIQEESSEDEGEYEEVRK2200000000000000	sp O60524 NEMF_HUMAN
2189	GSAPHSESDLPEQEEIILGSDDEQEDPNDYCK2.000002E+32	sp Q96SB4 SRPK1_HUMAN
2190	KHSPSPPPPTPTESR2020000000000000	sp Q92922 SMRC1_HUMAN
2191	NYQQNYQNSSEGEKNEGSESAPEGQAQQR3E+24	sp P67809 YBOX1_HUMAN
2192	TPSDDEEDNLFAPPK2000000000000000	sp Q641Q2 FA21A_HUMAN
2193	LPPTFFCGVCSDTDEDNGNGEDFQSELQK2.02E+19	sp Q9Y4E1 FA21C_HUMAN
2194	SVASSQPAKPTK20020000000000	sp P49792 RBP2_HUMAN
2195	ATSAGSSPSCSLAGR200000000000	sp Q07157 ZO1_HUMAN
2196	GRASPGGVSTSSSDGK2000000000000000	sp Q9BXB5 OSB10_HUMAN
2197	VEENPDADSDFDAKSSADDEIEETR200000220000000000	sp P54259 ATN1_HUMAN
2198	RASGQAFELILSPR2000000002000	sp Q9H0G5 NSRP1_HUMAN
2199	SLSGSSPCPK202000000000	sp P16949 STMN1_HUMAN
2200	SCTPSPDQISHR2000000000	sp Q9UQ35 SRRM2_HUMAN
2201	TLSSSAQEDIIR2000000000	sp Q7Z2W4 ZCCHV_HUMAN
2202	KPSPSESPKPFPAVSPPEPR20000000000200000	sp Q9H788 SH24A_HUMAN
2203	TTSSDLTSSSSSGPR2000000000000000	sp Q96JM3 CHAP1_HUMAN
2204	QRSPSPAPAPAAAAAGPPTR2.02E+19	sp Q5QP82 DCA10_HUMAN
2205	QAPGVGAVGGGSPER20000	sp Q9H7N4 SFR19_HUMAN
2206	GAQASSGSPALPR2000000	sp Q96G74 OTUD5_HUMAN
2207	KKPGDASSLPDAGLSPGSQVDSK2000000000	sp P42684 ABL2_HUMAN
2208	SPGHMVIDLQTK200010000000	sp Q13459 MYO9B_HUMAN
2209	LLKPGEEPSEYTDEEDTK20000000	sp Q9NWH9 SLTM_HUMAN
2210	DKSPVREPIDNLTPEER2000000000000000	sp O15173 PGRC2_HUMAN
2211	QNPEQSADEDAEKNEEDSEGSSDEDEDEDGVSAAATFLK2.000000000002E+33	sp Q14498 RBM39_HUMAN
2212	LTFDSSFSPNTGK2000000	sp B5ME19 EIFCL_HUMAN
2213	CVSVQTDPTDEIPTK2000000000000000	sp Q99613 EIF3C_HUMAN
2214	TASISSPSEGTPVGSYGCTPQSLPK2.2E+22	sp P21796 VDAC1_HUMAN
2215	HNSASVENVSLR200000000000	sp P18583 SON_HUMAN
2216	RLSPSSSAASSYSFSDLNSTR2E+20	sp Q6PKG0 LARP1_HUMAN
2217	KPSTSDSDSNFEK20200200000000	sp Q9Y4H2 IRS2_HUMAN
2218	SPSGPVKSPPLSPVGTTPVK2000020002000000000	sp P50402 EMD_HUMAN
2219	EEEEGISQESSEEEQ2002200000	sp P11388 TOP2A_HUMAN
2220	KGDRSPEPGQTWTR200000000000	sp Q9BVC5 ASHWN_HUMAN
2221	TGSGSPFAGNSPAR2000000020000	sp P17096 HMGA1_HUMAN
2222	SSLGQASETEEDTVSVSK20002020000000000000	sp Q09666 AHNK_HUMAN
2223	YSDSDELTVQR20000000000000	sp Q9UPT8 ZC3H4_HUMAN
2224	SPSPGNHTSNSSNASNATVVPQNSSAR200200000000000000	sp Q13435 SF3B2_HUMAN
2225	CTLPEHESPSQDISDACEAESTER2000000000000000	sp Q8NFC6 BD1L1_HUMAN
2226	ASGQAFELILSPR2000000002000	sp Q9BTA9 WAC_HUMAN
2227	KNSTDLDSAPEDPTSPK200000000020000	sp Q32MZ4 LRRF1_HUMAN
2228	RLSESSALKQPATPTAAESSEGEEGEGDDGGGETESR2.000000202E+31	sp P16949 STMN1_HUMAN
2229	SLHLSPQEQSASYQDR20000000000000	sp Q96RK0 CIC_HUMAN
2230	ESPRPLQLPGAEGPAISDGEEGGGEPGAGGAAGAAGAGR2E+24	sp Q96S55 WRIP1_HUMAN
		sp Q9H410 DSN1_HUMAN
		sp Q6PKG0 LARP1_HUMAN

2231	SDSGGSSEPFDR200200000000	sp	P48634	PRC2A_HUMAN
2232	SPEACCELTLQPLR20000000000000	sp	P06132	DCUP_HUMAN
2233	NSPNNISGISNPPGTPR2000000000000000	sp	Q9BWW4	SSBP3_HUMAN
2234	SRTASGSSVTSLDGTR2020000000000000	sp	Q92597	NDRG1_HUMAN
2235	GDQCCYSHSPPTPR2000000	sp	Q9NXH9	TRM1_HUMAN
2236	VQGEAVSNIQENTQTPTVQEESEEEEEVDGVEVK2E+29	sp	E9PAV3	NACAM_HUMAN
		sp	Q13765	NACA_HUMAN
2237	DATPPVSPINMEDQER2000200000000000	sp	P17275	JUNB_HUMAN
2238	SRSDIDVNAAGAK20000000000000	sp	075122	CLAP2_HUMAN
2239	AITSLGGGSPK2000	sp	Q6KC79	NIPBL_HUMAN
2240	HIKEEPLSEEPCTSTAIASPEK20000000000020000	sp	Q9Y2X3	NOP58_HUMAN
2241	SPSFASEWDEIEK200000000000	sp	Q92625	ANS1A_HUMAN
2242	HASSPESPAPAPGSHR200002000000000000	sp	Q9UQ35	SRRM2_HUMAN
2243	ALSSDSILSPAPDAR220000200000000	sp	Q8IVT2	MISP_HUMAN
2244	QPPVSPGTALVGSQK200000000000	sp	P17096	HMGA1_HUMAN
2245	SRSHTSEGAHLDITPNSGAAGNSAGPK2. 2E+23	sp	Q92597	NDRG1_HUMAN
		sp	P10412	H14_HUMAN
2246	KASGPPVSELITK200000000000	sp	P16403	H12_HUMAN
		sp	P16402	H13_HUMAN
2247	AAHPENSDEQDVLTKPVR2020000000000000	sp	Q9H2Y7	ZN106_HUMAN
2248	LDSSPSVSTLAAK22000000000000	sp	Q13428	TCOF_HUMAN
2249	TSPADHGGSVGSESGSAVDSVAGEHSVSGR2. 0000002E+30	sp	Q5T4S7	UBR4_HUMAN
2250	KPVTVSPTTPTSPTGEAS2000000000	sp	Q9Y6G9	DC1L1_HUMAN
2251	SGSSPDSEITELKFPSINH200200000000000000	sp	P17812	PYRG1_HUMAN
2252	GPSSPSTPPAAAAPAEQAPR200200000000000000	sp	Q86VQ1	GLCI1_HUMAN
2253	TQVLSPDSLFTAK2000000000	sp	P55196	AFAD_HUMAN
2254	ELLDTSFEDLSKPK2000000000	sp	P78345	RPP38_HUMAN
2255	GDSETDLEALFNAVMPK20000000010000	sp	P46937	YAP1_HUMAN
2256	SILSPGGSCGPIK200000000000	sp	P78347	GTF2I_HUMAN
2257	REESPMVDVQSPSAQDTQSIASDGTPQGEK1. 000000000002E+26	sp	Q7Z6Z7	HUWE1_HUMAN
2258	KSPSPAGGSSPYSR202000000200000	sp	Q14004	CDK13_HUMAN
2259	KSPVFSDESDLDLDFDISK200020000000000000	sp	000566	MPP10_HUMAN
2260	MYSEDKPLSSESLSESEYIEEVR2202020000000000	sp	Q5BKY9	F133B_HUMAN
2261	EFGDSDENEMEEHELK200000000000	sp	Q5F1R6	DJC21_HUMAN
2262	ILLVDSPGMGNADDEQQEEGTSSK2E+19	sp	Q9UKN8	TF3C4_HUMAN
2263	KPNAGGSPAPVR2000000	sp	Q3KQU3	MA7D1_HUMAN
2264	QPATPTAAESSEGEEGEGDDGGETESR2. 00002E+22	sp	Q96S55	WRIP1_HUMAN
2265	HAPSPEPAVQGTGVAGVPEESGDAAIIPAK2E+27	sp	P13051	UNG_HUMAN
2266	RFGDSSSDNGFSSTGTPAKPTVEK2. 202E+22	sp	Q8IU60	DCP2_HUMAN
2267	TPPPPTLSPAATVGR200002000000000000	sp	Q8NDX5	PHC3_HUMAN
2268	SPSTYLHTPTSEDAAIIPSK200000000000000000	sp	Q13111	CAF1A_HUMAN
2269	EPTPSIASDISLPIATQELR200002000000000000	sp	A0FGR8	ESYT2_HUMAN
2270	SNSSSEAVLQEELSAQAK2000000000000000	sp	Q9BXF6	RFIP5_HUMAN
2271	SKSPPKVPIVIQDDSLPAGPPPQIR2. 02E+25	sp	Q7Z422	SZRD1_HUMAN
2272	SLESSVIMDR22000000	sp	Q9H2Y7	ZN106_HUMAN
2273	KPEEESSESEEGSESEEEAPAGTR2. 022000202E+19	sp	Q13428	TCOF_HUMAN
2274	EPTPSIASDISLPIATQELR200000002000000000	sp	A0FGR8	ESYT2_HUMAN
		sp	P05386	RLA1_HUMAN
2275	KEESEESDDMGFLFD2002000000000000	sp	P05387	RLA2_HUMAN
		sp	Q8IY81	SPB1_HUMAN
2276	ALDISLSSGEEDEGDEEDSTAGTTK2. 200000000002E+19	sp	Q96T23	RSF1_HUMAN
2277	GRSTDEYSEADEEEEEEGKPSR2. 0032E+20	sp	Q9NYF8	BCLF1_HUMAN
2278	SSATSGDIWPLSAYDNSPR2000	sp	Q9NYF8	BCLF1_HUMAN
2279	LLSPQMSGEEEDSLAAK200000000000	sp	P42858	HD_HUMAN



2280	EEEEDEDFAHHSDNEQNR2000000020000000	sp Q641Q2 FA21A_HUMAN
2281	TRSPSPTLGESLAPHK2020000000000000	sp Q5SNT6 FA21B_HUMAN
2282	TASQGPQTDSVIQNSENIK2000000000000000	sp Q86UU1 PHLB1_HUMAN
2283	KLSSSSEPYEDEFNDDQSIKK2003000000000000	sp Q9NZN5 ARHGC_HUMAN
2284	AGSFTGTSDPEAAPAR2000000000000000	sp Q9H4L7 SMRCD_HUMAN
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2286	GHYEVTGSDDDETGK20000000	sp Q8WVM8 SCFD1_HUMAN
2287	LAALKDEPQTVPDVPSFGESPPLSPIDMDTQER2000200000000000	sp Q09666 AHNK_HUMAN
2288	LDADAVSPTVSQLSAVFEK2000000000000000	sp P17535 JUND_HUMAN
2289	KASSDSEDSSEEEEEVQGPPAK2. 22020022E+21	sp Q96SB3 NEB2_HUMAN
2290	NYQQNYQNSSEGEKNEGSESAPEGQAQQR2E+21	sp Q14978 NOLC1_HUMAN
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2292	VDIDTPDIDIHGPEGK2000000000000000	sp 095218 ZRAB2_HUMAN
2293	SPVPSAFSDQSR2000000000000000	sp Q09666 AHNK_HUMAN
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2295	LLGSDSEEEQK202000000	sp Q96RK0 CIC_HUMAN
2296	KYSASSGGLCEEATAAK2000000000000000	sp Q9BW71 HIRP3_HUMAN
2297	ICTLPPSPPLASLAPVADSSTR2000000000000000	sp Q9UHV7 MED13_HUMAN
2298	SLSDNGQPGTDPADSGGTSK2000000000000000	sp Q9UER7 DAXX_HUMAN
2299	MPQDGSDDDEEWPTLEK10000200000000000000	sp 075691 UTP20_HUMAN
2300	HDSPDPSPPR200020000	sp Q13895 BYST_HUMAN
2301	GEDSAEETEAKPAVVAPAVVEAVSTPSAAFPSDATAEQGPILTK2E+20	sp Q9BRD0 BUD13_HUMAN
2302	YSPSQNSPIHHIPSR2000020000000000	sp Q12906 ILF3_HUMAN
2303	ASVSGPNSPSETR20002000000	sp Q9NYF8 BCLF1_HUMAN
2304	SPSAMQQQDGLDR2000000000000000	sp 015344 TRI18_HUMAN
2305	SPQDPVGTPTIFKPKQSK20000000000000000000	sp 060343 TBCD4_HUMAN
2306	SAGAPASVSGQDADGSTSPR2000	sp P46013 KI67_HUMAN
2307	SIQDLTVGTGTEPGQVSSR2000000000000000	sp P42679 MATK_HUMAN
2308	SSSPGKPAVSSLNSSHRSR20000000000000000000	sp 043318 M3K7_HUMAN
2309	SNSPLVPPSK2000000000	sp Q9UHB7 AFF4_HUMAN
2310	LASPSGSTSSGLEVVAPEGTSAPGGGPGTLDDSATICR2. 022E+34	sp Q13247 SRSF6_HUMAN
2311	QGSITSPQANEQSVTPQR20000000000020000	sp Q13263 TIF1B_HUMAN
2312	TNSAEVTPPVLSVMGEATPVSIEPR2. 000000002E+23	sp Q9UQ35 SRRM2_HUMAN
2313	SPSVKSPTPK200000200000	sp Q6PJG2 EMSA1_HUMAN
2314	LQPSSPENSLDPFPPR2000000000000000	sp Q9NVD7 PARVA_HUMAN
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2316	EEPLSEEPCTSTAIASPEKK200000000000200000	sp Q7Z417 NUFP2_HUMAN
2317	LEASDCDHQQNSPTLERPGR2000000000	sp Q9Y2X3 NOP58_HUMAN
2318	LQQQAALSPTTAPAVSSVSK2000000000000000	sp Q9HAN9 NMNA1_HUMAN
2319	SGLTVPTSPK2000	sp Q8WXI9 P66B_HUMAN
2320	LLEDSEESSEETVSR200000000000	sp Q53EL6 PDCD4_HUMAN
2321	QVSTENDSTLVHR20000000000	sp 060231 DHX16_HUMAN
2322	EFVEDGEVDESISDFEDMDK200200001000	sp 075592 MYCB2_HUMAN
2323	SPSLGSDLTFATR2000000000000000	sp Q9BXY0 MAK16_HUMAN
2324	SSSPELVTHLK2000000000	sp Q13425 SNTB2_HUMAN
2325	KLSSSSEPYEDEFNDDQSIK22200000000000000000	sp Q07960 RHG01_HUMAN
2326	IESDEEEDFENVGK2000000000000000	sp Q9H4L7 SMRCD_HUMAN
2327	VPPAPVPCPPSPGPSAVPSSPK2000	sp Q96T23 RSF1_HUMAN
2328	KEPAITSQNSPEAR200000	sp 095817 BAG3_HUMAN
2329	KPPAPSPVQSQSPSTNWSPAVPVKK2. 000200000002E+20	sp P23193 TCEA1_HUMAN
2330	THSTSSSLGSGESPFGR2200000000000000	sp Q8IYB3 SRRM1_HUMAN
2331	SPFNPSPPQDSR20000000	sp Q9UGV2 NDRG3_HUMAN
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		sp Q9C0B5 ZDHC5_HUMAN

2333	HNSASVENVSLR200000000	sp Q9Y4H2 IRS2_HUMAN
2334	SPAVATSTAAPPPSSPLPSK2000000	sp Q14157 UBP2L_HUMAN
2335	DEETESEYDSEHENSEPVTNIR2.0020002E+20	sp Q15154 PCM1_HUMAN
2336	EYIPGQPPLSQSSDSSPTR20000000	sp P07814 SYEP_HUMAN
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2338	AGAPGALSPSYDGGLHGLQSK2000000000000000	sp P15923 TFE2_HUMAN
2339	TNPEVMVVEDLSDDGK200200000	sp Q96GN5 CDA7L_HUMAN
2340	SQSSDTEQQSPTSGGGK2200000000000000	sp P23588 IF4B_HUMAN
2341	RNSLTGEEQLAR2000000000000	sp Q9BX95 SGPP1_HUMAN
2342	STPSHGVSSSLNSTGSLSPK2E+20	sp Q9UBC2 EP15R_HUMAN
2343	LGSAEQSSPDL SFLVVK2000000000000000	sp Q96LW4 PRIPO_HUMAN
2344	DSL GDFIEHYAQLGPSSPEQLAAGAEEGGPR2000000000000000	sp Q96QR8 PURB_HUMAN
2345	EWYQSTIPQSPSPAPDDPEEGR20000000000000	sp Q08499 PDE4D_HUMAN
2346	SRSPLDKDTYPPSASVVGASVGGHR2.00000002E+25	sp Q96T37 RBM15_HUMAN
2347	TASTNNTAQR20000000000	sp Q9UBI6 GBG12_HUMAN
2348	SSQSPATAVPASDSPPTK200200000000000000	sp Q3KQU3 MA7D1_HUMAN
2349	GSPATSPHLGR20002000000	sp Q13671 RIN1_HUMAN
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		sp Q09160 1A80_HUMAN
2350	KGGSYSQAASSDSAQGSVDVSLTACK2E+20	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
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2352	NEEDSEGSSEDEDEDGVSAAATFLK22000000000000000000	sp B5ME19 EIFCL_HUMAN
		sp Q99613 EIF3C_HUMAN
2353	NDSVIVADQTPPTR2020000	sp P15336 ATF2_HUMAN
2354	LLGPSAAADILQLSSSLPLQSR2020000000	sp Q7Z6Z7 HUWE1_HUMAN
2355	SLSNSNPDISGTPTSPDDEV2.00000000000002E+21	sp Q96N67 DOCK7_HUMAN
2356	TVSPGSVSPIHGQGVVENLK2.0000002E+21	sp Q9NZM3 ITSN2_HUMAN
2357	LSASTASELSPK2000	sp Q3KQU3 MA7D1_HUMAN
2358	RPEGPGAQAPSSPR2000	sp P40222 TXLNA_HUMAN
2359	HASSPESPKPAPAGSHR22200000000000000000	sp Q9UQ35 SRRM2_HUMAN
2360	KGNAEGSSDEEGLVIDEPAK2000000000000000	sp P51858 HDGF_HUMAN
2361	TYSDEANQMR200000000	sp Q9BXF6 RFIP5_HUMAN
2362	GDMSDEDDENEFFDAPEIITMPENLGHK2E+25	sp P22059 OSBP1_HUMAN
2363	TEMDKSPFNPSPPQDSPR20000020000000	sp P08651 NFIC_HUMAN
2364	ESEDKPEIEDVGSDEEEKKGDK2E+23	sp P07900 HS90A_HUMAN
2365	LDSPPPSPITEASEAAEAEAGNLAVSSR2.002E+20	sp Q9C0C2 TB182_HUMAN
2366	ASAPSPNAQVACDHCLK20000000000000	sp Q14258 TRI25_HUMAN
2367	SPTAQSPTPSVEADSPDQK200000000000000000	sp Q96HH9 GRAM3_HUMAN
2368	TASGSSVTSLDGTR220000000000	sp Q92597 NDRG1_HUMAN
2369	AASSSPGSPVASSPSR2200000000000000	sp Q92538 GBF1_HUMAN
2370	DNSPPPAFKPEPPK20000000000000	sp Q9UDY2 ZO2_HUMAN
2371	SQSLPTLLSPVR200000020000	sp P28290 SSFA2_HUMAN
2372	LSSTDDGYIDLQFK20000000000000	sp Q96A57 TM230_HUMAN
2373	KEQSEVSVSPR202000	sp Q7L4I2 RSRC2_HUMAN
2374	RGESLDNLDSPR2000000000	sp Q8WWI1 LMO7_HUMAN
2375	YSPPAISPLVSEK20000000000000	sp Q14694 UBP10_HUMAN
2376	GVTQEVDGLSQT DGLTYFDK20000000000000	sp Q8NB49 AT11C_HUMAN
2377	SLSNSNPDISGTPTSPDDEV200002000000000000	sp Q96N67 DOCK7_HUMAN
2378	SLDGGSEPAVCEMCGIVGTR200000000000000000	sp Q969R5 LMBL2_HUMAN
2379	SANQSPQSVGSSGVDSGVSTSDGLR200000000000	sp Q14693 LPIN1_HUMAN
2380	GSAPHSEDLPEQEEIILGSDDEQEDPNDYCK2.00000000002E+26	sp Q96SB4 SRPK1_HUMAN

2381	AQSSPASATFPVSVQEPPTKPR2E+20	sp	P56524	HDAC4_HUMAN
2382	HSVTAATPPPSPTSGESGDLLSQSPSSAK2E+19	sp	Q96QU8	XPO6_HUMAN
2383	GAADVEVEPEPPPGVAVSR200	sp	Q8NCF5	NF2IP_HUMAN
2384	TLSPTPSAEGYQDVR20002000000000	sp	O14639	ABLM1_HUMAN
2385	RLSAQGGQAI SVVGLS SMSPLEEEAPQAK2E+27	sp	Q92797	SYMPK_HUMAN
2386	SPSGPPAMDLEGPR20000000000000	sp	O96018	APBA3_HUMAN
2387	LFDHPESPTPNPTEPLFLAQAEVYK200000000000000000	sp	P49327	FAS_HUMAN
2388	LLSSESEDEEEFIPLAQR200200000000000000	sp	Q96AY2	EME1_HUMAN
2389	DLESCSDDDNQGSK2000000000	sp	Q5T3I0	GPTC4_HUMAN
2390	IADPEHDHTGFLTEYVATR300000	sp	P27361	MK03_HUMAN
2391	SVEDVRPHHTDANNQSACFEAPDQK2E+25	sp	Q9Y520	PRC2C_HUMAN
2392	SLKESEQESEEEILAQK20002000000000	sp	Q9BW71	HIRP3_HUMAN
2393	HPASLTSSGSSGSPSSSIK20000000	sp	Q9COD5	TANC1_HUMAN
2394	SRSPTPPSSAGLGSNSAPPIPSR2. 020022E+22	sp	Q8IWX8	CHERP_HUMAN
2395	GRLTPSPDIIVLSDNEASSPR200000020000000000	sp	Q8WXI9	P66B_HUMAN
2396	AASPSQSVR2000000000	sp	Q8IYB3	SRRM1_HUMAN
2397	SVTSNQSDGTQESCESPDVLDLDR200000000000	sp	Q9UGV2	NDRG3_HUMAN
2398	STDEYSEADEEEEEEGKPSR2. 2E+21	sp	Q96T23	RSF1_HUMAN
2399	GSGDTSSLIDPDTLSSELR20000000000000000000	sp	Q9Y608	LRRF2_HUMAN
2400	SCTPSPDQISHR20200000000	sp	Q7Z2W4	ZCCHV_HUMAN
2401	GPPDFSSDEEREPTPVLGSGAAAAGR2. 00000002E+21	sp	P42167	LAP2B_HUMAN
2402	DSGRGDSVSDSGSDALR200000000000	sp	P42166	LAP2A_HUMAN
2403	DMESPTKLDVTLAK200000000000	sp	Q53EL6	PDCD4_HUMAN
2404	EHYPVSSPSPSPAQPGGVSRR2200000000000000	sp	P27816	MAP4_HUMAN
2405	AAVLSDSEDEEK202000000	sp	O75179	ANR17_HUMAN
2406	KPGSPGSESPK2000	sp	Q96ST2	IWS1_HUMAN
2407	SGTSSPQSPVFR200000	sp	Q96JM3	CHAP1_HUMAN
2408	SPCETISSPSTLESK202000000000	sp	Q8N556	AFAP1_HUMAN
2409	QSPPISTSK200000000	sp	Q9COD5	TANC1_HUMAN
2410	DSALAEPEGLSPAPPAR20000000	sp	P33981	TTK_HUMAN
2411	ESTQLSPADLTEGKPTDPSK20000000000000000000	sp	Q9H9J4	UBP42_HUMAN
2412	AHSPAEGASVSSSPGPK20000000000200000	sp	Q08J23	NSUN2_HUMAN
2413	AMDNHSDEEELAAFQQLDDSTVAR1. 000202E+25	sp	Q8NFG4	FLCN_HUMAN
2414	KDSSGSISEQLPCR20000000000000	sp	Q86VR2	F134C_HUMAN
2415	LAPVSPPEPQKAPVSPESVK20000000002000000	sp	Q8NEZ4	KMT2C_HUMAN
2416	KPISDNSFSSDEEQSTGPIK2022000000000000	sp	Q96JM3	CHAP1_HUMAN
2417	LPSTSGSEGVFR2000000000	sp	O60293	ZC3H1_HUMAN
2418	SAYQEYSDSDVPEELKR202000000000	sp	P29317	EPHA2_HUMAN
2419	SYLEGSSDNQLKDESTEPVDDR220000000000000000	sp	Q96JG6	CC132_HUMAN
2420	VAASPKSPTAALNESLVECPK20000200000000000000	sp	Q9UHI6	DDX20_HUMAN
2421	QGEDLAHVQHPTGAGPHAQEEDSQEEEEDEEAASR2E+25	sp	Q53EZ4	CEP55_HUMAN
2422	YLLGDAPVSPSSQK2000000	sp	Q9NW97	TMM51_HUMAN
2423	GKSSPICSTTGDDK200000000000	sp	Q9NYB0	TE2IP_HUMAN
2424	VLGSEGEEDEALSPAK2000000000000000	sp	Q12802	AKP13_HUMAN
2425	LGVSVSPSR2020000	sp	P18858	DNLI1_HUMAN
2426	AKPVVSDFDSEEQDER20002000000000	sp	Q86VM9	ZCH18_HUMAN
2427	HQQSDLSAACSDADLHR2002000200000000	sp	P51531	SMCA2_HUMAN
2428	GGSYSQAASSDSAQGSVSLTACK2000000000000000	sp	Q9H9J4	UBP42_HUMAN
		sp	P05534	1A24_HUMAN
		sp	P30447	1A23_HUMAN
		sp	P01891	1A68_HUMAN
		sp	P01892	1A02_HUMAN
		sp	Q09160	1A80_HUMAN
		sp	P10316	1A69_HUMAN

2429	NVRSDISDQEEDSEEGCPVSINLSK2. 002E+23	sp Q15154 PCM1_HUMAN
2430	AAAAAATAPPSPGPAQPGPR20000000000	sp Q6SPF0 SAMD1_HUMAN
2431	NVPHEICEDSDIDGDIR200000000	sp 000629 IMA3_HUMAN
2432	GSAPHSESDLPEQEEELGSDDDDEQEDPNDYCK2. 02000000000002E+28	sp Q96SB4 SRPK1_HUMAN
2433	MYSEDKPLSSESLSESEYIEEVR1. 0000000220202E+23	sp Q5BKY9 F133B_HUMAN
2434	ALSSDSILSPAPDAR200000000	sp Q8IVT2 MISP_HUMAN
		sp 015320 CTGE5_HUMAN
		sp A4D2H0 CTGEF_HUMAN
		sp Q86UF2 CTGE6_HUMAN
2435	SNSELEDEILCLEK2000000000000	sp P0CG41 CTGE8_HUMAN
		sp A4FU28 CTGE9_HUMAN
		sp Q8IX94 CTGE4_HUMAN
2436	DPSASPGDAGEQAIR200000000000000	sp P18206 VINC_HUMAN
2437	LPSDEDESGTEESDNTPLK2000020000000000000	sp Q9ULH0 KDIS_HUMAN
2438	GDPRLSPDPVAGSAVSQELR20000000000000000	sp Q9BUR4 WAP53_HUMAN
2439	DVDDGSGSPHSPHQLSSK200000000	sp Q9H4A3 WNK1_HUMAN
2440	ASGQAFELILSPR200000000000000	sp P16949 STMN1_HUMAN
2441	ELSESEDEELQLEEFMPLK2002000000000000000	sp Q99543 DNJC2_HUMAN
2442	ASPSTAGETPSGVK200000000000000	sp Q9UI10 EI2BD_HUMAN
2443	DSQQLINIPSLNDSSEIEDISEIESNR2. 0002E+19	sp Q9UDY2 ZO2_HUMAN
2444	MESEGGADDSAEGLDLDNEDR1. 020000002E+25	sp P07910 HNRPC_HUMAN
2445	SELSQDAEPAGSQETK200000000000000	sp Q9BVJ6 UT14A_HUMAN
2446	RPQSPGASPSQAER200020000000	sp Q96T58 MINT_HUMAN
2447	ELVGDGTSQEGDHEPSGSETEEDTSSSPHR20202000000000000	sp P0DJ93 SIM13_HUMAN
2448	QGSITSPQANEQSVTPQR20000000020000	sp Q9UQ35 SRRM2_HUMAN
2449	SRSPLDKDYPPSASVVGASVGGHR2. 02E+25	sp Q96T37 RBM15_HUMAN
2450	EFVEDGEVDESISDFEDMDK200200000000	sp Q9BXY0 MAK16_HUMAN
2451	KSSTVATLQGTPDHGDPR20000000000000000	sp Q8IVT2 MISP_HUMAN
2452	VSIGNITLSPAIR2000000	sp Q0JRZ9 FCHO2_HUMAN
2453	SNSTTQVSQPR200000000	sp Q9UPN4 CP131_HUMAN
2454	HNFLFEDNMALPSESVSSLTDLKPPTGSNQASPAR2. 02E+21	sp Q9BZQ8 NIBAN_HUMAN
2455	DSAI PVESDTDDEGAPR20000000000	sp Q96D46 NMD3_HUMAN
2456	SQSLPTLLSPVR202000000000000	sp P28290 SSFA2_HUMAN
2457	GPSQEEEDNHSDEEDRASEP200000020000	sp Q86VM9 ZCH18_HUMAN
2458	TPSSQNLALLAR2000000000000	sp 075385 ULK1_HUMAN
2459	DSELSDTDSGCCLGQSESDK2. 0000002E+19	sp Q96T88 UHRF1_HUMAN
2460	DLDEEGSEKELHENVLDK2000000000000	sp 043719 HTSF1_HUMAN
2461	NAPAAVDEGSI SPR202000	sp P28715 ERCC5_HUMAN
2462	RESEKLENETLNK2002000000000	sp P23588 IF4B_HUMAN
2463	AGVRPSSSGSAWEACSEAPSK20000000000000000	sp Q12789 TF3C1_HUMAN
2464	TLSTHSVPNISGATCSAFASPFGCPYSHR2. 02E+29	sp P28290 SSFA2_HUMAN
2465	NSPLDCGSASP20000	sp 075179 ANR17_HUMAN
2466	LLSSNEDDANILSSPTDR2000000000200000	sp 075448 MED24_HUMAN
2467	SPSPPTQHTGQPPGQPSAPLSAPR2. 02E+26	sp Q68EM7 RHG17_HUMAN
2468	LNETELTDLEGQESPPK20000	sp Q9ULF5 S39AA_HUMAN
2469	DVDDGSGSPHSPHQLSSK2002000000000	sp Q9H4A3 WNK1_HUMAN
2470	TSPADHGGVSGSESGSAVDSVAGEHSVSGR2. 0000002002E+30	sp Q5T4S7 UBR4_HUMAN
2471	EHYPVSSPSSPPAQP GGVSRR2002000000000000000	sp 075179 ANR17_HUMAN
2472	DTSATSQVNGSPQAEQPSLESTSK20000000000000000	sp Q86WB0 NIPA_HUMAN
2473	DKSPSSLLEDAK2000000000000	sp P42684 ABL2_HUMAN
2474	QLSESESSLEMDDER20000000000000000	sp Q16204 CCDC6_HUMAN
2475	EVSSRPSTPGLSVVSGISATSEIPNK2. 0002E+20	sp Q14C86 GAPD1_HUMAN
2476	VLDEEGSER2000	sp 043719 HTSF1_HUMAN

2477	AGGLDWPEATEVSPSR20000	sp Q9Y446 PKP3_HUMAN
2478	GNKSPSPDGS PAATPEIR20000002000000000	sp 000499 BIN1_HUMAN
2479	ALSPGLEESGLGLLAR200000000000000	sp 015417 TNC18_HUMAN
2480	TSSGTSLSAMHSSGSSGK2020000000000000	sp Q12888 TP53B_HUMAN
2481	NHSNAQFIESYVCR200000000000000	sp P41229 KDM5C_HUMAN
2482	SLPVSVPVWGFK200000000000000	sp Q96B36 AKTS1_HUMAN
2483	DTSPSSGSAVSSSK2000000000000000	sp Q8NEY8 PPHLN_HUMAN
2484	TGSNISGASSDISLDEQYK2000000000000000	sp P22059 OSBP1_HUMAN
2485	EDENDASDEDDDEKR20000000000	sp Q9Y5B6 PAXB1_HUMAN
2486	ASLGSLEGEAEAEASSPK20000000000002000	sp Q09666 AHNK_HUMAN
2487	ETPHSPGVEDAPIAK200000000000000	sp Q9UHB6 LIMA1_HUMAN
2488	KPPAPSPVQSQSPSTNWS PAVPK2. 000202000002E+19	sp Q8IYB3 SRRM1_HUMAN
2489	GHHVTDSENDEPLNLNASDSESEELHR2. 000000000202E+21	sp Q96ST2 IWS1_HUMAN
2490	GNDPLTSSPGR20000	sp P49736 MCM2_HUMAN
2491	TTQSMQDFPVVDSEEEAEFEFK200000000000	sp Q9UBB9 TFP11_HUMAN
2492	VISDSESDIGGSDVEFKPDK200000020000000000	sp P52701 MSH6_HUMAN
2493	DKEEIFGSDADSEDDADSDEDR20002000002000000	sp Q8N7H5 PAF1_HUMAN
2494	SDSASSEPVGIYQGF EK220000000000000	sp Q05655 KPCD_HUMAN
2495	TMFAQVESDDEEAK20000000	sp Q9NW82 WDR70_HUMAN
2496	QSVVYGGKSTIRDR2200000	sp Q8WWY3 PRP31_HUMAN
2497	SSEGGVGVGPGGGDEPPTS PR2000	sp Q9Y4H2 IRS2_HUMAN
2498	IAAPELHKGDS DSEDEPTK20200000000	sp Q92733 PRCC_HUMAN
2499	VQDTSNTGLGEDI IHQLSK200000000000000000	sp Q8IWA0 WDR75_HUMAN
2500	ALSSLHGDDQDSEDEVLTIPEVK200000000000000	sp P11717 MPRI_HUMAN
2501	LKSVEDEMDSPGEEPFTGQGR2. 0000002E+20	sp Q12857 NFIA_HUMAN
2502	TPSSPLQSCTPSLS PR20000000000002000	sp Q96AP0 ACD_HUMAN
2503	RGSQKSTDSPGADAELPESAAR2020000000000000	sp Q9Y4K1 AIM1_HUMAN
2504	LESSEGEI IQTVDR200000000000	sp Q99590 SCAFB_HUMAN
2505	TGVTSTSDSEEGDDQEGEK200020000000000000	sp 075475 PSIP1_HUMAN
2506	TSPTVATQTGASATSTR200000000000000000	sp Q6P1L5 F117B_HUMAN
2507	SPPGSGASTPVGPWDQAVQR200000000000000000	sp Q8TE77 SSH3_HUMAN
2508	SNSQTLSDAPFTTCPEGPAR200000000000000000	sp Q8IZW8 TENS4_HUMAN
2509	ADAPDAGAQS DSELPSYHQNDVSLDR200000000000000000	sp Q92538 GBF1_HUMAN
2510	SPESDSQSPEFESQSPR200200000000002000	sp Q9H5H4 ZN768_HUMAN
2511	AEELSPAALSPSLEPIR20000200000000	sp Q12774 ARHG5_HUMAN
2512	LPSPTSPFSSLSQDQAATSK2020000000000000000	sp Q8WWI1 LM07_HUMAN
2513	SGSALLQSQSSTEDPKDEPAELKPDSEDLSSQSSASK2E+26	sp Q5VTR2 BRE1A_HUMAN
2514	METADAPSPAPSLGER100000002000200000	sp Q12873 CHD3_HUMAN
2515	EFVSSDESSGENK220022000000	sp Q08945 SSRP1_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
2516	KGGSYSQAASSDSAQSDVSLTACK20002000000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
2517	RDSGDNSAPSGQER200000000000000	sp Q8IXQ3 CI040_HUMAN
2518	FDSSLSSDDETK20022000000	sp P28715 ERCC5_HUMAN
2519	QEEESPVERPLK200000000	sp Q96SB4 SRPK1_HUMAN
2520	SPSPDGS PAATPEIR20000002000000000	sp 000499 BIN1_HUMAN
2521	ATNESEDEIPQLVPIGK200000000000000000	sp 076021 RL1D1_HUMAN
2522	SGSSQELDVKPSASPQER2020000000000200000	sp Q9UQ35 SRRM2_HUMAN
2523	SEDGVEGDLGETQSR200000000000000000	sp Q09666 AHNK_HUMAN
2524	AVSPPHLDGPPSPR2000000002000	sp P29590 PML_HUMAN
2525	GLLAQGLRPESPPPAGPLLNGAPAGESPQPK2E+21	sp Q96TA1 NIBL1_HUMAN

2526	FQETEFLSPPR20000	sp	P42166	LAP2A_HUMAN
2527	SQDQDSEVNELSR200	sp	Q86VM9	ZCH18_HUMAN
2528	SPSFGEDYYGPSR20200000000200	sp	P49761	CLK3_HUMAN
2529	SKSPNNEGDVHFSR200000000000	sp	Q8NAP3	ZBT38_HUMAN
2530	KKEPAITSQNSPEAR200000	sp	P23193	TCEA1_HUMAN
2531	EKTPELPEPSVK20000000000	sp	Q8IYB3	SRRM1_HUMAN
2532	AWLDEDSNLSPSPLR2000020000	sp	Q9C086	IN80B_HUMAN
2533	GRLTPSPDIIVLSDNEASSPR2000000002000000000	sp	Q8WXI9	P66B_HUMAN
2534	TASGSSVTSLDGTR2020000000	sp	Q92597	NDRG1_HUMAN
2535	ALDISLSSGEEDEGDEEDSTAGTTKQPSK2. 2E+23	sp	Q8IY81	SPB1_HUMAN
2536	MESHSEDEDLAGAVGGLGWSNR2. 02E+20	sp	Q96MH2	HEX12_HUMAN
2537	DLSPIDLNSAAK200000000000	sp	Q9P2B4	CT2NL_HUMAN
2538	KRESESEDETTPAAPQLIK2020000000000000000	sp	060832	DKC1_HUMAN
2539	SDSQASLTVPLSPQVVR2002000002000000	sp	Q8TEV9	SMCR8_HUMAN
2540	VQQEIDSDSPVAR200000000	sp	Q6T4R5	NHS_HUMAN
2541	QPTPPSEAAASK20000000000	sp	Q86UP2	KTN1_HUMAN
2542	GAAAADLLSSSPESQHGGTQSPGGGQPLLQPTK2E+24	sp	Q9H3S7	PTN23_HUMAN
2543	LDTGSPEPETR20000000	sp	Q9Y2F5	ICE1_HUMAN
2544	VISDSESDIGGSDVEFKPDTK202000020000000000	sp	P52701	MSH6_HUMAN
2545	DSGSQVEEDTSCSDSK20000000000000	sp	Q08499	PDE4D_HUMAN
2546	SRSPLDKDITYPPSASVVGASVGGHR2. 000000003E+25	sp	Q96T37	RBM15_HUMAN
2547	SVSNGTAKPATSENFDEDLK2. 00002E+20	sp	Q96K49	TM87B_HUMAN
2548	GLMAGGRPEGQYSEDEDTDDEYK1. 000000002E+22	sp	Q9NPQ8	RIC8A_HUMAN
2549	SGGLQTPECLSR20000000	sp	P85037	FO XK1_HUMAN
2550	KPSDSL SVASSR22000	sp	Q5T0W9	FA83B_HUMAN
2551	TSPADHGGSVGSESGSAVDSVAGEHSVSGR2. 20000002002E+31	sp	Q5T4S7	UBR4_HUMAN
2552	APTLAELDSSESEEETLHK220200000000	sp	Q08AE8	SPIR1_HUMAN
2553	LYADSDSGDSDKR20200000000	sp	043896	KIF1C_HUMAN
2554	ENSFGPSLEFR2000000	sp	Q9UQ35	SRRM2_HUMAN
2555	LPLSPPLVEDSAFEPSPR2000000000000000	sp	096007	MOC2B_HUMAN
2556	YSDGDKSDDLVDVSNEDPATPR3. 0000002E+24	sp	Q04726	TLE3_HUMAN
2557	LRSPEALVQGR20000000000	sp	Q9UHB6	LIMA1_HUMAN
2558	TDGSISGDRQPVTVADYISR2. 00002E+20	sp	P51116	FXR2_HUMAN
2559	SSGDPEQIKEDSLSEESADAR20200200000	sp	Q9BVS4	RIOK2_HUMAN
2560	RPTETNPVTSNSDEECNETVK2. 000000002E+19	sp	P46100	ATRX_HUMAN
2561	QGSITSPQANEQSVTPQR200000000020000	sp	Q9UQ35	SRRM2_HUMAN
2562	KPPAPSPVQSQSPSTNWSPAVPVK2. 000002020002E+19	sp	Q8IYB3	SRRM1_HUMAN
2563	QSPASPPPLGGAPVR2002000000000000	sp	Q14160	SCRIB_HUMAN
2564	SVA AEGALLPQTPPSR2002000	sp	Q86X27	RGPS2_HUMAN
2565	AEAPAGPALGLPSPEAESGVDR20000000000	sp	Q53F19	CQ085_HUMAN
2566	EHYPVSSPSSPPAQP GGVS R2000200000000000000	sp	075179	ANR17_HUMAN
2567	LSPLPPGSPGSAAR20000020000000	sp	Q6ZN18	AEBP2_HUMAN
2568	ATLLNVPDLSDSIHSANASER20002000	sp	Q07157	ZO1_HUMAN
2569	VSGSPEQAVEENLSSYSLDR2000000000000000000	sp	Q9Y6A5	TACC3_HUMAN
2570	TDDTSGEDNDEKEAVASK2000200000000000000	sp	Q9Y618	NCOR2_HUMAN
2571	QEQINTEPLEDTVLSPTK200	sp	015446	RPA34_HUMAN
2572	SFSLASSNSPISQR20000000000000000	sp	Q9BXB4	OSB11_HUMAN
2573	KHGLAHDEMKS PR2000	sp	Q9Y2W1	TR150_HUMAN
2574	EGEETEGSEEDDKENDKTEEMPNDVLENK2. 002E+26	sp	P49321	NASP_HUMAN
2575	SLSPGGAALGYR20200000000000	sp	Q96T37	RBM15_HUMAN
2576	SRSVSPCSNVESR20202000000000	sp	Q9UQ35	SRRM2_HUMAN
2577	SESTD SLGGLSPSEVTAIQCK2000000000000000000	sp	060318	GANP_HUMAN
2578	DSHSSEDEASSQTDLSQTISK2. 2000002E+19	sp	Q5JTV8	TOIP1_HUMAN
2579	SYSSPDITQAIQEEEK2000000000000000	sp	P40818	UBP8_HUMAN

2580	IVEYSPPSAPR20000000	sp	095696	BRD1_HUMAN
2581	KPSDSLVSASSR20000	sp	Q5T0W9	FA83B_HUMAN
2582	AGLESGAEPGDGSDTTKK2000000	sp	060832	DKC1_HUMAN
2583	GSPQSAEELSSGLLASAAASSR2E+19	sp	Q12948	FOXC1_HUMAN
2584	AESSDSGAEESEEEEAQEEVK2000002000000000000	sp	Q5T8D3	ACBD5_HUMAN
2585	VTETEDDSDSDDDEDDVHVTIGDIK2. 000202E+24	sp	Q6UN15	FIP1_HUMAN
2586	SSSVLSLEGSEK20020000000	sp	A1L390	PKHG3_HUMAN
2587	SLDSESDDSK202000000000	sp	Q13185	CBX3_HUMAN
2588	ELKSDASSDQGPASPEK200220000020000	sp	Q6BDS2	URFB1_HUMAN
2589	TSSTCSNESLSVGGTSVTPR20020000000000000	sp	060343	TBCD4_HUMAN
2590	SPTGPSNSFLANMGTVAHK2000000000000000000	sp	Q96I25	SPF45_HUMAN
2591	SSTPLPTISSAENTR2200000000000000000	sp	P42167	LAP2B_HUMAN
2592	TCNSPQNSTDSVSDIVPDSPPFGALGSDTR2. 0202E+22	sp	P42166	LAP2A_HUMAN
2593	TVANLLSGKSPR2000	sp	Q14694	UBP10_HUMAN
2594	QHLYSSEDEPDDNPDVLSR3002000000000000000	sp	Q13428	TCOF_HUMAN
2595	RLSSASTGKPLSVEDDFEK2200000000000000000	sp	Q8NFC6	BD1L1_HUMAN
2596	LFEDDSNEK20000	sp	075152	ZC11A_HUMAN
2597	RLSYNTASNK200000000	sp	043719	HTSF1_HUMAN
2598	KPSTDEQTSSAEEDVPTCGYLNLSNSR2. 22E+21	sp	P49207	RL34_HUMAN
2599	SGSIVELIAGGGSSCSPVLSR2E+19	sp	Q8N556	AFAP1_HUMAN
2600	VSSPLPSPSAMTDAANSQAAAK2. 0002E+20	sp	P42858	HD_HUMAN
2601	TLDVSTDEEDKIHSSSEK2200000000000000000	sp	Q8WXI9	P66B_HUMAN
2602	LASPSGSTSSGLEVVAPEGTSAPGGGPGTLDDSATICR2. 0202E+36	sp	P16383	GCFC2_HUMAN
2603	GSVSPSTPSPKPTGIFQTSANSSFEVK2. 002202E+27	sp	Q13263	TIF1B_HUMAN
2604	MHNLHGTKPPPSEGSDEEEEEDEEDEEER1. 00000000000002E+30	sp	015027	SC16A_HUMAN
2605	GSPEEELPLPAFEK200000000000000	sp	Q9BQ67	GRWD1_HUMAN
2606	AGSCCLSPGGR200000	sp	Q5JTD0	TJAP1_HUMAN
2607	KGSDDAPYSPTAR200000200000	sp	Q969V6	MKL1_HUMAN
2608	ASPSQPSSQPLQIHR2000000000000000000	sp	075151	PHF2_HUMAN
2609	VLSSTSEDEPGVVK202200000000000	sp	Q9HC35	EMAL4_HUMAN
2610	SMVSPVSPSTGTISVPNSCPASPR2. 002E+24	sp	Q8NI08	NCOA7_HUMAN
2611	NYQQNYQNSSESGEKNEGSESAPEGQAQR3. 00002E+24	sp	P85037	FO XK1_HUMAN
2612	EVSSRPSTPGLSVVSGISATSEDI PNK2E+21	sp	P67809	YBOX1_HUMAN
2613	RNSSPVPSPASVPGQR202000000000000	sp	Q14C86	GAPD1_HUMAN
2614	ISPLSSPCSSPLQGTPASSLVSK2E+22	sp	Q9H1B7	I2BPL_HUMAN
2615	KSPSGPVKSPPLSPVGTTPVK2. 0000002E+20	sp	Q14432	PDE3A_HUMAN
2616	QDDSPSGASYGQDYDLSPSR200000000000020000	sp	Q9BVC5	ASHWN_HUMAN
2617	ASLEAAIADAEQR200000000000000	sp	Q9NYV4	CDK12_HUMAN
2618	EGDPVSLSTPLETEFGSPSELSR2000	sp	P05787	K2C8_HUMAN
2619	RVESEESGDEEGKK200200000000	sp	Q9BUR4	WAP53_HUMAN
2620	GNKSPSPDGSPAATPEIR20200000000000000	sp	Q15435	PP1R7_HUMAN
2621	VNSNGKESPGSSEFFQEA VSHGK20020000000000000	sp	000499	BIN1_HUMAN
2622	RLSSTSLASGHSVR220000000000000	sp	Q8N108	MIER1_HUMAN
2623	SLDMSIIAEVK200000000000000	sp	Q9BZL6	KPCD2_HUMAN
2624	TGSNISGASSDISLDEQYK20000000000000000	sp	P05787	K2C8_HUMAN
2625	EPAITSQNSPEAR200000	sp	P22059	OSBP1_HUMAN
2626	GGHSSVSTESESSSFHSS20000000000000000	sp	P23193	TCEA1_HUMAN
2627	KLSSSEPYEDEFNDDQSIK2. 22E+19	sp	P61073	CXCR4_HUMAN
2628	GPKPEPPGSGSPAPPR2000000	sp	Q9H4L7	SMRCD_HUMAN
2629	TLEEVMAEEDEGTPRGSPA2000	sp	Q9UKJ3	GPTC8_HUMAN
2630	SSGTASSVAFTPLQGLEIVNPQAAEK2. 0000020002E+26	sp	A6NKF1	SAC31_HUMAN
2631	LNHVAAGLVSPSLKSDTSSK200002000000	sp	Q8WWY3	PRP31_HUMAN
2632	LIRPEYAWIVQPVSGAVYDRPGASPK2000	sp	Q05519	SRS11_HUMAN
		sp	Q12769	NU160_HUMAN

2633	SPSSDSWTCADTSTER2000000000000000	sp Q8N6T3 ARFG1_HUMAN
2634	SQSSGSSATHPIISVPGAR2000000000000000	sp Q9Y4H2 IRS2_HUMAN
2635	LEKEEEEGISQESSEEEQ2002200000	sp P17096 HMGA1_HUMAN
2636	HSPIAPSSPSPQVLAQK2000000000000000	sp Q9NQS7 INCE_HUMAN
2637	IQPLEPDSPTGLSENPTATEK2000000000000000	sp Q9ULJ3 ZBT21_HUMAN
2638	SLHSAHSLASR200000000000	sp O15027 SC16A_HUMAN
2639	RPDPDSDEDEDYER2000000000	sp Q96I25 SPF45_HUMAN
2640	SATRPSPSPERSSTGPEPPAPTPLLAER2.020002E+23	sp Q9UQ35 SRRM2_HUMAN
2641	SLSSGSNFCSEK200000000000	sp O95243 MBD4_HUMAN
2642	NTPSQHSHSIQHSPER20000	sp Q9NYF8 BCLF1_HUMAN
2643	RPACDPEPGESGSSSDEGCTVVRPEK2022200000000000	sp O15541 R113A_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
2644	GGSYSQAASSDSAQGSVSLTACK2E+20	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
2645	CSGPGLSPGMVR2000000	sp P21333 FLNA_HUMAN
2646	EQGTESRSSTPLPTISSAENTR2200000000000000	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
2647	ESDQTLAALLSPK2000	sp P18583 SON_HUMAN
2648	KSSISSISGR200200000	sp Q86SQ0 PHLB2_HUMAN
2649	DGEQSPNVSLMQR2000000000	sp Q58WW2 DCAF6_HUMAN
2650	GYSPYSVSGSGSTAGSR200	sp Q15149 PLEC_HUMAN
2651	KAPAGQEEPPTPSSPLSAEQDR2000200000000000	sp P13051 UNG_HUMAN
2652	QLPALDGSLMGPEPPAQEEAPVSPHK2000000000000000	sp Q8N9T8 KRI1_HUMAN
2653	SPSPPTQHTGQPPGQPSAPQLSAPR2.002E+21	sp Q68EM7 RHG17_HUMAN
2654	GGRSPDEVTLTSIVPTR2000000000000000	sp Q9UKS6 PACN3_HUMAN
2655	VIGQDHFSESEEEAPAEASSGALR2002000000000000000	sp P23497 SP100_HUMAN
2656	RLSPSSAASSYSFSDLNSTR20000000000000000000	sp P50402 EMD_HUMAN
2657	TSAQHALTSVSR200	sp P61073 CXCR4_HUMAN
2658	RQEIESDSESDGELQDR2020200000000	sp Q9BYW2 SETD2_HUMAN
2659	ELSESVQQQSTPVPLISPK2000	sp Q9ULU4 PKCB1_HUMAN
2660	KQSVFSAPSLSAGASAAEPLDR2000000000000000	sp Q14573 ITPR3_HUMAN
2661	LFSQQQDVSNK2000000000	sp P55196 AFAD_HUMAN
2662	SQTTTERDSDTDVEEEELPVENR2020000000000000	sp Q14676 MDC1_HUMAN
2663	TAANKSPCETISSPSSTLESK2000000000000000	sp Q9COD5 TANC1_HUMAN
2664	SPGSTPTTPTSSQAPQK2200000000000000	sp P35658 NU214_HUMAN
2665	NEIIQSPISQVPSVEK200000000000	sp Q7Z3T8 ZFY16_HUMAN
2666	QPATPTAAESSEGEEGEGDDGGETESR2.0002E+22	sp Q96S55 WRIP1_HUMAN
2667	EISDDEAEEK2000000000	sp P08238 HS90B_HUMAN
		sp Q58FF7 H90B3_HUMAN
2668	AAPPPPPPPPLESSPR2000	sp Q5T4S7 UBR4_HUMAN
2669	NSLTGEEQLAR200000000000	sp Q9BX95 SGPP1_HUMAN
2670	VSPAGGLDDKPK200000000000	sp Q8WY36 BBX_HUMAN
2671	ESAAPASPAPSPAPSTPAPPQK200000002000000000	sp Q3KQU3 MA7D1_HUMAN
2672	NVSTGSAENEEK2002000000	sp O60271 JIP4_HUMAN
2673	SRTGSESSQTGTSTTSSR2020000000000000	sp P23588 IF4B_HUMAN
2674	ENPPVEDSDEDDKR2000000000	sp Q8TCJ2 STT3B_HUMAN
2675	TQTPPVSPAPQPTTEER2000000000	sp Q14247 SRC8_HUMAN
2676	SPPTMLLPASPAK200000000020000	sp Q9UGU5 HMGX4_HUMAN
2677	FSLSPSLGPQASR202000000000	sp Q9NYF3 FA53C_HUMAN
2678	ESEDELEEANGNNPIDIEVDQNK2E+22	sp Q9UKL0 RCOR1_HUMAN
2679	NNTAAETEDDESDEDR200002000000	sp Q6KC79 NIPBL_HUMAN



2680	ADEPSSEESDLEIDK20000000	sp P50502 F10A1_HUMAN
		sp Q8IZP2 ST134_HUMAN
2681	TGSTSSKEDDYESDAATIVQK300000000000	sp Q5T4S7 UBR4_HUMAN
2682	LYSSEESRPYTNK2000	sp Q14004 CDK13_HUMAN
2683	STLESEKPGSPEAAETSPPSNIIDHCEK2E+19	sp Q96T23 RSF1_HUMAN
2684	GPVSPSVSFQPLAR200000000000	sp Q7Z434 MAVS_HUMAN
2685	AHSSPASLQLGAVSPGTLTPTGVVSGPAATPTAQHLR2.0000002E+31	sp P46937 YAP1_HUMAN
2686	SLSNSNPDISGTPTSPDDEV2.0200000000002E+21	sp Q96N67 DOCK7_HUMAN
2687	VISDSESDIGGSDVEFKPDTK2.000000002E+19	sp P52701 MSH6_HUMAN
2688	HTDDEMTGYVATR100300000	sp Q16539 MK14_HUMAN
2689	QEKPAEKPAETPVATSPTATDSTSGDSSR2200000000000000	sp P54727 RD23B_HUMAN
2690	GYDSEDDTWEPEQHLVNCEEYIHFNR2E+24	sp Q9Y232 CDYL1_HUMAN
		sp P42167 LAP2B_HUMAN
2691	GPPDFSSDEEREPTPVLGSGAAAAGR2.0000002E+20	sp P42166 LAP2A_HUMAN
		sp Q15311 RBP1_HUMAN
2692	TPSSEEISPTKFPGLYR2000020000000000	sp Q6PKG0 LARP1_HUMAN
2693	TASISSPSEGTPTVGSYGCTPQSLPK2.2E+23	sp Q9H6F5 CCD86_HUMAN
2694	QQDLHLESPQR20000	sp Q8TF01 PNISR_HUMAN
2695	QLAQSSALASLTGLGGLGGYSGDSEDER200200000	sp P46013 KI67_HUMAN
2696	SPQPDPVDTFASTK2000000000000000	sp Q92797 SYMPK_HUMAN
2697	SPQTLAPVGEDAMKTPSPAEDAR2.0000000000002E+24	sp Q8IWX8 CHERP_HUMAN
2698	SRSPTPPSSAGLGSNSAPPIDSR2.02020022E+24	sp Q6PL18 ATAD2_HUMAN
2699	IESDTEETQDTSVDHNETGNTGESSVEENEK2E+29	sp Q8NC51 PAIRB_HUMAN
2700	FDQLFDDESDPFEVLK2000000000	sp O43294 TGF11_HUMAN
2701	SPKPAAPAPPFSSSSGVLGTGLCELDR2E+28	sp Q9HAS0 NJMU_HUMAN
2702	ELESSEEGGSAER200000000000	sp O75376 NCOR1_HUMAN
2703	HEAPSSPISGQPCGDDQNASPSK2000000000000020000	sp Q9Y608 LRRF2_HUMAN
2704	RGSGDTSSLIDPDTLSSEL2000000000000000	sp P19338 NUCL_HUMAN
2705	AAAAAPASEDEDEDDEDDEDDEDDDEEDDSEEEAMETTPAK2E+34	sp P62995 TRA2B_HUMAN
2706	RHSHSHSPMSTR202000000000	sp Q5T0W9 FA83B_HUMAN
2707	KPSDSLVAASSR20022000	sp Q13263 TIF1B_HUMAN
2708	LASPSGSTSSGLEVVAPEGTSAPGGGPGTLDDSATICR2.2E+31	sp O15027 SC16A_HUMAN
2709	SLHSAHSLASR200200000000	sp O94806 KPCD3_HUMAN
		sp Q15139 KPCD1_HUMAN
2710	TISPSTSNLIPLMR20002000000000	sp Q14157 UBP2L_HUMAN
		sp Q15154 PCM1_HUMAN
2711	STSAPQMSPGSSDNQSSSPQPAQQK2E+24	sp P27824 CALX_HUMAN
2712	KDEETESEYDSEHENSEPVTNIR2.0020002E+20	sp Q9HCD6 TANC2_HUMAN
2713	QKSDAEEDGGTVSQEEEDRKPK2.000000002E+20	sp Q9BQE9 BCL7B_HUMAN
2714	SCDELSPVSPTQGGYPSEPTR200200000000000000	sp Q9Y580 RBM7_HUMAN
2715	VDSSTNSSPSPQSESLSPAHTSDFR200020000000000000	sp Q9Y2X3 NOP58_HUMAN
2716	SFSSPENFQR2000000000	sp Q9Y4B6 VPRBP_HUMAN
2717	HIKEEPLSEEPCTSTAIASPEK200000000000000000	sp P49023 PAXI_HUMAN
2718	QKSDHGAYSQSPAIK2000000000000000	sp Q14160 SCRIB_HUMAN
2719	SSPGGQDEGGFMAQK2000000000000000	sp Q6PJG2 EMSA1_HUMAN
2720	NSLESISSIDR20000020000	sp O15014 ZN609_HUMAN
2721	SHE SNAPGSAGGQASEKPR200000000000	sp Q9NYF8 BCLF1_HUMAN
2722	TNSMGSATGPLPGTK2000000000000000	sp Q14152 EIF3A_HUMAN
2723	YSPSQNSPIHHIPSR2000000000000000	sp Q9UBC2 EP15R_HUMAN
2724	LGSSLSR20000	sp Q3B726 RPA43_HUMAN
2725	STPSHGVSSSLNSTGSLSPK2000	sp O60504 VINEX_HUMAN
2726	HQEVQDQDPVFGSDSSGYQSDHKKK220302000000	sp P42166 LAP2A_HUMAN
2727	LCDDGPQLPTSPR2000	sp P42167 LAP2B_HUMAN
		sp Q9NZL4 HPBP1_HUMAN
2728	EQGTESRSSTPLPTISSAENTR222000000000000000	
2729	LLQTCFSSPADDSMDR2000000000	

		sp	P05534	1A24_HUMAN
		sp	P30447	1A23_HUMAN
2730	KGGSYSQAASSDSAQGSVSLTACK2002000000	sp	P01891	1A68_HUMAN
		sp	P01892	1A02_HUMAN
		sp	Q09160	1A80_HUMAN
		sp	P10316	1A69_HUMAN
2731	SLGPSLATDK20000000000	sp	Q9H3N1	TMX1_HUMAN
2732	VGDSTPVSEKPVSAAVDANASESP2E+20	sp	Q9H8Y8	GORS2_HUMAN
2733	TPELPEPSVK20000000000	sp	Q8IYB3	SRRM1_HUMAN
2734	SNSFNPLGNR20000000000	sp	O95835	LATS1_HUMAN
2735	GRLTSPDIIVLSDNEASSPR2020000002000000000	sp	Q8WXI9	P66B_HUMAN
2736	IPCKSPPELTDATSTK2000000000000000	sp	P46013	KI67_HUMAN
2737	AASPESASTPESLQAR2000000000000000	sp	Q96BT3	CENPT_HUMAN
2738	TETPPPLASLVSK2000000000000000	sp	Q3MHD2	LSM12_HUMAN
2739	VSPASPAGSPSADFAVHGESLGR2. 002E+23	sp	Q96C92	SDCG3_HUMAN
2740	QPATPTAAESSEGEEGDDGGETESR2200000000000000000	sp	Q96S55	WRIP1_HUMAN
2741	SPSDLHISPLAK20000000000	sp	O95785	WIZ_HUMAN
2742	TPSSSPITPPASETK20002000000000	sp	Q9NZJ0	DTL_HUMAN
2743	ESSIIAPAPAEDVDTPPR20000	sp	Q15910	EZH2_HUMAN
2744	TGQAGSLSGSPKPFSPQLSAPITTK2. 02E+20	sp	Q9ULU4	PKCB1_HUMAN
2745	HSPIAPSSPSPQVLAQK2000020000000000000	sp	Q9NQS7	INCE_HUMAN
2746	LASPGSTSSGLEVVAPEGTSAPGGPGTLDDSATICR2. 00000000022E+29	sp	Q13263	TIF1B_HUMAN
2747	NSVERPAEPVAGAATPSLVEQQK2E+22	sp	Q9BTC0	DID01_HUMAN
2748	DYLLSESEDEGNDGERK2020000000000000	sp	Q9BVJ6	UT14A_HUMAN
2749	AASSSPGSPVASSPSR20000	sp	Q92538	GBF1_HUMAN
2750	SGKNSQEDSEDEKDKV2000200000000000	sp	Q9H1E3	NUCKS_HUMAN
2751	SCDPGEDCASCQQDEIDVVPESPLSDVGSSEVGTGPK2002000000000000000	sp	Q86W56	PARG_HUMAN
2752	SPHSPPIATPPSQSSSCSDAPMLSTVHSAK2. 020002002E+28	sp	Q9NYF8	BCLF1_HUMAN
2753	SFSADNFIGIQR200000000000	sp	Q8N7R7	CCYL1_HUMAN
2754	RDSFDRGPSLNPVLDYDHGSR2000000000000000	sp	P43243	MATR3_HUMAN
2755	QQPQPSQTESPQEAQIIQAK2000000000000	sp	O96020	CCNE2_HUMAN
2756	SPGSTPTTSSQAPQK2000000000000	sp	P35658	NU214_HUMAN
2757	DTSATSQSVNGSPQAEQPSLETSK2. 02E+20	sp	Q86WB0	NIPA_HUMAN
2758	DSGSEDFLMEDDDSDYGSSK2. 00000000002E+19	sp	Q9H1E3	NUCKS_HUMAN
2759	LASPGSTSSGLEVVAPEGTSAPGGPGTLDDSATICR2. 20000000002E+30	sp	Q13263	TIF1B_HUMAN
2760	SQDADSPGSSGAPENLTFK2002000000000000	sp	P55196	AFAD_HUMAN
2761	DSGSDQLDAGVR2020000000000000	sp	Q86VM9	ZCH18_HUMAN
2762	SANGSESDGEENIGWSTVNLDEEK2. 0000202E+25	sp	O43290	SNUT1_HUMAN
2763	KVHPYEDSDSEDEVDWQDTR2022000000000000	sp	O14730	RIOK3_HUMAN
2764	QILIIACSPVSSVR20000000	sp	Q5QJE6	TDIF2_HUMAN
2765	SPNELVDDLK2000000000000	sp	Q9UNZ2	NSF1C_HUMAN
2766	SPGFASPK2000000000	sp	P49790	NU153_HUMAN
2767	LGSTGSPNSEAESVPEVNPVPPK2. 2E+26	sp	Q8IZ21	PHAR4_HUMAN
2768	AGMSSNQSISPVLDVPR1020000220000000000	sp	Q9UQ35	SRRM2_HUMAN
2769	GTSRPPPEGGLGYSQLGDDDLK2E+20	sp	Q9UQ88	CD11A_HUMAN
		sp	P21127	CD11B_HUMAN
2770	KWSDSSKQDDSPSGASYGQDYDLSPSR2. 022E+25	sp	Q9NYV4	CDK12_HUMAN
2771	GLRDSHSSEDEASSQTDLSQTISK2. 20000022E+19	sp	Q5JTV8	TOIP1_HUMAN
2772	GSPTSAYPER2000000000	sp	Q6ZRV2	FA83H_HUMAN
2773	SALSPSKSPAK20000	sp	P49792	RBP2_HUMAN
		sp	Q7Z3J3	RGPD4_HUMAN

2774	VGSLDNVGHLPAGGAVK2000000000000000	sp P27816 MAP4_HUMAN
2775	VPPAPVPCPPSPGSAVPSSPK20000000000000	sp O95817 BAG3_HUMAN
2776	RPPSPDIVLSDNEQPSSPR20000000000002000	sp Q86YP4 P66A_HUMAN
2777	GPAATPGSPDPGPGPWGEEHLAER2E+20	sp Q5VV17 OTUD1_HUMAN
2778	AGPTLTDENDDLGLPPSPGDTSYQQDQVDDFHEAR2E+33	sp Q9NQZ2 SAS10_HUMAN
2779	QGSITSPQANEQSVTPQR220000000020000	sp Q9UQ35 SRRM2_HUMAN
2780	APEPEEEIEEVEESEDEGEDPALDLSQAIAFQQA2E+23	sp Q8TBF4 ZCRB1_HUMAN
2781	ERDHSPTPSVFNSEER20000000000000	sp Q6UN15 FIP1_HUMAN
2782	DLELPLSPLLGGGPER20000000000000	sp P19419 ELK1_HUMAN
2783	ADAPDAGAQSDSELPSYHQNDVSLDR2000000000000000	sp Q92538 GBF1_HUMAN
2784	KLSGDQPAAR2000000000	sp Q13428 TCOF_HUMAN
2785	EHYPVSSPSSPPAQGGVSR200000200000000000	sp O75179 ANR17_HUMAN
2786	SASASHQADIK200000000000	sp Q14739 LBR_HUMAN
2787	SYLEGSSDNQLKDESETPVDDR3.00002E+21	sp Q9UHI6 DDX20_HUMAN
2788	TGVTSTSDSEEEGDDQEGEK222000000000000000	sp O75475 PSIP1_HUMAN
2789	AELGMDSTSQSPPIKR2000000	sp Q9NR19 ACSA_HUMAN
2790	VFPEPTESGDEGEELGLPLLSTR200000000000000000	sp Q9BUR4 WAP53_HUMAN
2791	RLSPSASPPR200020000	sp Q8IYB3 SRRM1_HUMAN
2792	EGETQGVAFEHESPADFQNSQSPVQDQDK2E+26	sp P78332 RBM6_HUMAN
2793	KYESDEDSLSSGR200020000000	sp O00418 EF2K_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
2794	GGSYSQAASSDSAQGSVSLTACK20002002000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
2795	TFQQIQEEEDDDYPGSYSPQDPSAGPLLTEELIK200000000000000000	sp Q9NQG5 RPR1B_HUMAN
2796	DFLSGPDYDHTVGSDEDEEVEGEP2020000000000000	sp Q6ZW31 SYDE1_HUMAN
2797	QGGSQPSSFSQSQVTPQDQEK200000000000000000	sp Q9H0L4 CSTFT_HUMAN
2798	SEDEDSLEEAGSPAPGPCPR2000000000000000	sp Q8TBB5 KLDC4_HUMAN
2799	RNSSSPVSPASVPGQR22002000000000	sp Q9H1B7 I2BPL_HUMAN
2800	STSAQMSPGSSDNQSSSPQAQQK2E+25	sp Q14157 UBP2L_HUMAN
2801	CSSSSGGSSGDEDEGLDGLDAPGGGK2.22E+24	sp Q9P258 RCC2_HUMAN
2802	TPASTPVSGTPQASPMIER20002010000	sp Q9UHR4 BI2L1_HUMAN
2803	YSPSQNSPIHHIPSR2000000000000000	sp Q9NYF8 BCLF1_HUMAN
2804	DYTGCTSESLSPVK20000	sp O95297 MPZL1_HUMAN
2805	DVDDGSGSPHSPHLSSK200000000000	sp Q9H4A3 WNK1_HUMAN
2806	KDSEGYSESPDLEFEYADTDK20000000000000	sp Q5VSL9 STRP1_HUMAN
2807	YGMGTSVER10020000	sp P08559 ODPA_HUMAN
2808	GGNVFAALIQQSEEEEEEEK2000000000	sp Q8NE71 ABCF1_HUMAN
2809	AAYGDLSEEEEEENEPESLGVVYK220000000000000000	sp O15541 R113A_HUMAN
2810	HGSFHEDEDPIGSPR20000000002000	sp Q96T58 MINT_HUMAN
2811	EKTPSPKEEDEEPESPPEK202000000000200000	sp Q9H1E3 NUCKS_HUMAN
2812	KAENAEGQTPAIGPDGEPLDQMSDLPVK2002000000	sp P51532 SMCA4_HUMAN
2813	RDPSSNDINGMEPTPSTVSTPSADLLGLR22020000000000	sp O95782 AP2A1_HUMAN
2814	SDDSFQSLHGSLPLDSFSK2.002E+19	sp Q7Z5J4 RAI1_HUMAN
2815	VSPSGMEEAGCSR200000000000	sp Q96RL1 UIMC1_HUMAN
2816	VLDTSSLTQSAPASPTNK200000	sp Q8N122 RPTOR_HUMAN
2817	EHYPVSSPSSPPAQGGVSR2202000000000000	sp O75179 ANR17_HUMAN
2818	SPISDNSGCDAPGNSNPSLSPSSAESEK2.00200000002E+26	sp Q99590 SCAFB_HUMAN
2819	DFLSGPDYDHTVGSDEDEEVEGEP2000200000000000	sp Q6ZW31 SYDE1_HUMAN
2820	HLSESSGKPLSTK200000000000	sp Q8IVT2 MISP_HUMAN
2821	VSPASPAGSPSADFAVHGESLDR2.0002E+20	sp Q96C92 SDCG3_HUMAN

2822	GYSPYSVSGSGSTAGSR200200	sp Q15149 PLEC_HUMAN
2823	VMTIPYQMPASSPVICAGGQDR1.00000010002E+22	sp Q15365 PCBP1_HUMAN
2824	TSSVSNPQDSVGSPCR200000	sp P49023 PAXI_HUMAN
2825	KPSTSDSDSNFEK220020000000	sp P11388 TOP2A_HUMAN
2826	KPPAPSPVQSQSPSTNWSPAVPVK2.000000000002E+19	sp Q8IYB3 SRRM1_HUMAN
2827	FAEDEEKSENSEEDGDITDK20022000000000	sp Q5C9Z4 NOM1_HUMAN
2828	ALEEGDGSVSGSSPR2000	sp Q8N3X1 FNBP4_HUMAN
2829	TQMAEVLSPR2000	sp P11388 TOP2A_HUMAN
2830	ATSPLVSLYPALECR2000000000000000	sp Q96G74 OTUD5_HUMAN
2831	RTDALTSPPGR220000	sp P49736 MCM2_HUMAN
2832	ESEVCPVPTNSPSPPLPPPPPLPK200020000000000000	sp Q9ULJ3 ZBT21_HUMAN
2833	TPAQSGAWDPNNPNTPSR20000000000000000000	sp 000267 SPT5H_HUMAN
2834	EVSDDEAEKEDK200000000000	sp P07900 HS90A_HUMAN
2835	RPTETNPVTSNSDEECNETVK20200000000000	sp P46100 ATRX_HUMAN
2836	KLTSDEEGEPSGK220000000000	sp Q8WVC0 LEO1_HUMAN
2837	GKGNIESSEDEDDTADLFPEESGFHAWR2.22E+24	sp Q9H501 ESF1_HUMAN
2838	KPVTVSPPTPTSPTEGEAS2020000000000000	sp Q9Y6G9 DC1L1_HUMAN
2839	AVSLDSPVSVGSPPVK200000000000000000	sp Q9Y6Q9 NCOA3_HUMAN
2840	QGSITSPQANEQSVTPQR20200000000020000	sp Q9UQ35 SRRM2_HUMAN
2841	TSLMSAESPTPR201200200000	sp P49810 PSN2_HUMAN
2842	ASALLFSSDEEDQWNIPASQTHLASDSR2.2E+22	sp Q641Q2 FA21A_HUMAN
2843	DGDSYDPYDFSDEEEMPQVHTPK300000000000000000	sp Q5SNT6 FA21B_HUMAN
2844	ELSTSPCNPIVR20200000000	sp P25205 MCM3_HUMAN
2845	SANGGSESDGEENIGWSTVNLDEEK2.02E+20	sp Q9H2Y7 ZN106_HUMAN
2846	SSSPTQYGLTK2000000000	sp 043290 SNUT1_HUMAN
2847	LPALSHSEGEDEDEEEDGK202000000000000000	sp Q9NYL2 MLTK_HUMAN
2848	SSMSGLHLVK20000000	sp P55201 BRPF1_HUMAN
2849	GSPSPASVSSGR200000000000	sp Q13085 ACACA_HUMAN
2850	MSGGSTMSSGGGNTNNSNSK200000000000000000	sp P10398 ARAF_HUMAN
2851	SLGGAVGSVASGAR2000000000000000	sp Q86U70 LDB1_HUMAN
2852	IFDEVMGCFSDSPQSPFPPEAGHTSLYDEDKVR2.000002E+26	sp Q8NHG8 ZNR2_HUMAN
2853	VQSPEPPAPER2000000000	sp Q13085 ACACA_HUMAN
2854	SPPTQVAISSDSAR2000000000000000	sp Q14160 SCRIB_HUMAN
2855	NTPSQHSHSIQHSPER2000000000020000	sp Q6ZUT6 COO52_HUMAN
2856	VSNLQVSPK2000	sp Q9NYF8 BCLF1_HUMAN
2857	HSPSPPPPTPTESR20200000000000	sp 000443 P3C2A_HUMAN
2858	LAAQLGAPTSPIPDSAIVNTR20000000000000	sp Q92922 SMRC1_HUMAN
2859	MSGEPITVESIR20000000000000	sp Q9H8Y5 ANKZ1_HUMAN
2860	EGSPIPHDPEFGSK200000000000	sp Q5VT52 RPRD2_HUMAN
2861	SPSKPTLAYPESNSR20000000000000	sp P85037 FOXK1_HUMAN
2862	YTPTSPSYSPSSPEYTPSPK220000000000	sp Q86XR8 CEP57_HUMAN
2863	AGMSSNQSISSPVLDVPR102002002000000000	sp P24928 RPB1_HUMAN
2864	SHVSSEPYEPIPPQVPVVEHK200000000000	sp Q9UQ35 SRRM2_HUMAN
2865	EAENQGLDISSPGMSGHR2000000000	sp 075376 NCOR1_HUMAN
2866	DSGDENEPIQER200000000000	sp Q15545 TAF7_HUMAN
2867	NVALLSPLYHSPAR20000	sp Q8WX93 PALLD_HUMAN
2868	NEVNGTSEDIKSEGDTQSN200000200	sp Q15717 ELAV1_HUMAN
2869	ADEPSSEESDLEIDK220020000000	sp 095232 LC7L3_HUMAN
2870	RYPSSISSSPQK200000	sp P50502 F10A1_HUMAN
2871	KKSEPDELLEDFLNSSQK200000000000000000	sp Q8IZP2 ST134_HUMAN
2872	GLNSQSSDHLNKR2000000000	sp Q14157 UBP2L_HUMAN
		sp Q8TBA6 GOGA5_HUMAN
		sp A8CG34 P121C_HUMAN
		sp Q96HA1 P121A_HUMAN

2873	KLEGNPQGSNQGK20000000000	sp P61006 RAB8A_HUMAN
2874	SPGSTPTTSSQAPQK2020000000000	sp P35658 NU214_HUMAN
2875	SASPHVDLCLVSPCEFEHR2.02E+20	sp Q66K74 MAP1S_HUMAN
2876	SPISDNSGCDAPGNSNPSLSPSSAESEK2E+29	sp Q99590 SCAFB_HUMAN
2877	SIQFVDWCPTGFK20000000000000	sp P68363 TBA1B_HUMAN
		sp P68366 TBA4A_HUMAN
		sp Q9UQ88 CD11A_HUMAN
2878	DLLSDLQDISDSER202000	sp P21127 CD11B_HUMAN
		sp Q9H3Z4 DNJC5_HUMAN
2879	SLSTSGESLYHVLGLDK20000000000	sp Q96JM3 CHAP1_HUMAN
2880	DNQESSDAELSSSEYIK2000000000000	sp P49790 NU153_HUMAN
2881	IPSIVSSPLNSPLDR20000000000	sp Q8IYB3 SRRM1_HUMAN
2882	APQTSSSPPVVR202000000	sp Q9UQ35 SRRM2_HUMAN
2883	SRTPPSAPSQSR200000000000	sp P29317 EPHA2_HUMAN
2884	LPSTSGSEGVPR200000000	sp P17812 PYRG1_HUMAN
2885	SGSSPDSEITELKFPSINH2.022E+21	sp P80723 BASP1_HUMAN
2886	AQGPAAAEKPKVEAPAANSQTVTVK2000000000	sp P46013 KI67_HUMAN
2887	IPCKSPELEDTATSSK2000000000000	sp Q13107 UBP4_HUMAN
2888	EQLSETEGSGEDEPGNDPSETTQK2.002E+19	sp Q8N108 MIER1_HUMAN
2889	DSSGQEDTQSSNDDPSQSVASQDAQEIRPR2.000002E+30	sp Q8TEW0 PARD3_HUMAN
2890	FSPDSQYIDNR200000000000	sp Q9NR30 DDX21_HUMAN
2891	NGFPHPEDCNPSEAASEESNSEIEQIPVEQK2000000000000	sp Q96S38 KS6C1_HUMAN
2892	SPEESFDIK2000200000	sp Q5UIP0 RIF1_HUMAN
2893	ATSEEDVSIKSPICEK2000000	sp O15234 CASC3_HUMAN
2894	DPSPEADAPVLGSPEK200000000000000	sp Q9BTC0 DIDO1_HUMAN
2895	GGDDHDDTSDSDGLTLK2002020000000	sp Q92993 KAT5_HUMAN
2896	SNCLGTDEDSQDSSDGIPAPR2.000000002E+22	sp Q8IYP5 FUND1_HUMAN
2897	NPPPQDYESDDSYEVLDTYAR2000000000000	sp P18669 PGAM1_HUMAN
2898	FSGWYDADLSPAGHEEAK20000000000000000	sp Q96MG7 MAGG1_HUMAN
2899	GPGGSQGSQPSPQGAR2000000	sp Q96S55 WRIP1_HUMAN
2900	RPAAAAAAGSASPR2000	sp Q96B23 CR025_HUMAN
2901	RDSSSQLASTESDKPTTGR2200000000000000000	sp P55081 MFAP1_HUMAN
2902	KIVEPEVVGESDSEVEGDAWR202000000000	sp Q9UKJ3 GPTC8_HUMAN
2903	KPSVSEEVQATPNK2000000000000	sp Q07960 RHG01_HUMAN
2904	SDDSKSSPELVTHLK22000000000	sp Q9TNN7 1C05_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
2905	GGSCSQAASSNSAQGSDESLIACK2000000000	sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
		sp P30510 1C14_HUMAN
2906	VDSEGFSENDAAGDFR200000000000	sp O43823 AKAP8_HUMAN
2907	RASQGLLSIENSESDSSEAK2E+19	sp Q5UIP0 RIF1_HUMAN
2908	NDKSEEEQSSSVK200000000000	sp P07910 HNRPC_HUMAN
2909	LHYTPPLQSPIIDNDPLLQSPWR20000000000000000	sp Q5TKA1 LIN9_HUMAN
2910	REESPMVDQPSAQDTQSIASDGTQGEK1.00000002E+26	sp Q7Z6Z7 HUWE1_HUMAN
2911	YDWSGAEESDDENAVCAAQNCQRPK2.00002E+23	sp P29375 KDM5A_HUMAN
2912	TPPVALNSSR200000000000	sp Q9UQ35 SRRM2_HUMAN
2913	KLSPIIEDSR200000200	sp O60566 BUB1B_HUMAN
2914	VISDESIDIGSDVEFKPDTK2.000200002E+19	sp P52701 MSH6_HUMAN

2915	EGDVDVSDSDEDDNLPANFDTCHR2. 02E+19	sp Q9NYV6 RRN3_HUMAN
2916	GSGTASDDEFENLR20002000000000	sp Q7Z6Z7 HUWE1_HUMAN
2917	KPSDSLVAASSR2020022000	sp Q5T0W9 FA83B_HUMAN
2918	GHNEGNQLISAEPVIVSSDEEGPVEHK220000000000	sp Q9BQF6 SENP7_HUMAN
2919	IVSGSPISTPSPSPLPR20000000200000	sp Q9ULM3 YETS2_HUMAN
2920	SSSLKSAQGTGFELGQLQSIR2. 2E+20	sp Q9C0B5 ZDHC5_HUMAN
2921	GTSDSSSGNVSEGESPPDSQEDSFQGR200000000000000000	sp Q8WUB8 PHF10_HUMAN
2922	TIQNSSVSPTSSSSSSSTGETQTQSSSR2000000000000000	sp Q7Z417 NUFP2_HUMAN
2923	SAAEMYGSVTEHPSPSPLLSSSFDLDYDFQR200000000000000000	sp P07910 HNRPC_HUMAN
2924	KEESEESDDMGFLFD200200010000000	sp P05387 RLA2_HUMAN
		sp P05386 RLA1_HUMAN
2925	DLPTIPGVTSPSSDEPPMEASQSHLR20000000000000000000	sp Q13541 4EBP1_HUMAN
2926	ELDDSSSESSDLQLEGPSTLR22022000000000000000	sp P29590 PML_HUMAN
2927	TGSGSPFAGNSPAR200000000000	sp Q9UPT8 ZC3H4_HUMAN
2928	VSALEEDMDDVESSEEEEEDEK100002200000000000	sp Q8NAV1 PR38A_HUMAN
2929	SLDSEPSVPSAAKPPSPEK20000	sp Q7Z3K3 POGZ_HUMAN
2930	LSVPTSDEEDEVPAKPR20022000000000000000	sp Q8NE71 ABCF1_HUMAN
2931	GLFSDEEDEDLFSQSASNLK2. 00000000002E+19	sp Q9Y4E1 FA21C_HUMAN
2932	TPASINATPANINLADLTR2. 002E+19	sp Q7KZ85 SPT6H_HUMAN
2933	RGSGDTSSLIDPDTLSELR20000000000000000000	sp Q9Y608 LRRF2_HUMAN
2934	QSHSEPSLQSK20000000	sp Q9UQ35 SRM2_HUMAN
2935	LGSPDSTSPSSPTFWNYSR2000000000000000	sp Q96SK2 TM209_HUMAN
2936	GPSPAPAPAASPQPGSR2000000020000000	sp Q8IX07 FOG1_HUMAN
2937	LASPSGTSSGLEVVAPGTSAPGGGPGTLDDSATICR2. 22E+32	sp Q13263 TIF1B_HUMAN
2938	SVEPQDAGPLER20000000000000	sp Q9NZT2 OGFR_HUMAN
2939	HDSPDLAPNVYSLPR2000000000000000	sp Q9BRD0 BUD13_HUMAN
2940	DISEGGESPVVQSDEEEVQVDTALATSHTDR2. 00002E+24	sp Q96EV8 DTBP1_HUMAN
2941	TEARSSDEENGPPSSPDLDR220000000020000000	sp Q96B36 AKTS1_HUMAN
2942	LCSSAETLHSPDIGK2000000000000000	sp 096028 NSD2_HUMAN
2943	TGSYGALAEITASK2000000000000000	sp 014974 MYPT1_HUMAN
2944	SPPPEMDTPTSTR2000000000000000	sp P46013 KI67_HUMAN
2945	LSTTPSPTSSLHEDGVEDFR2. 20000002E+19	sp 094842 TOX4_HUMAN
2946	ASPITNDGEDEFVPSDGLDKDEYTFSPGK2E+28	sp Q02880 TOP2B_HUMAN
2947	QVPDAAATATAYLCGVK2000000000000000	sp P09923 PPBI_HUMAN
2948	LVDVSPDR20000	sp Q9NYZ3 GTSE1_HUMAN
2949	ASESSSEKDDYEIFVK20222000000000000000	sp P18583 SON_HUMAN
2950	VSDQNSPVLPK2000000	sp Q9UPU5 UBP24_HUMAN
2951	EISSPESCPPEIK220020000000	sp Q92889 XPF_HUMAN
2952	GRASSHSQTQGGGSVTK220000000000000000	sp P02545 LMNA_HUMAN
2953	SPGHMVIDQTK2000000000000000	sp Q9NWH9 SLTM_HUMAN
2954	SASPSVGEERPR20200000000000	sp Q9P1Y6 PHRF1_HUMAN
2955	IESDTEETQDTSVDHNETGNTGESSVEENEK2E+29	sp Q6PL18 ATAD2_HUMAN
2956	SKSQSSGSSATHPISVPGAR20000000000000	sp Q9Y4H2 IRS2_HUMAN
2957	LTFDITTFSPNTGK2000000	sp P45880 VDAC2_HUMAN
2958	LSAITMALQYGESEEDAALAAAR1. 00000202E+19	sp Q86U86 PB1_HUMAN
2959	SSTPLPTISSAENTR200000000000	sp P42166 LAP2A_HUMAN
		sp P42167 LAP2B_HUMAN
2960	KSPSGPVKSPPLSPVGTTPVK2000200000000000	sp Q9BVC5 ASHWN_HUMAN
2961	KRESESEDETTPAAPQLIK20202000000000000000	sp 060832 DKC1_HUMAN
2962	YSPTSPTYSPVYTPSPK2E+20	sp P24928 RPB1_HUMAN
2963	CPEILSDESSSDEDEK200222000000	sp P35659 DEK_HUMAN
2964	AASSSPGSPVASSPSR2002000200000	sp Q92538 GBF1_HUMAN
2965	EPEGPPSPGK20000	sp P58012 FOXL2_HUMAN
2966	VSEEQTQPPSPAGAMSTAMGR20000010001000	sp Q16666 IF16_HUMAN

2967	SASSGAEGDVSSEREP20000000000000	sp Q8TEA8 DTD1_HUMAN
2968	SPSFGEDYYGPSR20000000000000	sp P49761 CLK3_HUMAN
2969	GYSPYSVSGSGSTAGSR3000002000000000	sp Q15149 PLEC_HUMAN
2970	KQETAAVCGETDEEAGESGGEGIFR2000000200000000	sp Q9ULL5 PRR12_HUMAN
2971	RNSSDSAIDNPKPNK20000000000000	sp P46100 ATRX_HUMAN
2972	DGGSGNSTIIVSR200220000000	sp Q7Z6Z7 HUWE1_HUMAN
2973	SNSEVEDVGPTSHNR2000000000000000	sp Q13206 DDX10_HUMAN
2974	ADLNQGIGEPQSPSR20000	sp Q96C19 EFHD2_HUMAN
2975	TASGSSVTSLDGTR2000000000000000	sp Q92597 NDRG1_HUMAN
2976	SVSASSGPCSPAAPHPPPPR2.00002E+21	sp Q86UY5 FA83A_HUMAN
2977	LASTSDIEEK2000000000	sp O14974 MYPT1_HUMAN
2978	AMLDQLMGTSR200	sp Q9Y383 LC7L2_HUMAN
2979	LSSSDRYSDASDDSFSEPR20020020000000	sp Q9NRF8 PYRG2_HUMAN
2980	SHSPSSPDPTSPVGDSTR2000000000000000	sp Q13586 STIM1_HUMAN
2981	KSSISSISGR2000200000	sp Q86SQ0 PHLB2_HUMAN
2982	GLSQEGTGPPTSAGEGHSR200000000000000000	sp Q63HN8 RN213_HUMAN
2983	DMSPLSETEMALGK20000000000000	sp P27816 MAP4_HUMAN
2984	ASLVALPEQTASEEETPPPLLTK20000000000000	sp Q9UKV3 ACINU_HUMAN
2985	EEGPPPPSPDGASSDAEPEPPSGR200002200000000000	sp Q9NTJ3 SMC4_HUMAN
2986	RPESAPAESSPSK200000000000	sp Q07889 SOS1_HUMAN
2987	KPSPSEPEPWKPFPAVSPEPR202000000000000000	sp Q96JM3 CHAP1_HUMAN
2988	QEKAPEKPAETPVATSPTATDSTSGDSSR2000000000000000	sp P54727 RD23B_HUMAN
2989	SQASMVINSDDTDEEEVSAALTLAHLK1.000202E+23	sp Q14676 MDC1_HUMAN
2990	TPLSSLSATSNNSTVACIDR2020000000000000	sp P00533 EGFR_HUMAN
2991	SQEMVHLVNK200100000000	sp P16070 CD44_HUMAN
2992	GPSPSPVGPSPASVAQSR200000000000000000	sp O14497 ARI1A_HUMAN
2993	SDGACDSPSSDKENSSQIAQDHQK20000000200000000000	sp Q9UIG0 BAZ1B_HUMAN
2994	RSEDEPPAASASAAPPPQR20000000000000000000	sp Q96IZ0 PAWR_HUMAN
2995	MSPMGTASGSNSPTSASVQR2000000000000000	sp O76080 ZFAN5_HUMAN
2996	SPSPDDILER202000000000	sp Q13523 PRP4B_HUMAN
2997	IESDTEETQDTSVDHNETGNTGESSVEENEK2E+29	sp Q6PL18 ATAD2_HUMAN
2998	NNSNTCNIENELEDSRK2000000000000000	sp Q6PL18 ATAD2_HUMAN
2999	SSRSSPELTR200200000000	sp Q9UQ35 SRRM2_HUMAN
3000	IGHHSTDDSSAYR200000000000	sp P12694 ODBA_HUMAN
3001	MHDGELEEQEEDDEKSDSEGGDLK1E+25	sp Q9NU22 MDN1_HUMAN
3002	ANSDDERPVASDNDDEK200000002000000000	sp Q8WVC0 LEO1_HUMAN
3003	AVPPVSPELR2000000	sp Q96JM3 CHAP1_HUMAN
3004	KPPAPSPVQSQSPSTNWSPAVPVK2.000202E+19	sp Q8IYB3 SRRM1_HUMAN
3005	KPVTVSPTTPTSPTTEGEAS20200000000000	sp Q9Y6G9 DC1L1_HUMAN
3006	KSPSPGVKSPPLSPVGTTPVK20000200020000000000	sp Q9BVC5 ASHWN_HUMAN
3007	ITETGAVKPTGECSGEQSPDTNYEPPGEDK2000000000000000	sp P49454 CENPF_HUMAN
3008	TASSDSEEDPEALEK20022020000000000000	sp Q8N9T8 KRI1_HUMAN
3009	AAPEASSPPASPLQHLLPGK200020000000000000	sp Q96TA1 NIBL1_HUMAN
3010	GSGSACSLCCCGR2000000000000000	sp Q9GZU1 MCLN1_HUMAN
3011	APIDTSDVEEK20000000	sp Q06265 EXOS9_HUMAN
3012	TPAAPASPAAVPSEGSGSTGWR20000000000000000000	sp P07199 CENPB_HUMAN
3013	TDALTSSPGR20000	sp P49736 MCM2_HUMAN
3014	KPIDSLRDSR200	sp Q9UQ35 SRRM2_HUMAN
3015	SSGHSSSELSPAVEK200000000	sp Q9UQ35 SRRM2_HUMAN
3016	STAQQELDGKSPSTPVIVASHTANKEEK200000000000000000	sp P35606 COPB2_HUMAN
3017	LASPSGSTSSGLEVVPEGTSAPGGGPGTLDDSATICR2E+32	sp Q13263 TIF1B_HUMAN
3018	SQLDDHPESDDEENFIDANDEDMEK20000000000000000000	sp Q03701 CEBPZ_HUMAN
3019	QNPEQSADEDAEKNEEDSEGSSDEDEDEDGVSAAATFLK2.0022E+21	sp B5ME19 EIFCL_HUMAN
		sp Q99613 EIF3C_HUMAN

3020	QGSQSSQMEVQEGYGFSGDDPYSSAEPHVSGVK2E+33	sp Q13263 TIF1B_HUMAN
3021	SPSPPTQHTGQPPGQPSAPSQLSAPR2E+24	sp Q68EM7 RHG17_HUMAN
3022	SKSQSSGSSATHPI SVPGAR200000000000000000	sp Q9Y4H2 IRS2_HUMAN
3023	QSETVDQNSDSEMLAILK20000002000000000	sp P46100 ATRX_HUMAN
3024	SHSESASPSALSSSPNLSPTGWSQPK2200002000000000	sp P04049 RAF1_HUMAN
3025	EADDDEEVDDNIPEMPSPK102000	sp P26358 DNMT1_HUMAN
3026	SELSQDAEPAGSQETK200000000000000000	sp Q9BVJ6 UT14A_HUMAN
3027	SVSPSPSVLSEGR20200000000000	sp Q86UW7 CAPS2_HUMAN
3028	TDHGAEIVYKSPVVGDTSPR200020002000	sp TAU_HUMAN
		sp P10636 TAU_HUMAN
3029	GGPASPGGLQGLETR200000000000	sp Q8NCD3 HJURP_HUMAN
3030	TTSAGTGMWR2000000000	sp P60468 SC61B_HUMAN
3031	TSSSSPANSDVEIDGIGR2000200000000000	sp O60293 ZC3H1_HUMAN
3032	SATPEPVTDNR202000000000	sp Q5VT52 RPRD2_HUMAN
3033	SPGPAADGPIDLSK2000000000000000	sp Q8IX07 FOG1_HUMAN
3034	TDPSLDLHLSAETQPAQEELR2022000000000000	sp P11171 41_HUMAN
3035	SSSVGSSSSYPI SPAVSR200220000000000000	sp Q15149 PLEC_HUMAN
3036	VGDSTPVSEKPVSAAVDANASESP200	sp Q9H8Y8 GORS2_HUMAN
3037	ALEVEESNSEDEASFK20200002000	sp Q03111 ENL_HUMAN
3038	LEKEEEEGISQESSEEEQ2200000	sp P17096 HMGA1_HUMAN
3039	ERTPESEEEVWETNR200200000000000000	sp Q5T200 ZC3HD_HUMAN
3040	ELSNSPLRENSFGSPLEFR202000000000000000	sp Q9UQ35 SRRM2_HUMAN
3041	LSSKLSAVSLR2002000	sp Q15036 SNX17_HUMAN
3042	SQSSDTEQQSPTSGGGK200000000	sp P23588 IF4B_HUMAN
3043	GLNSQSSDHLNK200000000000	sp A8CG34 P121C_HUMAN
		sp Q96HA1 P121A_HUMAN
3044	SVSTPSEAGSQDSGDGAVGSR2000000000	sp Q13409 DC1I2_HUMAN
3045	RTDAL TSSPGR20000	sp P49736 MCM2_HUMAN
3046	SPAYTPQNL DSESESGSSIAEK2E+22	sp Q15648 MED1_HUMAN
3047	TSPADHGGSVGSESGSAVDSVAGEHSVSGR2E+30	sp Q5T4S7 UBR4_HUMAN
3048	KVEEDLKADEPSSEESDLEIDK220000000000	sp P50502 F10A1_HUMAN
3049	EGDPVSLSTPLETEFGSPSELSPR200000000	sp Q9BUR4 WAP53_HUMAN
3050	ETAVPGLGIEDISP NLS PDDK2000200000	sp Q12888 TP53B_HUMAN
3051	ESEDKPEIEDVGSDEEEK2E+19	sp P07900 HS90A_HUMAN
3052	GQKPALDCSQVSPRPATSPENNASLSDTSPMDSSPSGIPK2. 000002E+30	sp P18858 DNLI1_HUMAN
3053	VGPATPSAQVGK200000000	sp Q13428 TCOF_HUMAN
3054	QNPQSPQDSSVTSK200000000000	sp Q9UBN7 HDAC6_HUMAN
3055	APKPPTDGSTSPTSTPSEDQEALGK2000000000000000	sp Q8NHM5 KDM2B_HUMAN
3056	SVDEVNYWDK20000000000	sp P12694 ODBA_HUMAN
3057	QKSAEPSPTVMSTSLGSLNSELDR2. 00020001E+22	sp P49023 PAXI_HUMAN
3058	SGPSPTTPQPLR2002000000000	sp Q9Y4F5 C170B_HUMAN
3059	RKSLDSESDSK200020000000	sp Q13185 CBX3_HUMAN
3060	AFGPGLQGSAGSPAR20000	sp P21333 FLNA_HUMAN
3061	GILAADESTGSI AK20000000	sp P04075 ALDOA_HUMAN
		sp ALDOA_RABIT
3062	KASSDSEDSSEEEEEVQGPPAK2. 20020022E+21	sp Q14978 NOLC1_HUMAN
3063	YQPLASTASDNDFVTPEPR3. 00000002E+19	sp Q15021 CND1_HUMAN
3064	SVEEPTQPGGTGLSDSR200000000000000000	sp Q9H0B6 KLC2_HUMAN
3065	SNEDSLILEK20002000000	sp O94880 PHF14_HUMAN
3066	DLFDYSPPLHK2000000	sp Q9NYF8 BCLF1_HUMAN
3067	KFSLDEL AGPGAEGPSNLK200000000000000000	sp O76064 RNF8_HUMAN
3068	SSSLGSGSSRDSISSDSETSEPLSCR2020000000000000	sp O75164 KDM4A_HUMAN
3069	KPSTSDSDSNFEK2220020000000	sp P11388 TOP2A_HUMAN



3070	KLSSERPSSDGEVVENGITTCNGK220000000000000000	sp Q8N556 AFAP1_HUMAN
3071	KRESESEDETTPAAPQLIK202000000000000000	sp O60832 DKC1_HUMAN
3072	ILDNPNTGEPAPVLSSPPPADVSTFLAFPSPEK2E+19	sp P46060 RAGP1_HUMAN
3073	GLSASLPDLSENWIEVK2000000000000000	sp Q6PKG0 LARP1_HUMAN
3074	TSPSSPAPLPHQEATPR202000000000000000	sp P04920 B3A2_HUMAN
3075	SSSSSSGGGLLPYPR200002000000000000	sp O60293 ZC3H1_HUMAN
3076	LPDSPALAK2000000	sp Q13586 STIM1_HUMAN
3077	KKEPAITSQNSPEAR200000000	sp P23193 TCEA1_HUMAN
3078	SPGEPGGAAPER2000000000000000	sp Q96EB6 SIR1_HUMAN
3079	KPSDSLVSASSR2000020000	sp Q5T0W9 FA83B_HUMAN
3080	VLSSTSEEDPGVVK2200000000000000	sp Q8NI08 NCOA7_HUMAN
3081	TASGSSVTLSDGTR2020000000000000	sp Q92597 NDRG1_HUMAN
3082	ESSANNSVSPESLR20002000	sp Q04726 TLE3_HUMAN
3083	SSTVAPAQPDGAESEWTDVETR2E+22	sp Q02241 KIF23_HUMAN
3084	TPQEAIMDGTEIAVSPR2000	sp Q06587 RING1_HUMAN
3085	LPSSPVYEDAASF2000000000000000	sp Q14247 SRC8_HUMAN
3086	RGSSGSVDETLFALPAASEPVIR200000000000000000	sp Q9NQC3 RTN4_HUMAN
3087	SLGYHGSSYSEGVPEVSPVSSPSLTHDK2.00000002E+20	sp Q9Y618 NCOR2_HUMAN
3088	GQSLSNPTDDSWK2002000000000000	sp Q9Y6D5 BIG2_HUMAN
3089	THDHQLESSLSPVEVFAK2002000000000000	sp Q9H410 DSN1_HUMAN
3090	ASSPSPLTIGTPESQR200200000000000000	sp Q9NPI6 DCP1A_HUMAN
3091	MEQQYESSSDGTEKLPER2220000000000000	sp P46100 ATRX_HUMAN
3092	YSDGDKSDDLVVDVSNEDPATPR2.00002E+22	sp Q04726 TLE3_HUMAN
3093	EEGSLSDTEADAVSGQLPDPTNPSAGK2E+25	sp Q8N1G4 LRC47_HUMAN
3094	SAESPTSPVTSETGSTFK2000000000000000	sp Q02952 AKA12_HUMAN
3095	SDSPEIPFQAAAGPSDGLDASSPGNSFVGLR2E+29	sp Q12888 TP53B_HUMAN
3096	QHEVEIPSPTQK2000	sp Q08499 PDE4D_HUMAN
3097	AFAAVPTSHPPEDAPAPPTPGPAASPEQLSFR2000000000	sp Q14160 SCRIB_HUMAN
3098	AEQLQDAEEEKDSDNEENKDSLVDDEEEK200000002000000000	sp Q9H1E5 TMX4_HUMAN
3099	NLNSNLFSVNR200000	sp P52948 NUP98_HUMAN
3100	LPSTSGSEGVVFR2002000000000000	sp P29317 EPHA2_HUMAN
3101	NKSPAAVTEPETNK2000000000000000	sp O75449 KTNA1_HUMAN
3102	HAPTYTIPLSPVLSPTLPAEAPTAQVPPSLPR2.0002E+23	sp Q9NPI6 DCP1A_HUMAN
3103	EVIEIEDASPTK20000	sp P46100 ATRX_HUMAN
3104	GPPDFSSDEEREPTPVLGSGAAAAGR2E+21	sp P42167 LAP2B_HUMAN
3105	TNSSDSERSPDLGHSTQIPR200200000000000000	sp P42166 LAP2A_HUMAN
3106	RQSVSPPYKEPSAYQSSTR202000000000000000	sp Q6VY07 PACS1_HUMAN
3107	TESPPSPSSVEDAR2002002000000000	sp Q9NYV4 CDK12_HUMAN
3108	SERPPTILMTEEPSSPK2000	sp Q9BRK4 LZTS2_HUMAN
3109	KGGSYSQAASSDSAQGSVSLTACK2E+22	sp Q5VZK9 LR16A_HUMAN
3110	SETIQDQSLVGSSTR200000	sp Q09160 1A80_HUMAN
3111	TEMKSPFNPSQDSR100200000200000000	sp P05534 1A24_HUMAN
3112	TASISSPSEGTPVGSYGCTPQSLPK2.02E+25	sp P30447 1A23_HUMAN
3113	NAEESESEAEED2020000000	sp P01891 1A68_HUMAN
3114	SLSPSHLTEDR2020000000000000	sp P01892 1A02_HUMAN
3115	SQSKSPTGTPAR2020200000000000	sp P10316 1A69_HUMAN
3116	LASPSGSTSSGLEVVAPEGTSAPGGGPGTLDDSATICR2E+34	sp Q9NZN5 ARHGAP_HUMAN
3117	ALEYSNGIFDCSPTSPFMGLR2002000000000000	sp P08651 NFIC_HUMAN
		sp Q6PKG0 LARP1_HUMAN
		sp Q7L1Q6 BZW1_HUMAN
		sp Q5T200 ZC3HD_HUMAN
		sp Q13595 TRA2A_HUMAN
		sp Q13263 TIF1B_HUMAN
		sp Q86W92 LIPB1_HUMAN

		sp Q9TNN7 1C05_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
3118	GGSCSQAASSNSAQGSDESLIACK2E+20	sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
		sp P30510 1C14_HUMAN
3119	SSSPFLSK202000000	sp Q9NYV4 CDK12_HUMAN
3120	SYQNSPSSDDGIRPLPEYSTEK20020000000000000000	sp Q15648 MED1_HUMAN
3121	SGSSQELDVKPSASPQER2000000000200000	sp Q9UQ35 SRRM2_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
		sp P05534 1A24_HUMAN
3122	KGGSYSQAASSDSAQGSVDVSLTACK2000000000	sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
		sp P01892 1A02_HUMAN
		sp Q9BQE3 TBA1C_HUMAN
		sp Q71U36 TBA1A_HUMAN
3123	TIGGGDDSFNTFFSETGAGK20000000000000	sp P68363 TBA1B_HUMAN
		sp Q6PEY2 TBA3E_HUMAN
		sp Q13748 TBA3C_HUMAN
3124	LADSGDGAGPSPEEK200000	sp Q92917 GPKOW_HUMAN
3125	YSDGDKSDDLVDVSNEDPATPR2E+22	sp Q04726 TLE3_HUMAN
3126	MEQQYESSSDGTEKLPER2020000000000	sp P46100 ATRX_HUMAN
3127	SPSPPTQHTGQPPGQPSAPQLSAPR2E+26	sp Q68EM7 RHG17_HUMAN
3128	SSDSESNENEPHLPHGVSQR2E+19	sp Q9BQ52 RNZ2_HUMAN
3129	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR20000000000000000000	sp Q8WUZ0 BCL7C_HUMAN
3130	NSNSPPSPSSMNQR200200000000	sp Q7Z5L9 I2BP2_HUMAN
3131	VETVSPSESPKDTIDK20200000000	sp Q99590 SCAFB_HUMAN
3132	DKDVTLSPVK20000	sp Q8NFC6 BD1L1_HUMAN
3133	SEEEQSSSVKDETENVK20020000000000	sp P07910 HNRPC_HUMAN
3134	LAELPAAAQPSAEDSDTEDDSEAEQTER202000000000000	sp Q095714 HERC2_HUMAN
3135	SGEQIFTQDVRPEQLLEGK2E+19	sp Q8WXE0 CSKI2_HUMAN
3136	VLQSDSETEDTNASPEK202000000020000	sp Q9HAW4 CLSPN_HUMAN
3137	LENVSLSLDKSPTEK200000	sp Q86W56 PARG_HUMAN
3138	RYPSSISSSPQK200200000	sp Q14157 UBP2L_HUMAN
3139	VPPAPVPCPPSPGPSAVPSSPK2000000020000	sp Q095817 BAG3_HUMAN
3140	IGHHSTSDSSAYR20000000000	sp P12694 ODBA_HUMAN
3141	ASPITNDGEDEFVPSDGLDKDEYTFSPGK2E+28	sp Q02880 TOP2B_HUMAN
		sp Q09160 1A80_HUMAN
		sp P05534 1A24_HUMAN
3142	KGGSYSQAASSDSAQGSVDVSLTACK20000000000000	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
3143	GYNHGQGSYSYSNSYNSPGGGGSDYNYESK20000000000000000000	sp Q12906 ILF3_HUMAN
3144	VKPASPVAQPK20000000	sp Q96EV2 RBM33_HUMAN
3145	KAEGEPQEE SPLK20000	sp Q9NYF8 BCLF1_HUMAN
3146	LKEEHGIELSSPR2000	sp Q014647 CHD2_HUMAN
3147	ITEVSCKSPQDPVK200000000	sp P46013 KI67_HUMAN

3148	SHILEDDENSVDISMLK200000000200000000	sp 075717 WDHD1_HUMAN
3149	RNSVTPLASPEPTK200000000000	sp Q16875 F263_HUMAN
3150	AAFDDAIAELDTLSEESYK2000000	sp P62258 1433E_HUMAN
3151	SEDESETEDEEEKSQEDQEQR202000000000000000	sp P25205 MCM3_HUMAN
3152	LVEPHSPSPSSK20000000	sp 015014 ZN609_HUMAN
3153	TPSSPLQSCPTSLSPR20000000000000	sp Q96AP0 ACD_HUMAN
3154	GGGGYTCQSGSGWDEFTK20000000000	sp P15924 DESP_HUMAN
3155	SSTPLPTISSAENR22000000000000000	sp P42166 LAP2A_HUMAN
		sp P42167 LAP2B_HUMAN
3156	YHGHSMSDPGVSYR20000002000	sp P29803 ODPAT_HUMAN
		sp P08559 ODPA_HUMAN
3157	FTMDLSDSEDFSDFEKTDDEDFVPSDASPPK1.000200002E+30	sp P11388 TOP2A_HUMAN
3158	SRSRDSGDENEPIQER202002000000000000	sp Q8WX93 PALLD_HUMAN
3159	SHSESASPSALSSSPNNLSPTGWSQPK2.0022E+19	sp P04049 RAF1_HUMAN
3160	EHIEIIAPSPQR20000	sp Q9NZJ5 E2AK3_HUMAN
3161	YLSADSGDADDSDADLGS AVK20000000000	sp Q15361 TTF1_HUMAN
3162	GSSDGRGSDSEDLPHR20000000000	sp Q9UPQ0 LIMC1_HUMAN
3163	GSGGSSGDELREDDEPVK200220000000000000	sp Q7Z4V5 HDGR2_HUMAN
3164	DSSKGEDSAEETEAKPAVVAPAPVVEAVSTPSAAFPSDATAEQGPILTK2E+42	sp Q12906 ILF3_HUMAN
3165	GHTDTEGRPPSPPTSTPEK20000200000	sp Q00613 HSF1_HUMAN
3166	GSLTPHPAACDKDGTDSAAEESSQETSK2020000000000000	sp P55201 BRPF1_HUMAN
3167	SQSDLDDQHDYDSVASDEDTDQEPLR302002000000000000	sp Q9Y2X7 GIT1_HUMAN
3168	GLMAGGRPEGQYSEDEDTDTDEYKEAK2000020000000000	sp Q9NPQ8 RIC8A_HUMAN
3169	KPSPQPSSPR2000	sp 075909 CCNK_HUMAN
3170	GDESEEEDELEVPVPSR2000000000000000	sp P13861 KAP2_HUMAN
3171	SKSPKSPPEEGAVSS202000000000000000	sp Q01130 SRSF2_HUMAN
3172	TASRPDDIPDPSPPK22000	sp Q5VZK9 LR16A_HUMAN
3173	SSPVCSTAPVETEPK2000000000000000	sp Q96BY7 ATG2B_HUMAN
3174	AGPTLTDENGDDLGLPPSPGDTSYQQDQVDDFHEAR2E+31	sp Q9NQZ2 SAS10_HUMAN
3175	SLSPSHLTEDR20000000000	sp Q5T200 ZC3HD_HUMAN
3176	SRSFDYNYR20000000	sp 075494 SRS10_HUMAN
3177	KEENADSDDEGELQDLLSQDWR200000000000000000	sp Q96NB3 ZN830_HUMAN
		sp P10636 TAU_HUMAN
3178	SGYSSPGSPGTPGSR200000000	sp TAU_HUMAN
3179	VHPYEDSDSSEDEVDWQDTR2022000000000000	sp 014730 RIOK3_HUMAN
3180	RLSTSPDVIQGHQPR200000000000	sp Q9Y385 UB2J1_HUMAN
3181	SASVSSISLTK20000000000	sp Q07889 SOS1_HUMAN
3182	KPSPQPSSPR200000000	sp 075909 CCNK_HUMAN
3183	IDAGTMAEPSASPSK20000	sp Q8IXQ3 CI040_HUMAN
3184	SSSVGSSSSYPISPA VSR200200030000000000	sp Q15149 PLEC_HUMAN
3185	SQSSEGVSSLSSSPNSLETQSQSLSR2.2E+20	sp 095155 UBE4B_HUMAN
3186	FSREEFPTLQAAGDQDK200000000000000000	sp P48634 PRC2A_HUMAN
3187	TSFVSGSDELGP IR20000000000	sp Q9NRY4 RHG35_HUMAN
3188	AEAPLPSPK2000	sp Q09666 AHNK_HUMAN
3189	NLSSDEATNPISR200000000000	sp Q7Z417 NUFP2_HUMAN
3190	GLGSSPDLELLLR220000000000	sp Q9NQQ7 S35C2_HUMAN
3191	LSTTPSPTSSLHEDGVEDFR2220000000000000	sp 094842 TOX4_HUMAN
3192	QPPVSPLTLSPGPEAHQGFSR2000020000000000000	sp Q6UUV7 CRTC3_HUMAN
3193	VATLNSEESDPPTYK200000000000	sp Q00341 VIGLN_HUMAN
3194	APSDSSLGTPSDGRPELR200200000000000000	sp Q15642 CIP4_HUMAN
3195	SGSSSPDSEITELK220020000000	sp P17812 PYRG1_HUMAN
3196	QGSFSEDVISHK200000000000	sp Q9UPN3 MACF1_HUMAN
3197	VQGGVPAGSDEYEDECPHLIALSSLNR2E+19	sp Q9BVS4 RIOK2_HUMAN

3198	AASSSPGSPVASSPSR2200000000000	sp Q92538 GBF1_HUMAN
3199	SPPASPESWK20002000000	sp Q96JM3 CHAP1_HUMAN
3200	EGAASPAPETPQPTSPETSPK200002000	sp Q9Y3X0 CCDC9_HUMAN
3201	QRPSYDIFEDSDDSEPGPPAPR2002000000000	sp Q9UMN6 KMT2B_HUMAN
3202	KKASSDSEDSSEEEEEVQGPPAK2. 2020022E+20	sp Q14978 NOLC1_HUMAN
3203	TESPATAAETASEELDNR2000000000000000000	sp Q9NTJ3 SMC4_HUMAN
3204	YHGHSMSDPGVSYSR20000002000	sp P08559 ODPA_HUMAN
3205	YSHSYLSDSDTEAK202000000	sp P29803 ODPAT_HUMAN
3206	ESLKEEDESDDNM20000000000010	sp Q92614 MY18A_HUMAN
3207	SRSSSPVTELASR20220000000000	sp P25788 PSA3_HUMAN
3208	LGSPKPER2000000	sp Q9UQ35 SRRM2_HUMAN
3209	SAEPSPTVMSTSLGSNLSELDR2020000000000000000	sp Q86VM9 ZCH18_HUMAN
3210	SPQGSPVLAEDSEGE200000	sp P49023 PAXI_HUMAN
3211	EVEEFSEDDDEDDSDSEAEK20000000200200000	sp Q99638 RAD9A_HUMAN
3212	ILDPTGEPAPVLSSPPADVSTFLAFPSPEK2000000000000020000	sp Q9Y2W2 WBP11_HUMAN
3213	SASSDTSEELNSQDSPK20220200000000000000	sp P46060 RAGP1_HUMAN
3214	TSPADHGGSVGSESGSAVDSVAGEHSVSGR2. 00202002E+23	sp 014745 NHRF1_HUMAN
3215	APSASPLALHASR202000000000	sp Q5T4S7 UBR4_HUMAN
3216	VNDAEPGSPEAPQGK200000000	sp Q8N3F8 MILK1_HUMAN
3217	SLGYHGSSYSPEGVEPVSPVSSPSLTHDK2. 00000002002E+20	sp Q8N8A6 DDX51_HUMAN
3218	EEETSIDVAGKPNEVTK20000000000000	sp Q9Y618 NCOR2_HUMAN
3219	SLSPGKENSALDMEK2020000000000000	sp P53985 MOT1_HUMAN
3220	RPTPQDSPIFLPVDDTSFR20000000000000	sp Q9H307 PININ_HUMAN
3221	KPLAAPGDGEGLGQTAQPSPPAR200000	sp P49327 FAS_HUMAN
3222	SPPKSPEEEGAVSS200020000000020	sp Q9Y4F5 C170B_HUMAN
3223	SPQGSPVLAEDSEGE20000000000200000	sp Q01130 SRSF2_HUMAN
3224	LSSSLGR20000	sp Q99638 RAD9A_HUMAN
3225	VDDDSLGEFPVTNSR200000000000	sp Q9BST9 RTKN_HUMAN
3226	TLTDEVNSPDSRRR20000000	sp Q92785 REQU_HUMAN
3227	GRLTPSPDIVLSDNEASSPR20000002000002000	sp Q8TAQ2 SMRC2_HUMAN
3228	MEEEGTEDNGLEDDSR200	sp Q8WXI9 P66B_HUMAN
3229	LKFSDEEDGRDSDEEGAEGHR2000000020000000000	sp Q14151 SAFB2_HUMAN
3230	HGGSPQPLATTPLSQEPVNPPEASPTR2E+25	sp P48634 PRC2A_HUMAN
3231	ASESSSEEKDDYEIFVK20022000000000000	sp Q9UQ35 SRRM2_HUMAN
3232	SPAPLLHVAALGQK2000000000000000	sp P18583 SON_HUMAN
3233	DDISEIQLASDHSGR2002002000	sp Q8N163 CCAR2_HUMAN
3234	KGGSYSQAASSDSAQGSVDVSLTACK2E+20	sp Q07157 ZO1_HUMAN
3235	SYQNSPSSDDGIRPLPEYSTEK2. 0000002E+22	sp P01891 1A68_HUMAN
3236	METEADAPSPAPSLGER2000000000	sp P01892 1A02_HUMAN
3237	DTSPSSGSAVSSSK2000000000000	sp P05534 1A24_HUMAN
3238	TVDSQGPTPVCTPTFLER2000000020000000	sp Q09160 1A80_HUMAN
3239	MESEGGADDSAEEDLLDDDDNEDRGDDQLELIKDDEK1. 020000002E+38	sp P10316 1A69_HUMAN
3240	RPETVVPGEATETDSER202000	sp P30447 1A23_HUMAN
3241	GAPSSPATGVLSPQK2000000000000000	sp Q15648 MED1_HUMAN
3242	FFTTGSDSESESSLGEEELVTKPVGGNYGK2. 020202202E+25	sp Q12873 CHD3_HUMAN
3243	DELHIVEAEAMNYEGSPIK30000000	sp Q8NEY8 PPHLN_HUMAN
		sp Q9BYG3 MK67I_HUMAN
		sp P07910 HNRPC_HUMAN
		sp Q6QNY0 BL1S3_HUMAN
		sp Q5SRE5 NU188_HUMAN
		sp B5ME19 EIFCL_HUMAN
		sp Q99613 EIF3C_HUMAN
		sp P06748 NPM_HUMAN

3244	DSSTSPGDYVLSVSENSR20000000000000	sp P46108 CRK_HUMAN
3245	SEAGHASSPDSEVTSLCQK20000000000000	sp Q6NZY4 ZCHC8_HUMAN
3246	VGSSKPDSSPVLSPGNK2000000000	sp Q8N4C8 MINK1_HUMAN
3247	SPGSPGPPELPIETALDDR2000000000000000	sp Q8TEW0 PARD3_HUMAN
3248	GTSRPPPEGGLGYSQLGDDDLK2E+20	sp P21127 CD11B_HUMAN
3249	SPATSPISSNSHR20020000000000	sp Q9UQ88 CD11A_HUMAN
3250	DDISEIQSLASDHSGR2002000	sp Q9NR09 BIRC6_HUMAN
3251	ASESTARSSSESEDEDVIPATQCLTPGIR2. 22202E+23	sp Q07157 ZO1_HUMAN
		sp Q13428 TCOF_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
3252	GGSYSQAASSDSAQGSVSLTACKV20020000000	sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
3253	KPSPEPEGEVGPPK20000000000000	sp Q7Z5L9 I2BP2_HUMAN
3254	GYSPYVSGSGSTAGSR200000000000	sp Q15149 PLEC_HUMAN
3255	VQEHEDSGDSEVENEAK200000000000	sp Q9Y5J1 UTP18_HUMAN
3256	CTLPEHESPSQDISDACEAESTER2E+23	sp Q32MZ4 LRRF1_HUMAN
3257	CPEILSDESSSDEDEKK2002220000000	sp P35659 DEK_HUMAN
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3259	SRSPLDKDTYPPSASVVGASVGGHR202000000000000	sp Q96T37 RBM15_HUMAN
3260	AMSTTSSPPQPGK20000000000000	sp Q9UJU6 DBNL_HUMAN
3261	SPVSTRPLPSASQK2000000000000000	sp Q8ND56 LS14A_HUMAN
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3264	TDSV I IADQTPPTR20000	sp P17544 ATF7_HUMAN
3265	GNIETSEDEDDTADLFPEESGFEHAWR2. 2000002E+23	sp Q9H501 ESF1_HUMAN
3266	TNSDSALHQSTMPTQPESFSSGSQDVHQK2E+21	sp Q6UUV9 CRTC1_HUMAN
3267	ALQLSNSPGASSAFLK200000000000	sp Q12802 AKP13_HUMAN
3268	AGSPLCNSNLQDSEEDIEPSYSLFR2000000000000000	sp Q9NRY4 RHG35_HUMAN
3269	ESESEDETTPAAPQLIK2020000000000000	sp O60832 DKC1_HUMAN
3270	SLSSSLDDTEVKK20000000000000	sp O95292 VAPB_HUMAN
3271	AKSPTSPSPPR20200020000	sp Q8IYB3 SRRM1_HUMAN
3272	VNPHKVPASSVDSNIPSSQGYK200220000000000000	sp Q9UHB7 AFF4_HUMAN
3273	ATVTPSPVK20000	sp Q9H1E3 NUCKS_HUMAN
3274	LGSTSGEESDLER2220000000000	sp Q9BW71 HIRP3_HUMAN
3275	VKEEHLDVASPDKASSPELPSVENIK220000000000000	sp Q969R5 LMBL2_HUMAN
3276	LIESLSPDFCHQNK2000000000	sp Q9H6R7 CBO44_HUMAN
3277	YNLDASEEEDSNKK2000000000	sp O95218 ZRAB2_HUMAN
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3280	KGSSGNASEVSVACLTER2000000000000000	sp Q69YQ0 CYTSA_HUMAN
3281	TRPGSFQSLSDALSDTPAK200000000000	sp Q9H9C1 SPE39_HUMAN
3282	SPAQILISLESEER2000000000000000	sp P11532 DMD_HUMAN
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3284	AEAPSSPDVAPAGK2000000000	sp Q9P1Y6 PHRF1_HUMAN
3285	DLSEVSETTESTDVK20020000000000	sp P41227 NAA10_HUMAN
3286	VDSPSHGLVTSSLCIPSPAR20000000000000000000	sp Q9UER7 DAXX_HUMAN
3287	LSSLRSTSK200020000	sp P62753 RS6_HUMAN
3288	TSGGDHAPDPSGENSPAPQGR2000000000000000	sp Q96NY9 MUS81_HUMAN
3289	DHNVGSPKSPHASPQHTQTEK20002000000000000000	sp A0JNW5 UH1BL_HUMAN
3290	TLSTHSVPNISGATCSAFASPFGPCPYSHR2. 00002E+24	sp P28290 SSFA2_HUMAN
3291	SPISDNSGCDAPGNSNPSLSPSSAESEK2E+26	sp Q99590 SCAFB_HUMAN

3292	QRPDLGSAQK20000	sp P33991 MCM4_HUMAN
3293	TEMKSPFNPSQDSR2000000000000000	sp P08651 NFIC_HUMAN
3294	ASVALTPTAPSR200020000000	sp P28290 SSFA2_HUMAN
3295	QKSDAEEDGGTVSQEEDRKP2E+20	sp P27824 CALX_HUMAN
3296	LDFIESDPCSSEALSK2000000000	sp 043432 IF4G3_HUMAN
3297	KWDGSEEDNSK2000000000	sp 060841 IF2P_HUMAN
3298	GPSPSSPTPPAAAAPAEQAPR200000000000000000	sp Q86VQ1 GLC11_HUMAN
		sp Q641Q2 FA21A_HUMAN
3299	SRPTSFADELAAR20002000000000	sp Q9Y4E1 FA21C_HUMAN
		sp Q5SNT6 FA21B_HUMAN
3300	SLSPGKENVSAIDMEK2020000000001000	sp Q9H307 PININ_HUMAN
3301	SSSPAPADIAQTVQEDLR220000000000000000	sp Q13283 G3BP1_HUMAN
3302	IPSEMPSPNFGK2000000	sp Q9H2J7 S6A15_HUMAN
3303	GSGTASDDEFENLR202000000000	sp Q7Z6Z7 HUWE1_HUMAN
3304	IHDLEDDLEMSDASDASGEEGRVPK220020020000000000	sp P35269 T2FA_HUMAN
3305	NTSVGSTSTGVESILEPCDEIEQK2E+22	sp Q5T8I3 F102B_HUMAN
3306	KSPAQSPNPSTPR2000200000000	sp 076021 RL1D1_HUMAN
		sp TAU_HUMAN
3307	TDHGAEIVYKSPVSGDTSR200020000000	sp P10636 TAU_HUMAN
3308	AAQQSDSDGEALGGNPMVAGFQDDVDLEDQPR2.020000000001E+28	sp Q3YEC7 RABL6_HUMAN
3309	GAASTLVPGVSETSASPGSPSVR200000	sp Q6ZRS2 SRCAP_HUMAN
3310	TQSPGGCSAEAVLAR20000200000000	sp Q96MH2 HEX12_HUMAN
3311	SSDEENGPPSPDLDR220000000000000000	sp Q96B36 AKTS1_HUMAN
3312	NYQQNYQNSSEGEKNEGSESAPEGQAQR3.00000000002E+24	sp P67809 YBOX1_HUMAN
3313	SGSSPDSEITELKFPSINH22002000000000000000	sp P17812 PYRG1_HUMAN
3314	METVSNASSSNPSSPGR1000000000000200000	sp Q9NTI5 PDS5B_HUMAN
3315	APGSPAAPR2000000	sp Q9NQX5 NPDC1_HUMAN
3316	ESEDKPEIEDVGSDEEEK20000000000000000000	sp P07900 HS90A_HUMAN
3317	DMPGGFLFEGLSDDDDFHPNTR1.000000002E+22	sp Q96S82 UBL7_HUMAN
3318	VTTEIQLPSQSPVEEQSPASLSSLR2E+23	sp Q8WWI1 LMO7_HUMAN
3319	NYQQNYQNSSEGEKNEGSESAPEGQAQR200000000000	sp P67809 YBOX1_HUMAN
3320	SSSPTAGPSTEGAEGPEEK2.2E+19	sp Q8NHM5 KDM2B_HUMAN
3321	RPTETNPVTSNSDEECNETVK2200000000000000	sp P46100 ATRX_HUMAN
3322	AAVQELSSILAGEDPEER2200000000000000	sp P49768 PSN1_HUMAN
3323	VDSPSHGLVTSSLCIPSPAR200000000000020000	sp Q9UER7 DAXX_HUMAN
3324	FVSEGDGGR20000000	sp Q53EL6 PDCD4_HUMAN
3325	STQIENQHGAQDTSDLMSPSKR200000	sp Q9NVR2 INT10_HUMAN
3326	GPPDFSSDEEREPTPVLGSGAAAAGR20000000000000	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
3327	AGAASPLNSPLSSAVPSLGK200020000000000000	sp 075962 TRIO_HUMAN
3328	YSDGDKSDDLVDVSNEDPATPR2E+22	sp Q04726 TLE3_HUMAN
3329	DDSFDSLDSFGSR200200000000	sp Q9UPQ0 LIMC1_HUMAN
3330	EEGELEDGEISDDNNSQIR200000000000	sp 060293 ZC3H1_HUMAN
3331	AVSEGCASEDEVEGEA2000020000000000	sp Q9BYX2 TBD2A_HUMAN
3332	GGLSPANDTGAK2000000000	sp P30419 NMT1_HUMAN
3333	SVSASSGPCSPAAPHPPPPPR2.00022E+21	sp Q86UY5 FA83A_HUMAN
3334	SGAQASSTPLSPTR20020000	sp P02545 LMNA_HUMAN
		sp 014950 ML12B_HUMAN
3335	ATSNVFMFDQSQIQEFK200001000000000000	sp P19105 ML12A_HUMAN
		sp P24844 MYL9_HUMAN
3336	SRSFDYNYR2020000000	sp 075494 SRS10_HUMAN
3337	YGPADVEDTTGSGATDSKDDDDIDLFGSDDEESEEAKR20000000000000	sp P24534 EF1B_HUMAN
3338	EYVSNDAAQSDDEEKLQSQPTDTDGGR2E+24	sp Q5JSH3 WDR44_HUMAN

3339	IDISPSTFR2000000	sp Q9Y2W1 TR150_HUMAN
3340	NVRSDISDQEDEESEGC	sp Q15154 PCM1_HUMAN
	PVSINLSK2.00200000002E+23	sp P30510 1C14_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
3341	GGSCSQAASSNSAQGSDES	sp Q07000 1C15_HUMAN
	LIACKA20020000000	sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
3342	VVFSDESSEEDLSAEMTEDE	sp Q15021 CND1_HUMAN
	TPKK2.2002200000001E+22	sp Q96QT6 PHF12_HUMAN
3343	TTSPSSDTLLDR200000000000	sp P49321 NASP_HUMAN
3344	EGEETEGSEEDDKENDKTE	sp Q96B36 AKTS1_HUMAN
	EMPNDVLENK2.002E+26	sp Q5T200 ZC3HD_HUMAN
3345	TEARSSDEENGPPSSPDLDR	sp P35269 T2FA_HUMAN
	20000000022000000	sp Q14244 MAP7_HUMAN
3346	LRSPNSAHR2000000000	sp Q16204 CCDC6_HUMAN
3347	IHDLEDDLEMSSDASDASG	sp P46013 KI67_HUMAN
	EEGRVPK1220020020000000000	sp Q5T200 ZC3HD_HUMAN
3348	RLSSSSATLLNSPDR20000000020000	sp Q9P1T7 MDF1C_HUMAN
3349	LDQPVSAPPSPR20002000	sp P07910 HNRPC_HUMAN
3350	IPCDSPQSDPVDTPSTK200000000000000	sp Q69YN4 VIR_HUMAN
3351	STSPAGQHHSPISSR20000002000000	sp O60841 IF2P_HUMAN
3352	IQSSLSVNSDISK200200000	sp Q02880 TOP2B_HUMAN
3353	MESEGGADDSAEGLDLLDD	sp Q9HC35 EMAL4_HUMAN
	DDNEDRGDDQLELIKDDEK2.0000002E+36	sp Q8NDT2 RB15B_HUMAN
3354	SFLSESPSPGR20000	sp O75179 ANR17_HUMAN
3355	WDGSEEDDNSKK20000000000	sp P02545 LMNA_HUMAN
3356	KVVEAVNSDSDSEFGIPK202000000000	sp P30510 1C14_HUMAN
3357	APVSTESVIQSNTPTPPSQ	sp Q29865 1C18_HUMAN
	PLNETAEESR2.2E+28	sp P30504 1C04_HUMAN
3358	RNSLEGYSR20000000	sp Q9TNN7 1C05_HUMAN
3359	EHYPVSSPSPSPAQPGGVS	sp P04222 1C03_HUMAN
	R2000000000000000000000000	sp Q07000 1C15_HUMAN
3360	SGAASSTPLSPTR200020000	sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
3361	GGSCSQAASSNSAQGSDES	sp Q7Z2W4 ZCCHV_HUMAN
	LIACK20020000000	sp Q5T200 ZC3HD_HUMAN
		sp Q8WYL5 SSH1_HUMAN
		sp Q9HCD5 NCOA5_HUMAN
		sp O75475 PSIP1_HUMAN
		sp Q7Z4S6 KI21A_HUMAN
		sp P22059 OSBP1_HUMAN
		sp Q01433 AMPD2_HUMAN
		sp Q9P2B4 CT2NL_HUMAN
		sp P67809 YBOX1_HUMAN
		sp P25054 APC_HUMAN
3362	ASLEDAPVDDLTR2000000000000	
3363	TPSPPPPPIPEDIALGKK2020000000000000000	
3364	SSSSDSIHSVR2000000000	
3365	SSTDSLPGPISR20200000000	
3366	TGVTSTSDSEEGDDQEGEK22000000000000000	
3367	EKELSPPPGLPSK2000000000	
3368	TGSNISGASSDISLDEQYK2000002200200000000	
3369	TDSDDLQLYK2000000000	
3370	FQSQADQDQQASGLQSPPSR200000	
3371	NYQQNYQNSGESGEK2000000	
3372	SGECSVPVPMGSFPR2000200000000000	

3373	SLDLGIIAEVK2000000000000	sp	P08729	K2C7_HUMAN
3374	VDSPSHGLVTSSLCIPSPAR2000000000000000	sp	Q9UER7	DAXX_HUMAN
3375	AGVRPSSSGSAWEACSEAPSK2000000000000000	sp	Q12789	TF3C1_HUMAN
3376	YSPTSPTYSPPTPK300000002000000	sp	P24928	RPB1_HUMAN
3377	KEDSDEEEDDDSEDEEDEDDEDEDEDEIEPAAMK2.00000002E+32	sp	P19338	NUCL_HUMAN
3378	LEDAESSNPNLQSLSTDALPSASK2E+19	sp	Q9Y294	ASF1A_HUMAN
3379	HRSPPATPPPK2000200000	sp	Q8IYB3	SRRM1_HUMAN
3380	KSDGACDSPSSDKENSSQIAQDHQK2.00000202E+24	sp	Q9UIG0	BAZ1B_HUMAN
3381	GQKPALDCSQVSPRPATSPENNASLSDTSPMDSPPSGIPK20000000000000000000	sp	P18858	DNLI1_HUMAN
3382	SSSSGSDDYAYTQALLLHQR30000000000000	sp	Q8N5C8	TAB3_HUMAN
3383	SVSPCSNVESR202000000000	sp	Q9UQ35	SRRM2_HUMAN
3384	LMHNASDSEVDQDDVVEWK10002020000000000000	sp	Q9H2P0	ADNP_HUMAN
3385	AGMSSNQSISSPVLDAVPR10200000200000000000	sp	Q9UQ35	SRRM2_HUMAN
3386	EFVSSDESSGENK220002000000	sp	Q08945	SSRP1_HUMAN
3387	LPLQESEEEER2000000	sp	P23497	SP100_HUMAN
3388	EILSPVDIIDR200000000	sp	O14495	LPP3_HUMAN
3389	SNSPLPVPPSK202000000000	sp	Q13247	SRSF6_HUMAN
3390	GLMAGGRPEGQYSEDEDTDDEYK20000200000000	sp	Q9NPQ8	RIC8A_HUMAN
3391	STPESGSDKESVGSSTSNEGGR2.002E+23	sp	Q6GQQ9	OTU7B_HUMAN
3392	LQLSPWESSVNR20000000000	sp	Q14244	MAP7_HUMAN
3393	SFAGNLNTYK20000000000	sp	Q01813	PFKAP_HUMAN
3394	RAGDLEDSPK2000	sp	P51858	HDFG_HUMAN
3395	TGSPGSPGAGGVQSTAK2000000000000	sp	O43237	DC1L2_HUMAN
3396	GPPDFSSDEEREPTPVLGSGAAAAGR2.00000002E+21	sp	P42166	LAP2A_HUMAN
3397	YLVGDKPNAGSEEISSEDELVEEK3E+26	sp	P42167	LAP2B_HUMAN
3398	KTAVQYIESSDSEEIETSELPQK2202000000000000	sp	Q9NY61	AATF_HUMAN
3399	SDSQASLTVPLSPQVVR2000000002000000	sp	Q14527	HLTF_HUMAN
3400	SHSPSASQSGQLR2000000000000	sp	Q8TEV9	SMCR8_HUMAN
3401	SGGSGHAVAEPASPEQELDQNK2.00000002E+19	sp	Q8WWI1	LMO7_HUMAN
3402	AAGDSGEDSDSDYENLIDPTEPSNSEYSHSK2.00203E+24	sp	P46013	KI67_HUMAN
3403	RPSSTSVPLGDK20000000000	sp	Q15018	F175B_HUMAN
3404	LEGSDDLLEDSSEEHSR202000000	sp	Q9NXR1	NDE1_HUMAN
3405	GHSSLTNSPLDSSCK200000000	sp	Q9NR09	BIRC6_HUMAN
3406	SHSDASVGSSTESEHGSSSPR2000	sp	Q9Y6Q9	NCOA3_HUMAN
3407	AESPAEKVPEESVPLVQK20000000000000000000	sp	Q6XZF7	DNMBP_HUMAN
3408	AKEVELVSE200	sp	Q9UKV3	ACINU_HUMAN
3409	TSPLKDNPSPEPQLDDIKR20000002000000000000	sp	O60832	DKC1_HUMAN
3410	SIDTQTPSVQER2000020000000	sp	Q96JC9	EAF1_HUMAN
3411	RLSQPESAEK200000000	sp	Q86VQ1	GLCI1_HUMAN
3412	SKFDSDEEEDTENVEAASSGKVTR2E+25	sp	Q9UKV3	ACINU_HUMAN
3413	EGMNPSYDEYADSDQHDAYLER1.0003000002E+22	sp	Q8TF01	PNISR_HUMAN
3414	DVQSPGFLNEPLSSK2000000000000	sp	Q08945	SSRP1_HUMAN
3415	SASSDTSEELNSQDSPPK20200000000000000000	sp	Q8NG31	CASC5_HUMAN
3416	SPVQLEEAPFSR20000000000000	sp	O14745	NHRF1_HUMAN
3417	RLSDCESTDVK20000000000	sp	Q9BZL4	PP12C_HUMAN
3418	SSSLDGEHHDGYHR2000000000000	sp	Q7Z333	SETX_HUMAN
3419	SSSPGKPAVSSLNSSHRSR2E+19	sp	Q9C0J8	WDR33_HUMAN
3420	QGSITSPQANEQSVTPQR20020000000020000	sp	Q9UHB7	AFF4_HUMAN
3421	SPGNTSQPPAFFSK2000000000000000	sp	Q9UQ35	SRRM2_HUMAN
3422	IAQVSPGPR200000	sp	Q9Y618	NCOR2_HUMAN
3423	LTGSTSSLNK20000000	sp	Q9NQS7	INCE_HUMAN
3424	KSSISSISGR200000000	sp	Q9NV70	EXOC1_HUMAN
		sp	Q86SQ0	PHLB2_HUMAN



3425	SPEAVGSECPSR2000000000000	sp Q9Y4K1 AIM1_HUMAN
3426	EEDEEPESPPEKK2000000	sp Q9H1E3 NUCKS_HUMAN
3427	LLKPGEEPSEYTDEEDTK20000000000	sp O15173 PGR2_HUMAN
3428	GTSRPPPEGGLGYSQLGDDDLK2. 2E+21	sp P21127 CD11B_HUMAN
		sp Q9UQ88 CD11A_HUMAN
3429	GLNSQSSDHLNKR200200000000	sp Q96HA1 P121A_HUMAN
		sp A8CG34 P121C_HUMAN
3430	QISLPDLSQEEPQLK2000000000000	sp Q15390 MTFR1_HUMAN
3431	SRPTSEGSDIESTEPQK2000000000000000000	sp Q4G0J3 LARP7_HUMAN
3432	SGSSPDSEITELKFPSINHD20000000000000000	sp P17812 PYRG1_HUMAN
3433	LLDLSdstsvasfPSADETDGNLPESR2. 002E+19	sp Q13464 ROCK1_HUMAN
3434	NLEHLSSFSDEDDPGYSQDAYK2002000000000000000	sp Q2KHR3 QSER1_HUMAN
3435	AGMSSNQSISSPVLDAVPR20000020000000000	sp Q9UQ35 SRRM2_HUMAN
3436	TASFSESR2000000	sp P53396 ACLY_HUMAN
3437	STSAPQMSPGSSDNQSSSPQPAQQK2. 0001E+23	sp Q14157 UBP2L_HUMAN
3438	SPGLVPPSPEFAPR200000020000000	sp Q9H5H4 ZN768_HUMAN
3439	SPQPDPVGTPTIFKPKQSK20000000200000000000	sp P46013 KI67_HUMAN
3440	SASLYPDR20000000	sp Q6P4Q7 CNNM4_HUMAN
3441	LDSSPVSSPR20000000	sp O60293 ZC3H1_HUMAN
3442	RGSGSACSLCCCGR200000000000000	sp Q9GZU1 MCLN1_HUMAN
3443	AVATAAAQQTGPEEDSGSSEEEESDSEEEAETLAQVKPSGK2. 022000202E+25	sp Q13428 TCOF_HUMAN
3444	QLPALDGSLMGPEPPAQEEEEAPVSPHK2E+21	sp Q8N9T8 KRI1_HUMAN
3445	DHSPTPSVFNSDEER2000000000	sp Q6UN15 FIP1_HUMAN
3446	DTVIIVSEPEDEESQGLPTMAR200200000000001000	sp Q9H4L7 SMRCD_HUMAN
3447	DEVFHTDQDDPSSSDDEGMPYTRPVK220000000000000	sp Q5JSH3 WDR44_HUMAN
3448	SASEPSLHR20000000	sp P10398 ARAF_HUMAN
		sp P04049 RAF1_HUMAN
3449	ESTSSKSPPR20000	sp P78347 GTF2I_HUMAN
3450	ELDDSSSESSDLQLEGPSTLR2022000000000000000	sp P29590 PML_HUMAN
3451	SQSLPGADSLAKPIDK2000000000000000000	sp Q9Y4A5 TRRAP_HUMAN
3452	DHASQLSPVLSR2000000	sp Q8IXZ2 ZC3H3_HUMAN
3453	QNSVQEQPGTACLK2000000000000000000	sp Q9NQW6 ANLN_HUMAN
3454	TASGSSVTSLDGTR22020000000	sp Q92597 NDRG1_HUMAN
3455	SPGAGSLGSPASQR2000000	sp P53667 LIMK1_HUMAN
3456	SEGSEYEEIPK2000000000000	sp P48634 PRC2A_HUMAN
3457	IDISPSTLR2000000	sp Q9NYF8 BCLF1_HUMAN
3458	YSPTSPTYSPVYTPSPK2020000000000000000	sp P24928 RPB1_HUMAN
3459	SRSRSTPPSSAGLGSNSAPPIDSR2. 0202020022E+26	sp Q8IWX8 CHERP_HUMAN
3460	ASPITNDGEDEFVPSDGLDKDEYTFSPGK2. 0000000000002E+28	sp Q02880 TOP2B_HUMAN
3461	VGVSSKPDSSPVLSPGNK20000000000	sp Q8N4C8 MINK1_HUMAN
3462	TIQNSSVSPTSSSSSSSSTGETQTQSSSR2E+25	sp Q7Z417 NUFP2_HUMAN
3463	SPSPAPAPAAAAAGPPTRK2. 02E+20	sp Q9H7N4 SFR19_HUMAN
3464	SPEPGQWTR20000000000	sp Q09666 AHNK_HUMAN
3465	RTDALTSPPGR200000	sp P49736 MCM2_HUMAN
3466	SSPGPDTLVVLEFNASK200000000000000000000	sp O60303 K0556_HUMAN
3467	KLSSSEPYEEDEFNDDQSIK2. 002E+19	sp Q9H4L7 SMRCD_HUMAN
3468	TASISSPSEGTPVGSYGCTPQSLPK2. 22E+23	sp Q6PKG0 LARP1_HUMAN
3469	IACKSPQDPVDTASTK2000000000000000000	sp P46013 KI67_HUMAN
3470	SHSTEPNLSSFLNDPNPMK2000000000000000000	sp Q9UPQ0 LIMC1_HUMAN
3471	SRSSPVTELASR202220000000000	sp Q9UQ35 SRRM2_HUMAN
3472	VELSPGPPKAGR200000000000	sp Q8IZ73 RUSD2_HUMAN
3473	GESAPTLSTSPSSPSPTSPSPTLGR202000000	sp Q9H330 TM245_HUMAN
3474	KESESDSSDEPLIK2020022000000000	sp P35659 DEK_HUMAN

3475	LSSTDDGYIDLQFK2020000000000	sp Q96A57 TM230_HUMAN
3476	SPHQLLSPSSFPSATPSQK2.00000202E+20	sp Q14181 DPOA2_HUMAN
3477	HIVSNDSSDDESHEPK2002000000000	sp P47712 PA24A_HUMAN
3478	RNSSEASSGDFLDLK2200000000000	sp Q9UK76 HN1_HUMAN
3479	SQSIEQESQEK2020000000000	sp Q5VTL8 PR38B_HUMAN
3480	LHVGNISPTCTNK20000000	sp Q9BWF3 RBM4_HUMAN
3481	RPSQEQSASASSGQPAPLNR2000000000000	sp Q5T1M5 FKB15_HUMAN
3482	VKEPPSPQSPR20002000	sp Q00613 HSF1_HUMAN
3483	SASEPSLHR20000000	sp P04049 RAF1_HUMAN
		sp P10398 ARAF_HUMAN
3484	TIASDSEEEAGKELSDKK202000000000000	sp Q96ST2 IWS1_HUMAN
3485	AAEETPGKSPEK20000	sp Q76021 RL1D1_HUMAN
3486	KGSDDAPYSPTAR200000002000	sp Q75151 PHF2_HUMAN
3487	QFTSSSIK2000	sp Q04695 K1C17_HUMAN
3488	HNSPSGHLTHSR20000000200	sp Q8IWB9 TEX2_HUMAN
3489	SSSEAKPTSLGLAGGHK200000000000000	sp Q96RK0 CIC_HUMAN
3490	HFFHHADEDEEEDDPPER200000	sp Q43314 VIP2_HUMAN
3491	KTSLSDLNDPIILSSDDDDNDR2.00000000022E+21	sp Q9GZR1 SENP6_HUMAN
3492	TPTSSPASSPLVAK200002000000	sp Q14684 RRP1B_HUMAN
3493	IACEEFSSEEGEGGRK2020000000000	sp Q13547 HDAC1_HUMAN
3494	AFKPEETSSNSDPPSPVLNNSHPVPR2000000000000	sp Q60293 ZC3H1_HUMAN
3495	SSSFSELCHR200000000	sp Q5T8I3 F102B_HUMAN
3496	SASSGAEGDVSSEREP20000000200000	sp Q8TEA8 DTD1_HUMAN
3497	NSSSPVPASVPGQR2002000000000	sp Q9H1B7 I2BPL_HUMAN
3498	SKTPPPPPQTAQTK2000000000000	sp P25054 APC_HUMAN
3499	NSPNNISGISNPPGTPR20000000000002000	sp Q9BWW4 SSBP3_HUMAN
3500	LSMEDSKSPPK200000	sp Q9NQG5 RPR1B_HUMAN
3501	EVSSRPSTPGLSVVSGISATSEIPNKIEDLR2.2E+30	sp Q14C86 GAPD1_HUMAN
3502	TDHGAEIVYKSPVSGDTSR20022000	sp TAU_HUMAN
		sp P10636 TAU_HUMAN
3503	TGSESSQTGTSTTSSR202000000000000	sp P23588 IF4B_HUMAN
3504	GVNTFSPEGR200000	sp P28066 PSA5_HUMAN
3505	NQSPVLEPVGR2000000000	sp P51812 KS6A3_HUMAN
3506	SPSPTTPGSR2020000000000	sp P49450 CENPA_HUMAN
3507	SLSPGKENSALDMEK2000000000001000	sp Q9H307 PININ_HUMAN
3508	TDDEDFVPSDASPPK2000000020020000	sp P11388 TOP2A_HUMAN
3509	SGYSSPGSPGTPGSR200000000	sp TAU_HUMAN
		sp P10636 TAU_HUMAN
3510	GNSRPGTPSAEGGSTSTLR200000000000000	sp P35269 T2FA_HUMAN
3511	GTPSQSPVGR2000000	sp P52594 AGFG1_HUMAN
3512	LLEDGEDFNLGDALDSSNSMQTIQK2001000000	sp P05783 K1C18_HUMAN
3513	RESPSPAPKPR2000000000	sp Q8IYB3 SRRM1_HUMAN
3514	TDHGAEIVYKSPVSGDTSR200020000000	sp P10636 TAU_HUMAN
		sp TAU_HUMAN
3515	HIVSNDSSDDESHEPK2200000000000	sp P47712 PA24A_HUMAN
3516	IQQGEEEEKESSGEIEAAPVTGTGR220000000000000	sp Q6AI08 HEAT6_HUMAN
3517	QEAGDSPPPAPGTPK20000002000	sp P49848 TAF6_HUMAN
3518	SHVEDGDIAGAPASSPEAPAEQDPVQLK2200000000000000	sp Q9P2E9 RRBP1_HUMAN
3519	SPQLSDFGLER200000000000	sp Q8IX90 SKA3_HUMAN
3520	RNSSEASSGDFLDLK2000000000	sp Q9UK76 HN1_HUMAN
3521	EPESPASAPIPLFGADTIGQR2E+19	sp Q96AY4 TTC28_HUMAN
3522	SPSLDNPTFPNLGSENPLK2E+19	sp Q14653 IRF3_HUMAN
3523	VTNDISPESPGVGR22000000	sp Q15154 PCM1_HUMAN

3524	TDHGAEIVYKSPVVSGDTSR200020002000	sp P10636 TAU_HUMAN
		sp TAU_HUMAN
3525	SLSGSSPCPK20000200000	sp Q9UQ35 SRRM2_HUMAN
3526	FASDDEHDEHDENGATGPVKR2E+19	sp P05455 LA_HUMAN
3527	AAGDSGEDSDSDYENLIDPTEPSNSEYSHSK2.000200003E+28	sp Q15018 F175B_HUMAN
3528	SASSESEAEENLEAQPQSTVRPEEIPPIPENR2.202E+29	sp Q13427 PPIG_HUMAN
3529	GLVAAYSGESDSEEEQER2000000000	sp P98175 RBM10_HUMAN
3530	SLSPGKENVSALDMEK20000000001000	sp Q9H307 PININ_HUMAN
3531	NQVEEDAEDSGEADDEEKPEIHKPGQNSFSK2E+22	sp 075717 WDHD1_HUMAN
3532	RLSTSPDVIQGHQPR20000000000000	sp Q9Y385 UB2J1_HUMAN
3533	GSGSQSSVPSVDQFTGVGIR2000000000000000	sp Q9UQE7 SMC3_HUMAN
3534	SPGSTPTTPTSSQAPQK200000000000000000	sp P35658 NU214_HUMAN
3535	KSPSGPVKSPPLSPVGTTPVK20000200000000000000	sp Q9BVC5 ASHWN_HUMAN
3536	SRTASGSSVTSLDGTR202020000000000000	sp Q92597 NDRG1_HUMAN
3537	NSCQDSEADEETSPGFDEQEDGSSSQTANKPSR2.000002E+28	sp Q86W56 PARG_HUMAN
3538	EDLSPAFDHSPNK20000	sp P17706 PTN2_HUMAN
3539	SQSSHSHYDDSTLPLIDR20000200000000000000	sp 060716 CTND1_HUMAN
3540	RRSEVESTTESQDK2000000000000000	sp Q5UIP0 RIF1_HUMAN
3541	VLSEPIGPESPSSTECPR2000000000	sp Q56N19 ESCO2_HUMAN
3542	SDSASSEPVGIYQGFQK200000000000000000	sp Q05655 KPCD_HUMAN
3543	LQDISSELLATGAGLSESEAEAPDGDHNITELPQAVAGR2.0000002E+28	sp Q9NQ55 SSF1_HUMAN
3544	SASSESEAEENLEAQPQSTVRPEEIPPIPENR2.02202E+31	sp Q13427 PPIG_HUMAN
3545	THTTALAGRSPSPASGR202000000	sp Q9UQ35 SRRM2_HUMAN
3546	SSLGQSASETEEDTVSVSK2.20002E+19	sp Q13435 SF3B2_HUMAN
3547	ERSPSPTTIQSK20200000000	sp Q14687 GSE1_HUMAN
3548	EQLSETEGSGEDEPGNDPSETTQK2.00002E+21	sp Q13107 UBP4_HUMAN
3549	SLSPVAAPPLR202000000000	sp Q8NDT2 RB15B_HUMAN
3550	NSSLGSPSNLCGSPPGSIR20000000000200000000	sp Q9C0B0 UNK_HUMAN
3551	NVPQEESELESDVDADFK200000000	sp 000505 IMA4_HUMAN
3552	RNSSSPVSPASVPGQR2020020000000000	sp Q9H1B7 I2BPL_HUMAN
3553	GRSPQPPAEEDDDFDDLVAIDTYNCDLHFK200000000000000000	sp Q9BUJ2 HNRL1_HUMAN
3554	TTYDSAEENKENLYAGK2000000000000000	sp Q9HAW4 CLSPN_HUMAN
3555	GLPSPYNMSSAPGSR102000200	sp P15924 DESP_HUMAN
3556	QSETVDQNSDSDEMLAILK200000000000	sp P46100 ATRX_HUMAN
3557	RDSSSQLASTESDKPTTGR200000000000000000	sp Q96B23 CRO25_HUMAN
3558	SFTLDDESLK200000000000	sp Q86WR7 PRSR2_HUMAN
3559	SLSPQEDALTGSR200000000000	sp Q96EN8 MOCOS_HUMAN
3560	LKSVEDEMDSPGEEPFTYTGQR2E+20	sp Q12857 NFIA_HUMAN
3561	SESDLSQPESDEEGYALSGR20000002000000000000	sp Q15751 HERC1_HUMAN
3562	STLESEKPGSPEAAETSPPSNIIDHCEK2.000002E+19	sp Q96T23 RSF1_HUMAN
3563	EVDPSTGELQSLQMPESGSSLDPSQEGPTGLK2000000000000000	sp Q13501 SQSTM_HUMAN
3564	DKSDSPAQLR20000000	sp Q96ST3 SIN3A_HUMAN
3565	ALPSLNTGSSSPR2000	sp 014545 TRAD1_HUMAN
3566	ESSAAQAGSHATHPGTSVLEGGAGSMSPSR20000	sp Q5SYE7 NHSL1_HUMAN
3567	ASLGSLEGEAEAEASSPK2000000000000000	sp Q09666 AHNK_HUMAN
3568	VFIDQNLSPGK20000	sp Q14186 TFDP1_HUMAN
3569	SGSSPDSEITELKFPSINHD2.0222E+21	sp P17812 PYRG1_HUMAN
3570	VISDESSEDIGSDVEFKPDTK20200000000000000000	sp P52701 MSH6_HUMAN
3571	VADPDHDHTGFLTEYVATR300000	sp P28482 MKO1_HUMAN
3572	SPGAPGPLTLK20000000000000	sp Q15942 ZYG_HUMAN
3573	SDSLLSFR2002000	sp Q00587 BORG5_HUMAN
3574	LESSEGEIIQTVDR22000000000000	sp Q99590 SCAFB_HUMAN
3575	TASESISNLSEAGSIK20000000020002000	sp P30622 CLIP1_HUMAN
3576	HDTVTVSSDLQFTK2000000000	sp P30414 NKTR_HUMAN

3577	TTCMSSQGSDDDEQIKR200000000	sp Q9P0V9 SEP10_HUMAN
3578	SDHGAYSQSPAIK20000000000000	sp Q9Y4B6 VPRBP_HUMAN
3579	RGSPSAAFTFPDPTDDDFGK2000000000000000	sp Q9ULT0 TTC7A_HUMAN
3580	SDGAPASDSKPGSSEAAPSSK20002000000000	sp P80723 BASP1_HUMAN
3581	EEDCHSPTS KPPKPDQLK2000000000000000	sp P23588 IF4B_HUMAN
3582	GHTDTEGRPPSPPTSTPEK20000020000	sp Q00613 HSF1_HUMAN
3583	EGEPTVYSDEEEK20000000	sp 000264 PGRC1_HUMAN
3584	KVEEDLKADEPSSEESDLEIDK200020000000	sp P50502 F10A1_HUMAN
3585	NYQQNYQNSESGEKNESESAPEGQAQR2.000000002E+21	sp P67809 YBOX1_HUMAN
3586	SKPPPTYESEEEK2000000	sp 060885 BRD4_HUMAN
3587	HSHSHSPMSTR20002000000	sp P62995 TRA2B_HUMAN
3588	AQSYPDNHQEFSDYDNPIFEK2E+19	sp Q9Y2U5 M3K2_HUMAN
3589	RGSDASDFDISEIQSVCSDVETVPQTHRPTPR2E+30	sp Q03001 DYST_HUMAN
3590	DSSKGEDSAEETEAKPAVVAPVPVEAVSTPSAAFPSDATAEQGPILTK2E+38	sp Q12906 ILF3_HUMAN
3591	RLPQDHADSCVVSSDDEELSR2200000000	sp P78317 RNF4_HUMAN
3592	STAPSAASASASAAASSPAGGGAEELEHCGVCR2E+36	sp Q13263 TIF1B_HUMAN
3593	SQTPSPSTLNIDHMEQK202020000000000000	sp Q6GYQ0 RGPA1_HUMAN
3594	AALAPAKESPR2000	sp Q13428 TCOF_HUMAN
3595	ELDPSLVSANDSPSGMQTR200000000	sp Q5UIP0 RIF1_HUMAN
3596	TPLTSADEHVHVK20002000000000	sp Q5SW79 CE170_HUMAN
3597	LSPDFAEELR2000000000	sp Q14160 SCRIB_HUMAN
3598	VASETHSEGSEYEELPK20200000000000	sp Q5JSZ5 PRC2B_HUMAN
3599	KPPAPSPVQSQSPSTNWSPAVPVKK2020000000000000	sp Q8IYB3 SRRM1_HUMAN
3600	AGMSSNQSISSPVLDAVPR20000022000000000	sp Q9UQ35 SRRM2_HUMAN
3601	ESSEPPAPASSPEAASPTTEQGPAGTSK220000000000000000	sp Q92766 RREB1_HUMAN
3602	LSSQISAGEEK2000000	sp Q86UE4 LYRIC_HUMAN
3603	APSVATVGSICDLNLK2000000000000000	sp 075369 FLNB_HUMAN
3604	ILSDVTHSAVFGVPASK2000000000000000	sp Q2M2I8 AAK1_HUMAN
3605	DDSHAEDSEDEKEDHK2000000000	sp Q9H1E3 NUCKS_HUMAN
3606	AADPPAENSSAPEAEQGGAE200000000000	sp P67809 YBOX1_HUMAN
3607	TAGSSVTSLDGTR20000000000	sp Q92597 NDRG1_HUMAN
3608	AVLSDSPQLSEK2000000000	sp Q7Z4H7 HAUS6_HUMAN
3609	KPSLVDLPWEGAAPQSPSFGSEDSGSPK2020000000000000	sp Q13425 SNTB2_HUMAN
3610	KPPAPSPVQSQSPSTNWSPAVPVK2020200000000000	sp Q8IYB3 SRRM1_HUMAN
3611	SLTNSHLEK2000000000	sp Q9H2H9 S38A1_HUMAN
3612	QTPSPDVVLR20000000	sp Q9UPQ0 LIMC1_HUMAN
3613	DQSPPPSPPSYHPPPPPTK20002000000000000000	sp Q9P206 K1522_HUMAN
3614	DMAECSTPLPEDCSPHSPR20000000	sp Q12802 AKP13_HUMAN
3615	TPVQYSQQQNSPK20000	sp P46013 KI67_HUMAN
3616	ASPGTPLSPGSLR2002000	sp Q96BD0 SO4A1_HUMAN
3617	EHYVSSPSSPPAQP GGVS R202200000000000000	sp 075179 ANR17_HUMAN
3618	SLGYHGSSYSPGVEPVSPVSSPLTHDK2002000000000	sp Q9Y618 NCOR2_HUMAN
3619	EVDATSPAPSTSSTVK20000000000000	sp Q16666 IF16_HUMAN
3620	DGDSYDPYDFSDTEEMPQVHTPK2020000000000000	sp P25205 MCM3_HUMAN
3621	EPGSPSVQR20000000	sp Q9H6F5 CCD86_HUMAN
3622	SLGSASPGGQPPLSSPTR20000000000200000	sp Q9C0B5 ZDHC5_HUMAN
3623	SGSSSPDSEITELKFPSINH2.0222002E+21	sp P17812 PYRG1_HUMAN
3624	VIGQDHDFSESSEEAPEASSGALR220000000000000000	sp P23497 SP100_HUMAN
3625	MESEGGADDAEGLDLDLDDNEDRGDDQLELIKDDEK2E+29	sp P07910 HNRPC_HUMAN
3626	EEGRCSSDDDTDVDMEGLR22200020000000000	sp Q96AQ6 PBIP1_HUMAN
3627	GNIETSEDEDDTADLFPEESGFEHAWR2.22E+24	sp Q9H501 ESF1_HUMAN
3628	RFSDSEGEETVPEPR2000000000000000	sp Q13286 CLN3_HUMAN
3629	RKPSVPDASPADDSFVDPGER2E+19	sp P16333 NCK1_HUMAN

3630	MLPHAPGVQMQAIPEDAIPEESGDEDEDDPKR200000000000	sp Q13547 HDAC1_HUMAN
3631	DGTAPPPQSPGSPGTGQDEEWSDEESPR2E+26	sp Q9UKS6 PACN3_HUMAN
3632	SSLSSHSHSQSIYR220000000000	sp O15027 SC16A_HUMAN
3633	ELSDQATASPIVAR2000000	sp Q5JSH3 WDR44_HUMAN
3634	HIKEEPLSEEEPTSTAIASPEKK200000000000200000	sp Q9Y2X3 NOP58_HUMAN
3635	RGSDIDNPTLTVMDISPPSR2000000000000000000	sp Q9Y2U5 M3K2_HUMAN
3636	ATEDGEEDEVSAGEK200000	sp Q9BXW9 FACD2_HUMAN
3637	SQSLFLPTEAAQVSPPPYPAPQELTQPLLQQPR2E+21	sp Q6UUV7 CRTC3_HUMAN
3638	SLYESFVSSDR2000000000000	sp P18615 NELFE_HUMAN
3639	GAGGSPVGVVEGLVNVGTGQK2000000000000000000	sp Q7Z5J4 RAI1_HUMAN
3640	VETLSPESALQGSFR200000002000	sp Q99958 FOXC2_HUMAN
3641	LAGGQTSQPTTPLTSPQR200000	sp O95155 UBE4B_HUMAN
3642	ATSSSNPSSPAPDWYK2000002000000000	sp P46100 ATRX_HUMAN
3643	AQTPPISSLPTSPSDEVGR220020000000000	sp Q15751 HERC1_HUMAN
3644	EGETQGVAFEHESPADFQNSQSPVQDQDK200000000	sp P78332 RBM6_HUMAN
3645	ALSLGTIPSLTR200000000000	sp Q5SNT2 TM201_HUMAN
3646	LENSPLGEALR200000000	sp Q9NX40 OCAD1_HUMAN
3647	GDAASSPAPAASVGSQGGAR20000000020000000	sp Q8WVB6 CTF18_HUMAN
3648	EESDEEEEDDEESGR200000000000000	sp Q96EV2 RBM33_HUMAN
3649	GPVSGSPDSMNASR2000000000	sp O14920 IKKB_HUMAN
3650	KEVLASPR20000	sp Q9C0C2 TB182_HUMAN
3651	GAGATSGSPAGR2000000	sp P16455 MGMT_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
		sp Q29960 1C16_HUMAN
3652	GGSCSQAASSNSAQGSDESLIACK2002000000	sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
3653	VEHSPGPPVVDK2000000000	sp Q9H869 YYAP1_HUMAN
3654	SFQGDDSDLLK2000000000000	sp Q9UPQ0 LIMC1_HUMAN
3655	RSSQPSPTAVPASDSPPTK2000000000000000000	sp Q3KQU3 MA7D1_HUMAN
3656	QPPGVPNGPSSPTNESAPELPQR200000000000000	sp Q8TF74 WIPF2_HUMAN
3657	DSSTSYTETKDPSSGQEVATPPVQLQVCEPK2. 2E+30	sp Q9UKV3 ACINU_HUMAN
3658	KLEDVKNSPTFK200000	sp P55327 TPD52_HUMAN
3659	RKPSVPDSASPADDSFVDPGER200000000000000	sp P16333 NCK1_HUMAN
3660	APSPPTASNSNSQSEK2000000000000000000	sp Q8N108 MIER1_HUMAN
3661	WESQQDVSQTTVSR2000000000000	sp Q9NQT8 KI13B_HUMAN
3662	GRLTPSPDIVLSDNEASSPR2020000000000000000	sp Q8WXI9 P66B_HUMAN
3663	SPSPASTSTVPNMTDAPTAPK2. 02E+21	sp Q659A1 ICE2_HUMAN
3664	ASPPPAHAHEHGGVLAR2020000000000000000	sp O75385 ULK1_HUMAN
3665	SLPTPAVLLSPTK20000	sp Q8WX93 PALLD_HUMAN
3666	GGGSCSGSAGGGGSLPSQR200000000	sp O15234 CASC3_HUMAN
3667	SPHQLLSPSSFSPSATPSQK2. 000002E+20	sp Q14181 DPOA2_HUMAN
3668	LDGLVETPTGYIESLPR200000000000	sp P55209 NP1L1_HUMAN
3669	SPSLSLAPLSPR20000002000	sp Q969H4 CNKR1_HUMAN
3670	VQSGMGLILQGYESGSEEEGEIHEK1. 0000000202E+21	sp Q13523 PRP4B_HUMAN
3671	SLPTTVPESPNYR200000000000000	sp Q6PKG0 LARP1_HUMAN
3672	GYSPYSVSGSGSTAGSR20020000000000000	sp Q15149 PLEC_HUMAN
3673	SPSLLQSGAK20000000000	sp Q9BQG0 MBB1A_HUMAN

3674	SKGSDKDFVIVR2002000000000	sp Q13309 SKP2_HUMAN
3675	SQSSDTEQQSPTSGGGK2000000000000	sp P23588 IF4B_HUMAN
		sp P67809 YBOX1_HUMAN
3676	SVGDGETVEFDVVEGEK20000000000000000	sp P16989 YBOX3_HUMAN
		sp Q9Y2T7 YBOX2_HUMAN
3677	SGSSPDSEITELKFPSINH2. 02E+21	sp P17812 PYRG1_HUMAN
3678	NHSPSPVPTGAAPSLASPK2. 02E+19	sp Q5SYE7 NHSL1_HUMAN
3679	SPAGPAATPAQAQAASPR2000	sp Q13428 TCOF_HUMAN
3680	SNSVGIQDAFNDGSDSTFQK20000000000000000	sp O14497 ARI1A_HUMAN
3681	HGAPAAPSPPPR200000	sp Q4KMP7 TB10B_HUMAN
		sp P42166 LAP2A_HUMAN
3682	SSTPLPTISSAENTR20000000000000000	sp P42167 LAP2B_HUMAN
		sp Q13247 SRSF6_HUMAN
3683	ARVSPPPKR202000000	sp Q9Y613 FHOD1_HUMAN
3684	SLEGGGCPAR200000000000	sp Q13425 SNTB2_HUMAN
3685	KPSLVSDLPWEGAAPQSPSFGSEDSGSPK20000000000000000	sp Q02952 AKA12_HUMAN
3686	SAESPTSPVTSETGSTFK20000000000000000	sp Q15276 RABE1_HUMAN
3687	AQSTDSLGTSGSLQSK20200000000000000	sp Q9Y2X7 GIT1_HUMAN
3688	HGSGADSDYENTQSGDPLLGLEGK2. 000000002E+22	sp Q71F23 CENPU_HUMAN
3689	LRPISDDSEIEESDTR200202000000000	sp P29401 TKT_HUMAN
3690	ILATPPQEDAPSVDIANIR20000000000000000	sp Q9C0B5 ZDHC5_HUMAN
3691	SLGSASPGPGQPPLSSPTR200000000200000	sp Q6PJF5 RHDF2_HUMAN
3692	QASLSQSIR200000000	sp Q9BVS4 RIOK2_HUMAN
3693	SFEMTEFNQALEEIK20010000000000000	sp Q9NQC1 JADE2_HUMAN
3694	KYISSDNSDTTDSHATSTSASR2. 022E+21	sp Q7Z6B7 SRGP1_HUMAN
3695	STSSSDTMSTFKPMVAPR22000000000000000	sp P23588 IF4B_HUMAN
3696	SLENETLNK20000000000	sp P30510 1C14_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
3697	GGSCSQAASSNSAQGSDESLIACK2002000000	sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
3698	LQSIGTENTEENR2000000000000	sp ALDOA_RABIT
		sp P04075 ALDOA_HUMAN
3699	SKSPLPPEEEAK200000000000	sp Q1KMD3 HNRL2_HUMAN
3700	SSPPLRTPDVLESSGPAVR200002000000000000000	sp O75676 KS6A4_HUMAN
		sp P55795 HNRH2_HUMAN
3701	HTGPNSPDTANDGFVR2000000000000	sp P31943 HNRH1_HUMAN
		sp Q9Y2F5 ICE1_HUMAN
3702	VPGEDGTLPPTQGSPLR20000	sp P36915 GNL1_HUMAN
3703	EEQTDTSDGESVTHHIR20000000000000000	sp Q92922 SMRC1_HUMAN
3704	KHSPSPPTPTESR202000020000000	sp Q9HB20 PKHA3_HUMAN
3705	TYSDTDSCSDIPLDPDRPVHCSK2. 0002E+22	sp Q8IY81 SPB1_HUMAN
3706	TEIMSPLYQDEAPK2000000000000	sp Q7Z589 EMSY_HUMAN
3707	TNSSSSSPVVLK20200000000000000	sp P28290 SSFA2_HUMAN
3708	TGSVQTPPDLESSEEVDAEAGAPEVVGPK2. 0020000002E+27	sp P33527 MRP1_HUMAN
3709	QLSSSSYSGDISR2220000000000	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
3710	SSTPLPTISSAENTR200000000000000200	sp Q8IYB3 SRRM1_HUMAN
3711	KPPAPSPVQSQSPSTNWSPAVPVK2. 000200020002E+19	

3712	EAARSPDKPGGSPSASR20000002000000	sp Q9Y3T9 NOC2L_HUMAN
3713	SQLQALHIGLDSSSIGSGPGDAEADDGFPESR2.2E+20	sp O75116 ROCK2_HUMAN
3714	LDSTQLQLHNQVLEQHQQLNPPSSPK2000	sp Q86TC9 MYPN_HUMAN
3715	GAAAADLLSSSPESQHGGTQSPGGGQPLLQPTK2E+25	sp Q9H3S7 PTN23_HUMAN
3716	SRSRSPTPSSAGLGSNSAPPIDSR2.020202002E+26	sp Q8IWX8 CHERP_HUMAN
3717	CAPSAGSPAAAVGRESGAAATSSSGPQAQQHR2.000000002E+27	sp Q7L2J0 MEPCE_HUMAN
3718	WLDESDAEMELR200010000	sp Q9P035 HACD3_HUMAN
3719	ATSPSSVSGDFDDGHHSVSTPGPSR2.02E+24	sp Q86U86 PB1_HUMAN
3720	VVSSVSSSPR2000	sp P29375 KDM5A_HUMAN
3721	TLSDSELCSGR2000000000	sp O60292 SI1L3_HUMAN
3722	VPQALNFSPEESDSTFSK200000000000	sp Q9NYZ3 GTSE1_HUMAN
3723	SPDKPGGSPSASR20000000000000	sp Q9Y3T9 NOC2L_HUMAN
3724	VSITSEVESTSNPSSSSLQK2000000000	sp Q04656 ATP7A_HUMAN
3725	GNAEGSSDEEGKLVIDEPAKEK220000000000000000	sp P51858 HDGF_HUMAN
3726	DSDQVAQSDGEESPAAEQLLGEHIK2E+19	sp O94979 SC31A_HUMAN
3727	TENGTCPSPPLSPAAALGSGSAAAVPK2.000000002E+25	sp Q12948 FOXC1_HUMAN
3728	LSPPVASGGIPHQSPPTK200000000000200000	sp Q96T58 MINT_HUMAN
3729	GVVSEDIPLNLSR200	sp Q12931 TRAP1_HUMAN
3730	ESAAPASPAPSPAPSPPTAPPQK200020002000000000	sp Q3KQU3 MA7D1_HUMAN
3731	TASRPDDIPSPSSPK2020000	sp Q5VZK9 LR16A_HUMAN
3732	IYHLPDAESDEDEDFK2000000000	sp Q15019 SEPT2_HUMAN
		sp P05534 1A24_HUMAN
		sp Q09160 1A80_HUMAN
3733	GGSYSQAASSDSAQGSVDVSLTACK2000000000000000	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
		sp P42166 LAP2A_HUMAN
3734	SSTPLPTISSAENTR2000000000000000	sp P42167 LAP2B_HUMAN
		sp Q5T4S7 UBR4_HUMAN
3735	TSPADHGGVSGSESGSAVDSVAGEHSVSGR2.20000002E+31	sp P25205 MCM3_HUMAN
3736	RSEDESETEDEEEKSQEDQEK202000000000000000	sp O15027 SC16A_HUMAN
3737	RSSLSSHSHQSQIYR20200000000000	sp Q6ZNJ1 NBEL2_HUMAN
3738	RISQVSSGETEYNPTEAR200000000000000000	sp Q12996 CSTF3_HUMAN
3739	RPNEDSDEDEEKGA VPPVHDIYR2E+19	sp Q15477 SKIV2_HUMAN
3740	ASSLEDLVLK200000000	sp Q13435 SF3B2_HUMAN
3741	GFEEHKSDDSSDDEQEK2000220000000	sp Q9Y679 AUP1_HUMAN
3742	LRPQSAQSSFPSPGSPDVQLATLAQR2000000000000000	sp Q7Z417 NUFP2_HUMAN
3743	NDSWGSFDLR200000000	sp Q13596 SNX1_HUMAN
3744	LPPFPGLEPESEGAAGGSEPEAGSDTEGEDIFTGAAVVSK202000000000000000	sp P47712 PA24A_HUMAN
	000000	sp Q5JPI9 MET10_HUMAN
3745	HIVSNDSSDDESHEPK2202000000000	sp Q8IYB3 SRRM1_HUMAN
3746	SDKSGPGEDGFVPSALGTR2000000000000000	sp P19532 TFE3_HUMAN
3747	SPSPAPPPR2020000000	sp Q92597 NDRG1_HUMAN
3748	AASDPLLSSVPAVSK2000000	sp Q3KR37 GRM1B_HUMAN
3749	TASGSSVTSLDGTR2022020000000	sp O43818 U3IP2_HUMAN
3750	VPHLEEVMSPVTTPTDEDVGH1200020000000000	sp Q9UPP1 PHF8_HUMAN
3751	MNEISSDSESESLAPR2202000200000	sp Q9NYF3 FA53C_HUMAN
3752	DAEYIYPSLESDDDDPALK2000000000	sp Q9C0C2 TB182_HUMAN
3753	RFLSPLSLGPQASR2020000000000	sp O43237 DC1L2_HUMAN
3754	VSGAGFSPSSK200000	sp Q7L1V2 MON1B_HUMAN
3755	DFQDYMEPEEGCQSPQR20000	sp P02545 LMNA_HUMAN
3756	DKDQPPSPSPPPQSEALSSTR202000000000000000	sp Q15149 PLEC_HUMAN
3757	LRLSPSPTSQR202000000	
3758	SSSVGSSSYPIPAVSR220022000000000000	

3759	SELISEGRPVETDSESEFPVFPTALTSER2020000000000000000	sp Q5T5U3 RHG21_HUMAN
3760	LPNLSSPSAEGPPGPPSGPAPR2000000000000000000	sp O60784 TOM1_HUMAN
3761	SPQPVCSPAGSEGTK2000002000000000000	sp Q6KC79 NIPBL_HUMAN
3762	AGSPLCNSNLQDSEEDIEPSYSLFR2.00002E+23	sp Q9NRY4 RHG35_HUMAN
3763	RVSLSEIGFGK2000000000	sp Q00536 CDK16_HUMAN
3764	VASETHSEGSEYEELPK202002000000000	sp Q5JSZ5 PRC2B_HUMAN
3765	ISSKSPGHMVILDQTK202000100000000	sp Q9NWH9 SLTM_HUMAN
3766	KPVTVSPTTPTSPTTEGEAS20	sp Q9Y6G9 DC1L1_HUMAN
3767	VSGAQEMVSSAK2000000000000	sp O60664 PLIN3_HUMAN
3768	EESDDEDEKESSEEPK2200000000000000000	sp P35659 DEK_HUMAN
3769	KTSDANETEDHLESLICK2000000000000	sp Q09161 NCBP1_HUMAN
3770	STPLASPSPPGR2000200000000	sp Q8WYP5 ELYS_HUMAN
3771	KIPEPSPVTR200000	sp Q7Z4S6 KI21A_HUMAN
3772	SPSLLQSGAK20000020000	sp Q9BQG0 MBB1A_HUMAN
3773	STAQQELDGKASPTPVIVASHTANKEEK2000000000000000000	sp P35606 COPB2_HUMAN
3774	EDSGTFLGK200000000	sp O75569 PRKRA_HUMAN
3775	KTSSDDESEEDDLLQR2220000000000000000	sp Q9Y5J1 UTP18_HUMAN
3776	QHEVEIPSPTQK200000	sp Q08499 PDE4D_HUMAN
3777	HSSISPSTLTK2202000000000	sp Q14004 CDK13_HUMAN
3778	ASPVTSAAAFTASPANK2000000000000200000	sp Q9UIF9 BAZ2A_HUMAN
3779	VSGYSTCHSR200	sp Q5T8I3 F102B_HUMAN
3780	RPPSQLTSSSLSSPDIFDSPSPEEDKEEHISLAHR2000000000200000	sp Q96B97 SH3K1_HUMAN
3781	AATTATPAATTSPK2000	sp Q92766 RREB1_HUMAN
3782	TPGPPPPDDEDDPVPLPVSGDKEEDAPHR2E+31	sp Q69YN4 VIR_HUMAN
3783	TRSPSPTLGESLAPHK200020000000000	sp Q86UU1 PHLB1_HUMAN
3784	DVDDGSGSPHSPHQLSSK200002000000000	sp Q9H4A3 WNK1_HUMAN
3785	GEFSASPMLK200000	sp Q9UQ35 SRRM2_HUMAN
3786	ASTASPCNNNINAATAVALQEPR2E+21	sp Q71RC2 LARP4_HUMAN
3787	VMHSSPSSSDGK20000000	sp Q969R2 OSBP2_HUMAN
3788	GSDALSETSSVSHIEDLEK202200000000000	sp Q6P996 PDXD1_HUMAN
3789	SPTPTVNPR2000000000	sp Q8WY36 BBX_HUMAN
3790	TASPPPPPK20000000	sp Q8IYB3 SRRM1_HUMAN
3791	GGSYSQAASSDSAQGSVDVSLTACK20002000000000	sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
		sp P01892 1A02_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
3792	EFGYDSPHDLSD200	sp Q96CW6 S7A60_HUMAN
3793	DPVYDESTDEESENQTDLSGLASAAVSK2.0002E+21	sp Q13426 XRCC4_HUMAN
3794	ISFVEEDVHPK20000000000	sp Q8WYP5 ELYS_HUMAN
3795	YSPTSPTYSPK300000002000000	sp P24928 RPB1_HUMAN
3796	AAFDDAIAELDTLSEESYK2000	sp P62258 1433E_HUMAN
3797	DDEKEAEEGEDDRDSANGEDDS20	sp P07910 HNRPC_HUMAN
3798	SNMSPHGLPAR200000000	sp Q13330 MTA1_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P05534 1A24_HUMAN
3799	GGSYSQAASSDSAQGSVDVSLTACK20002000000000	sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
		sp P30447 1A23_HUMAN
3800	SSPSARPPDVPQQPQAAK2.2E+19	sp Q96JP5 ZFP91_HUMAN
3801	VSPASPAGSPSADFAVHGESLGDR2.0020002E+23	sp Q96C92 SDCG3_HUMAN



3802	NTSSDNSDVEVMPAQSPR2002000000000000	sp Q9HCK8 CHD8_HUMAN
3803	TETPPPLASLNVSK2000000000000000	sp Q3MHD2 LSM12_HUMAN
3804	TVDSQGPTPVCTPTFLER200020000000	sp Q9BYG3 MK67I_HUMAN
3805	KGNSPNEPPTPK20000000000	sp P48634 PRC2A_HUMAN
3806	DHNSDEDEDKYADDIDMPGQNFDSK2E+23	sp 095391 SLU7_HUMAN
3807	VKPETPPR20000	sp Q9UQ35 SRRM2_HUMAN
3808	QDDSPSGASYGQDYDLSPSR2000000000020000	sp Q9NYV4 CDK12_HUMAN
3809	QQPQPSQTESPQEAQIIQAK2000000000000000	sp 096020 CCNE2_HUMAN
3810	QLPALDGLMGPESPPAQEEEEAPVSPHK2. 01E+21	sp Q8N9T8 KRI1_HUMAN
3811	LFDSSDDDESSEDSDSNRFK220000202000000000	sp Q76FK4 NOL8_HUMAN
3812	AAEDDSASPPGAASDAEPGDEERPGLQVDCVCGDK2. 00000002E+31	sp P10588 NR2F6_HUMAN
3813	MDSNFGLAGLPGSPK2000	sp 043463 SUV91_HUMAN
3814	SATDGNTSTPPTSAK20000000	sp Q13620 CUL4B_HUMAN
3815	SASSSEEEELYLGPSPTR2. 0222E+19	sp Q6QNY0 BL1S3_HUMAN
3816	RPESPSEISPIK2000000000	sp Q7Z5K2 WAPL_HUMAN
3817	WLDDLLASPPPSGGGAR20000000000	sp Q9C0C2 TB182_HUMAN
3818	NNSFTAPSTVGK20000000000	sp 095453 PARN_HUMAN
3819	TVTCTVVEPEAPSPDVLQAATHR200000000000	sp Q9P1Y6 PHRF1_HUMAN
3820	SFSKEVEER20000000	sp Q04637 IF4G1_HUMAN
3821	SPLQSVVVR2000000000	sp Q9Y2W1 TR150_HUMAN
3822	SPPPGPDGHAK200000000000	sp P57081 WDR4_HUMAN
3823	TSAVSSPLLDQQR20002000000000	sp Q5T200 ZC3HD_HUMAN
3824	EKGPTTGEGALDLSDVHSPPKSPEGK2000200000	sp 095684 FR10P_HUMAN
3825	ASPITNDGEDEFVPSDGLDKDEYTFSPGK2. 0000000000002E+28	sp Q02880 TOP2B_HUMAN
3826	ESLGSEESGKDWDELEEEAR2. 0020002E+20	sp Q9Y5B9 SP16H_HUMAN
3827	FGSEEVEMEVESEDEDDKQEK1000200000000000	sp Q15459 SF3A1_HUMAN
3828	MPLDLSPLATPIIR2000000000	sp P15336 ATF2_HUMAN
3829	ENSPAVSPTTNSTAPFGLKPR2002000000000000	sp Q9Y2L9 LRCH1_HUMAN
3830	GPPSPAPVMHSPSR2000000020000	sp Q13573 SNW1_HUMAN
3831	MESEGGADDSAEGLDLDLDDDDNEDRGDDQLELIKDDEK2E+36	sp P07910 HNRPC_HUMAN
3832	HASSSPSPKAPAPGSHR2002000000000000	sp Q9UQ35 SRRM2_HUMAN
3833	NGSLDSPGKQDTEEDEEEDEKDK2. 000000002E+21	sp Q9Y266 NUDC_HUMAN
3834	KDEETEESYDSEHENSEPVTNIR2. 002000200002E+20	sp Q15154 PCM1_HUMAN
3835	GYGFVDFDSPAQAQK20000000	sp Q15434 RBMS2_HUMAN
3836	GPTSTSIDNIDGTPVRDER20000000	sp Q5VT52 RPRD2_HUMAN
3837	TVSTSSQPEENVDR2002000000000	sp P42684 ABL2_HUMAN
3838	NLDPDPEPPSPDSPTETFAAPAEVR2002000000000000	sp Q9BUA3 CK084_HUMAN
3839	SQLPTLEQDGGTQNPVSSPGMSQELR2000000000	sp P46937 YAP1_HUMAN
3840	SHHAPSPGSSGGGQPLAR1200000000000000	sp 014497 ARI1A_HUMAN
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3843	NSPEDLGLSLTGDSCK2000000200000000	sp Q12888 TP53B_HUMAN
3844	VVSDADSDSDAVSDK200002020002000	sp Q96ST2 IWS1_HUMAN
3845	STFSTNYR20000000	sp P05783 K1C18_HUMAN
3846	SGSSQELDVKPSASPQER22000000000200000	sp Q9UQ35 SRRM2_HUMAN
3847	VGSSNFR200000	sp P05787 K2C8_HUMAN
3848	QLSGQSTSSDTTYK2000000000000	sp Q9UGU0 TCF20_HUMAN
3849	KASPEAASTPR2000000000	sp P29590 PML_HUMAN
3850	VVSSTSEEEAFTEK22020000000000	sp Q8N573 OXR1_HUMAN
3851	LLPYPTLASPASD200200	sp P0C1Z6 TFPT_HUMAN
3852	KADSDSEDKGEESKPK20000000000000	sp P83916 CBX1_HUMAN
3853	LHGGFSDCSEDGEALNGEPELDLTSK2. 002E+21	sp 000178 GTPB1_HUMAN
3854	REESPMVDVQPSQAQDTQSIASDGTQGEK20000000000000	sp Q7Z6Z7 HUWE1_HUMAN
3855	QGSITSPQANEQSVTPQRR200200000000200000	sp Q9UQ35 SRRM2_HUMAN

3856	KFDHESSPGTDEDK2200200000	sp Q13427 PPIG_HUMAN
3857	YFDTNSEVEEESEDEDYIPSEDWK2.000002E+20	sp Q8N108 MIER1_HUMAN
3858	SLSQIHEAAVR2000000000	sp Q9NZM1 MYOF_HUMAN
3859	ASAVSELSPR2000002000	sp Q9Y2W1 TR150_HUMAN
3860	FSISPEDESSSYSSNSDFNYSYPTK22000000000000	sp Q96QD8 S38A2_HUMAN
3861	KASPEPPDSAEGALK20000000000000	sp Q9H1B7 I2BPL_HUMAN
3862	NDSPTQIPVSSDVCR20000000000000	sp Q8TC07 TBC15_HUMAN
3863	TSSSSPANSDVEIDGIGR2000020000000000	sp 060293 ZC3H1_HUMAN
3864	QIDSSPVGGETDETTVSQNYR200000000000	sp 015027 SC16A_HUMAN
3865	GTDTQTPAVLSPSK20000	sp P46087 NOP2_HUMAN
3866	RPPSPEPSTK20000000	sp Q01082 SPTB2_HUMAN
3867	KASQAASLASK2000000000	sp Q15154 PCM1_HUMAN
3868	GLFQDESDSCDCSYR202000000	sp Q8NEM2 SHCBP_HUMAN
3869	AGSISTLDSLDFAR2000002000000	sp Q9P260 K1468_HUMAN
3870	AQSYDIQAWK200000000	sp Q8WWQ0 PHIP_HUMAN
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3872	LPSDITDSDASEDPGGAER2020020020000000000	sp Q9UJK0 TSR3_HUMAN
3873	QSLPASPR2000	sp Q3KQU3 MA7D1_HUMAN
3874	LLGPSAAADILQLSSSLPLQSR20000000	sp Q7Z6Z7 HUWE1_HUMAN
3875	VVLGDGVQLPPGDYSTTPGGTLFSTTPGGTR3000000000002000000	sp Q13541 4EBP1_HUMAN
3876	TPASTPVSGTPQASPMIER2010000	sp Q9UHR4 BI2L1_HUMAN
3877	NSYNSQAPSPGLGSK20000000	sp Q9UHB7 AFF4_HUMAN
3878	SPQTLAPVGEDAMKTPSPAAEDAR2.0000000010202E+21	sp Q92797 SYMPK_HUMAN
3879	IPSIVSSPLNSPLDR2000200000	sp P49790 NU153_HUMAN
3880	MSSPEDSDTKR220000000000	sp Q15424 SAFB1_HUMAN
3881	I IPTLEGLQLPSPTATSQLPLESDAVECLNYQHYK2E+20	sp P61978 HNRPK_HUMAN
3882	RDSSDWEIPDGQITVGQR200000000000000000	sp P15056 BRAF_HUMAN
3883	FFTTGSDSESESSLGEEELVTKPVGGNYGK2.0202202E+23	sp B5ME19 EIFCL_HUMAN
3884	SASPHDVLCLVSPCFEHR2000000000000000000	sp Q99613 EIF3C_HUMAN
3885	LGAGGSPEKSPSAQELK2000200000000	sp Q66K74 MAP1S_HUMAN
3886	SVSGSPEPAAK200020000000	sp Q9UNE7 CHIP_HUMAN
3887	AQPFGFIDSDDAEER2020000000	sp Q8IYB3 SRRM1_HUMAN
3888	FNYSGSGGR20000	sp Q14676 MDC1_HUMAN
3889	EVDDILGEGSDSDSEK200002000	sp Q12906 ILF3_HUMAN
3890	SSSPFLSK22000000	sp Q9Y5B0 CTDP1_HUMAN
3891	DRPVGGSPGGPR2000000	sp Q9NYV4 CDK12_HUMAN
3892	TPLSSLSATSNNSTVACIDR2000000000000000000	sp Q8NHG8 ZNRF2_HUMAN
3893	KPSGSPDLWKLSPDQR200000000200000	sp P00533 EGFR_HUMAN
3894	AAYGDLSEEEEEENEPESLGVVYK3.00002E+22	sp Q96JM3 CHAP1_HUMAN
3895	AASSSPGSPVASSPSR20000000000000	sp 015541 R113A_HUMAN
3896	LPSPTSPFSSLSQQAATSK2000000000000000	sp Q92538 GBF1_HUMAN
3897	SGISTNHADYSSSPAGSPGAQVSLYNPSVASPAR2.2E+24	sp Q8WWI1 LMO7_HUMAN
3898	QVSASELHTSGILGPETLR2000000000000000	sp Q8WV41 SNX33_HUMAN
3899	TPTPSTPGDTQPNTAPVPPAEDGIK2E+21	sp P58107 EPIPL_HUMAN
3900	TESPPSPSSDVEDAR20000220000000	sp Q14839 CHD4_HUMAN
3901	QPSWDPSVSSVTPAPSPLSAAAVSPSK2E+26	sp Q9BRK4 LZTS2_HUMAN
3902	SGSSQELDVKPSASPQER2020000000000000000	sp Q9NQC3 RTN4_HUMAN
3903	SPFNSPSPQDSPR20000020000000	sp Q9UQ35 SRRM2_HUMAN
3904	EGDPVSLSTPLETEFGSPSELSPR202000000	sp P08651 NFIC_HUMAN
3905	SLSGSSPCPK20200200000	sp Q9BUR4 WAP53_HUMAN
3906	KRSSSEDAESLAPR2020000000000	sp Q9UQ35 SRRM2_HUMAN
3907	KRSDAGSSTPTTSTR20000000000000	sp Q4G0J3 LARP7_HUMAN
3908	SVASNQSEMEFSSLQDMPK20000100000001000	sp P51532 SMCA4_HUMAN
		sp Q8N163 CCAR2_HUMAN

3909	KVVVSPTK20000	sp P19338 NUCL_HUMAN
3910	SPSSEWLVR2000000000	sp Q9NWQ4 GPT2L_HUMAN
3911	SFQQSSLSRDSQGHGR2002000000	sp P43243 MATR3_HUMAN
3912	AGEPNSPDAEEANSPDVTAGCDPAGVHPPR200020000000000000	sp Q08J23 NSUN2_HUMAN
3913	TSSKESSPIPSPTSDR220002000000	sp Q01082 SPTB2_HUMAN
3914	AAKNSEEEEEEEK200000000	sp Q14978 NOLC1_HUMAN
3915	VSSPLSPSAMTDAANSQAAAK2E+21	sp Q8WXI9 P66B_HUMAN
3916	SNSTTQVSQPR200000000000	sp Q9UPN4 CP131_HUMAN
3917	VELSESEEDKGGK20200000000	sp Q8IYB3 SRRM1_HUMAN
3918	TEARSSDEENGPPSSPDLDR22000000020000000	sp Q96B36 AKTS1_HUMAN
3919	SQSLPTLLSPVR20000000000000	sp P28290 SSFA2_HUMAN
3920	SFEEEGEHLGSR20000000000000	sp Q9BXW6 OSBL1_HUMAN
3921	SPSVPKSPTPK2000200000	sp Q9NVD7 PARVA_HUMAN
3922	AQSLVISPPAPSPR200002000	sp P46013 KI67_HUMAN
3923	KPSKYPLCSADAAVSTTPPGSPPPPPLPEPPVLK2.0002E+19	sp Q9BTC0 DIDO1_HUMAN
3924	QGSGRESPLASR20000000	sp Q8WWM7 ATX2L_HUMAN
3925	NSLESISSIDR20020020000	sp Q14160 SCRIB_HUMAN
		sp Q29960 1C16_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
3926	GGSCSQAASSNSAQGSDESLIACK2002000000	sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
		sp P30510 1C14_HUMAN
3927	STPESGSDKESVGSSTSNEGGR200020000000000000	sp Q6GQQ9 OTU7B_HUMAN
3928	SISGTPAETPACK200000000000	sp Q9Y4D8 HECD4_HUMAN
3929	KKPSTDEQTSSAEEDVPTCGYLNVLNSNR2.20002E+26	sp Q8N556 AFAP1_HUMAN
3930	AELGMDSTSQSPPIK200000	sp Q9NR19 ACSA_HUMAN
3931	KEPAITSQNSPEAR200000000	sp P23193 TCEA1_HUMAN
3932	KFDHESSPGTDEDK200200000	sp Q13427 PPIG_HUMAN
3933	NGFPHPEDCNPSEAASEESNSEIEQEIPVEQK2.000200202E+21	sp Q9NR30 DDX21_HUMAN
3934	GSSGSPAHAESYSSGGGQK2E+20	sp P15408 FOSL2_HUMAN
3935	EQGTESRSSTPLPTISSAENTR2.0202E+20	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
3936	SPGSSWQEYGPSDTCGSR200000000000000000	sp Q5JSZ5 PRC2B_HUMAN
3937	TDSVIIADQTPPTR2020000	sp P17544 ATF7_HUMAN
3938	SPSPYYSR202000000	sp Q13595 TRA2A_HUMAN
		sp P62995 TRA2B_HUMAN
3939	SKEFVSSDESSGENK220022000000	sp Q08945 SSRP1_HUMAN
3940	TASRPDDIPDSSPK200000002000000	sp Q5VZK9 LR16A_HUMAN
3941	SFSKEELMSSDLEETAGSTIPK200000000000000000	sp O00567 NOP56_HUMAN
3942	YSPTSPTYSPTPK300000000020000	sp P24928 RPB1_HUMAN
3943	SSSVGSSSYPISPAISR22000000000000000000	sp Q15149 PLEC_HUMAN
3944	FGQAATMEGIGAIGGTPPAFNR10000000020000000	sp Q15233 NONO_HUMAN
3945	TASQEGCQELTQEQR200000000000000000	sp O43933 PEX1_HUMAN
3946	SGSPAPETTNEVPFAQHSSLDSR2E+22	sp O15047 SET1A_HUMAN
3947	QNSDPTSEPGPSPNPPAWVRPDNEAPPK2E+27	sp Q8N4C8 MINK1_HUMAN
3948	TQTPPLGQTPQLGLK20000000000000	sp P78344 IF4G2_HUMAN
3949	QSSGSATNVASTPDNR2000000000000000	sp Q7Z460 CLAP1_HUMAN
3950	RPASPSPEHLPATPAESPAQR220000000000000000	sp Q9H7L9 SDS3_HUMAN

3951	DGDSYDPYDFSDTEEEMPQVHTPK300000000100002000	sp P25205 MCM3_HUMAN
3952	DGDHVCSTGPASSSGEDDDED22000000000	sp Q8NEL9 DDHD1_HUMAN
3953	IALESEGRPEEQMESDNCSGGDDWTHLSSK200000000000000000	sp Q13501 SQSTM_HUMAN
3954	SNQIPTEVR2000000000	sp Q9BWH2 FUND2_HUMAN
3955	QGSITSPQANEQSVTPQR20000002000000	sp Q9UQ35 SRRM2_HUMAN
3956	SPPGAGSPPGAAVAPSALQLSPAAAAAAAHR2E+32	sp P78415 IRX3_HUMAN
3957	KPPAPSPVQSQSPSTNWSPAVPVKK2.000000002E+20	sp Q8IYB3 SRRM1_HUMAN
3958	ICPVPVSTVTGDENISDSEIEQYLR20000000000	sp Q9HAW0 BRF2_HUMAN
3959	FFTTGSDSESESSLGEEELVTKPVGGNYGK2.020202E+27	sp B5ME19 EIFCL_HUMAN
		sp Q99613 EIF3C_HUMAN
3960	FSTYSQSPDTPSLR2000000000	sp Q6ZS17 FA65A_HUMAN
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		sp P20700 LMNB1_HUMAN
3963	NVSSFPDDATSPLQENR20000000	sp P52292 IMA1_HUMAN
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		sp Q09160 1A80_HUMAN
3964	GGSYSQAASSDSAQGSVSLTACK2000000000	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
3965	RPDYAPMESSDEEDEFQFIKK2200000000000000	sp P55081 MFAP1_HUMAN
3966	SGSSPDSEITELK200000000000	sp P17812 PYRG1_HUMAN
3967	SGSLDSELSVSPK2000	sp Q12802 AKP13_HUMAN
3968	TASESISNLSEAGSIK20020002000	sp P30622 CLIP1_HUMAN
3969	SGGLQTPECLSR200	sp P85037 FO XK1_HUMAN
3970	SSTDSLPGPISR200000000000	sp Q9HCD5 NCOA5_HUMAN
3971	SSTPLPTISSAENTR2000200000000000	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
3972	QSSVTQVTEQSPK2000	sp Q14966 ZN638_HUMAN
3973	GLVAAYSGESDSEEEQER2000020000000	sp P98175 RBM10_HUMAN
3974	KPSDSLVAASSR2000000200	sp Q5T0W9 FA83B_HUMAN
3975	KPSPQPSSPR22000	sp 075909 CCNK_HUMAN
3976	QGPPCSDSEEVER2020000000	sp 043159 RRP8_HUMAN
3977	EIAIVHSDAEK200000	sp Q9H307 PININ_HUMAN
3978	HASSPESPKPAPAPGSHR220002000000000000	sp Q9UQ35 SRRM2_HUMAN
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3980	TSSKESPIPSPTSDR200220002000000	sp Q01082 SPTB2_HUMAN
3981	FYSDSHHLK20000000	sp Q92576 PHF3_HUMAN
3982	EIPSATQSPISK200000	sp Q9BQG0 MBB1A_HUMAN
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3984	RPQSPGASPSQAER200000000000	sp Q96T58 MINT_HUMAN
3985	ASYHFSPEELDENTSPLLGDAR200000000	sp 075410 TACC1_HUMAN
3986	MVEPENAVTITPLRPEDDYSPR2000	sp Q14160 SCRIB_HUMAN
3987	KDSSGISEQLPCR2002000000000	sp Q8NEZ4 KMT2C_HUMAN
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3990	SGSSQELDVKPSASPQR2000000020000000	sp Q9UQ35 SRRM2_HUMAN
3991	GIQYIDLSSDSEDVVPNCNTVQEK2.202E+19	sp Q9H4L7 SMRCD_HUMAN
3992	KEESEESDDMGFLFD200010000000	sp P05387 RLA2_HUMAN
		sp P05386 RLA1_HUMAN
3993	ALASNTSFFSGCSPIEEEAH200000000	sp 095239 KIF4A_HUMAN
		sp Q2VIQ3 KIF4B_HUMAN

3994	QKSDAEEDGGTVSQEEEDR2000000000	sp P27824 CALX_HUMAN
3995	LTPVSPESSTEELK20002000000	sp Q13501 SQSTM_HUMAN
3996	NDSPTQIPVSSDVCR200000000000	sp Q8TC07 TBC15_HUMAN
3997	RASQEANLLTLAQK2000000000000	sp Q6PJG2 EMSA1_HUMAN
3998	VGTEGLVDSFSQGLER2000000000	sp 075170 PP6R2_HUMAN
3999	SGIDITDFQAK2000000000000	sp P49790 NU153_HUMAN
4000	ASASGSGAQVGGPISSGSSASSVTVTR2020000000	sp P02545 LMNA_HUMAN
4001	AGMSSNQSISSPVLDVPR2022000000000	sp Q9UQ35 SRRM2_HUMAN
4002	EISPGSGPGEIR200000000000	sp 043491 E41L2_HUMAN
4003	SPQPPSR200000000	sp Q86VM9 ZCH18_HUMAN
4004	TPLTSADEHVHVK20022000000000	sp Q5SW79 CE170_HUMAN
4005	VTQHESDNENEIQIQNK2000000000000	sp Q5VZL5 ZMYM4_HUMAN
		sp P30505 1C08_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp Q29960 1C16_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
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		sp Q29963 1C06_HUMAN
		sp P30510 1C14_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
4007	GPAAPLTPGPQSPPTPLAPGQEK2000020000000000000	sp 075398 DEAF1_HUMAN
4008	SASPHVDLCLVSPCFEHR2000000000200000000	sp Q66K74 MAP1S_HUMAN
4009	GLGPPSPAPP20000000	sp Q13425 SNTB2_HUMAN
4010	VVEAVNSDSEFGIPKK2020000000000	sp Q02880 TOP2B_HUMAN
4011	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2.00000002E+29	sp Q8WUZ0 BCL7C_HUMAN
4012	ERSDSGSSSEPFDR200200000000	sp P48634 PRC2A_HUMAN
4013	LPSPTSPFSSLSQDQAATSK20020000000000000	sp Q8WWI1 LM07_HUMAN
4014	TVSPGVSPIHGQGVVENLK2.00002E+19	sp Q9NZM3 ITSN2_HUMAN
4015	LQAALDDEEAGGRPAMEPGNGSLDLGGDSAGR2000000000000	sp Q00839 HNRPU_HUMAN
4016	LPDGPTGSSEEEEFLEIPPFNK2200000000000000000	sp Q12888 TP53B_HUMAN
4017	EAPGSPPLSPR20000000	sp Q86UU0 BCL9L_HUMAN
4018	NSLEISSIDR200000000000	sp Q14160 SCRIB_HUMAN
4019	ASGEMASQYITAALR20010000000000000	sp P54578 UBP14_HUMAN
4020	LGMLSPEGTCK20000000	sp P49327 FAS_HUMAN
4021	GRLTPSPDIIVLSDNEASSPR200000000000002000	sp Q8WXI9 P66B_HUMAN
4022	IILDLISESPIK20000	sp P61978 HNRPK_HUMAN
4023	EWYQSTIPQSPAPDDPEEGR202000000000000	sp Q08499 PDE4D_HUMAN
4024	ENSPAVSPTTNSTAPFGLKPR2020000000000000000	sp Q9Y2L9 LRCH1_HUMAN
4025	RFSSSDFSDLSNGEELQETCSSLR2.22002002E+23	sp P23497 SP100_HUMAN
4026	HQQQLLASPGSSTVDNK20000000000	sp Q9C0E2 XPO4_HUMAN
4027	LPSTSGSEGVPR202020000000	sp P29317 EPA2_HUMAN
4028	TSSVSSISQVSPER200000000000000	sp P42356 PI4KA_HUMAN
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4030	GGSSVEAPCPDVTPEDDR2000000	sp Q9BY89 K1671_HUMAN
4031	RPSTSQTVSTPAPVPVIESTEAEAK2E+23	sp P12270 TPR_HUMAN
4032	SLSPSHLTEDR2000000000000	sp Q5T200 ZC3HD_HUMAN
4033	DEVFHTDQDDPSSSDDEGMPYTRPVK2220000100000000	sp Q5JSH3 WDR44_HUMAN
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4035	SPGVEKPIVKPTAGAGPQETNMK2E+23	sp Q76L83 ASXL2_HUMAN

		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
4036	LNQSGTSVGTDEESDVTQEEER2.022E+19	sp A6NKT7 RGPD3_HUMAN
		sp PODJD0 RGPD1_HUMAN
		sp PODJD1 RGPD2_HUMAN
4037	SNSRLEDQVIGVALQASK200000000000000000	sp Q2KHR3 QSER1_HUMAN
4038	TTWGDGGENSPCNVSK200000000	sp 000161 SNP23_HUMAN
4039	DGTAPPQSPGSPGTQDEEWSDEESPR200000000020000000	sp Q9UKS6 PACN3_HUMAN
4040	SVSGSPEPAAK202020000000	sp Q8IYB3 SRRM1_HUMAN
4041	SVNEILGLAESSPNEPK2000000	sp Q9H8G2 CAAP1_HUMAN
4042	LGSTSGEESDLER220000200000	sp Q9BW71 HIRP3_HUMAN
4043	DNLGEVPLTPTEEASLPLAVTK2000000000000000	sp Q6ZJ1 NBEL2_HUMAN
4044	TPSSPLQSCTPSLSR2000000000000000	sp Q96AP0 ACD_HUMAN
4045	EDTDHEEKASNEVDTK20000000	sp 075475 PSIP1_HUMAN
4046	QTDPSTPQQESSKPLGGIQPSSQTIQPK2E+24	sp Q14004 CDK13_HUMAN
4047	TDHGAEIVYKSPVVGDTSPR20022000	sp P10636 TAU_HUMAN
		sp TAU_HUMAN
4048	YSHSYLSDSDTEAK2000000	sp Q92614 MY18A_HUMAN
4049	SISLGDSEGPVATLAQPLR2000000000000000	sp 043379 WDR62_HUMAN
4050	EAEQAASEAAGGDTTPGSSPSSLYEEPLGQPPR2E+28	sp Q86TC9 MYPN_HUMAN
4051	MEREDSSEEEEEIDDEEIER220000000000000000	sp P55081 MFAP1_HUMAN
4052	LASTSDIEEK202000000	sp 014974 MYPT1_HUMAN
4053	ENSTVDFSK20000000	sp Q6U841 S4A10_HUMAN
		sp Q9Y6M7 S4A7_HUMAN
4054	REPESPPASAPIPLFGADTIGQR2E+19	sp Q96AY4 TTC28_HUMAN
4055	ELQGDGPPSSPTNDPTVK200000000000	sp Q12873 CHD3_HUMAN
4056	KKSSSEAKPTSLGLAGGHK202000000000000000	sp Q96RK0 CIC_HUMAN
4057	LPPAAPIADSENNSTMASASEGEMECGQELK2020000000000000	sp Q63HN8 RN213_HUMAN
4058	SPFEGAVTESQSLFSDNFR2E+19	sp 095628 CNOT4_HUMAN
4059	SPQPAEEDDDFDLVAIDTYNCDLHFK2E+30	sp Q9BUJ2 HNRL1_HUMAN
4060	LDSSPSVSSTLAAK2000000000000000	sp Q13428 TCOF_HUMAN
4061	LGSTSGEESDLER202000200000	sp Q9BW71 HIRP3_HUMAN
4062	CPEILSDESSDEDEKK20022000000000	sp P35659 DEK_HUMAN
4063	STSPAGQHHSPISR2200000020000000	sp Q5T200 ZC3HD_HUMAN
4064	GKASPFEEDQNR2000000000	sp Q8TB72 PUM2_HUMAN
4065	RPESPSEISPIK2000020000	sp Q7Z5K2 WAPL_HUMAN
4066	SKFDSDEEEEDTENVEAASSGKVTR2.0000000000002E+21	sp Q8TF01 PNISR_HUMAN
4067	AAALQALQAQAPTSPPPPPPLK200000000000	sp Q8NAF0 ZN579_HUMAN
4068	TTSISPALAR202000000	sp Q9UKE5 TNIK_HUMAN
4069	EVDPTGELQSLQMPESGSSLDPSQEGPTGLK20000000000000	sp Q13501 SQSTM_HUMAN
4070	TPPVVIK20000000	sp Q96A49 SYAP1_HUMAN
4071	DDISEIQSLASDHSGR2000	sp Q07157 ZO1_HUMAN
		sp P30505 1C08_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp Q29960 1C16_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
4072	GGSCSQAASSNSAQGSDESLIACK20000000000000	sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp P30510 1C14_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
4073	AESSSSISSASLSAANTPTVGVSR220202000000000000	sp Q0JRZ9 FCHO2_HUMAN

4074	APAGQEEPGTPSSPLSAEQ LDR200200000000000	sp	P13051	UNG_HUMAN
4075	ASETVSEASPGSTASQTGVPTQVVQQVQGTQQR2E+28	sp	P22670	RFX1_HUMAN
4076	SEESPVYNSSSVR20000000000	sp	Q03188	CENPC_HUMAN
4077	EVEEDESEDEEMSEDEEDSSGEEVVIPQK2. 00001200000022E+24	sp	P19338	NUCL_HUMAN
4078	ETSSPGTDDVFTPAPSDSPSSQR2000000	sp	P19634	SL9A1_HUMAN
4079	SGGLQTPECLSR20000200	sp	P85037	FOXK1_HUMAN
4080	GSPRPPPPPPPLLGTVSSPSSSPTHLWTGEVSAAPPPAR2. 0000000000002E+38	sp	Q6P1L5	F117B_HUMAN
4081	SPSGPSGSLENGTK2000000000000	sp	Q14134	TRI29_HUMAN
4082	KTSLSDLNDPIILSSDDDDNDNR2. 0000000022E+19	sp	Q9GZR1	SENP6_HUMAN
4083	SSDIVSSVR200000000	sp	Q14C86	GAPD1_HUMAN
4084	AAVVTSPPTTAPHK20000000000	sp	P35611	ADDA_HUMAN
4085	VAHSDKPGSTSTASFR200000000	sp	Q9P0L0	VAPA_HUMAN
4086	YHGHSMSDPGVSYR21000002000	sp	P08559	ODPA_HUMAN
4087	VDSDSDSSEDDINSVMK20202000000000000	sp	P29803	ODPAT_HUMAN
4088	SPSAMQQQDGLDR200000000000000	sp	Q6KC79	NIPBL_HUMAN
4089	KPLTTSGFHHSEEGTSSSGSK2000000000000	sp	060343	TBCD4_HUMAN
4090	NSQEDSEDEDKDVK20000000	sp	Q5SW79	CE170_HUMAN
4091	SSSVGSSSYPI SPAVSR200000000000000	sp	Q9H1E3	NUCKS_HUMAN
4092	VNPHKVPASSVDSNIPSSQGYK220000000000000	sp	Q15149	PLEC_HUMAN
4093	YESQEPLAGQESPLPLATR200000000	sp	Q9UHB7	AFF4_HUMAN
4094	SPELEGEKLSPLPGGPGAGDPR2. 000000002E+22	sp	Q9COC2	TB182_HUMAN
4095	DSSKGEDSAEETEAKPAVVAPAPVVEAVSTPSAAFPSDATAEQGPILTK2E+20	sp	Q15742	NAB2_HUMAN
4096	RNSSPVSPASVPGQR220000000000000	sp	Q12906	ILF3_HUMAN
4097	TDHGAEIVYKSPVSGDTSR200000002000	sp	Q9H1B7	I2BPL_HUMAN
4098	NQSLGQSAPSLTAGLK20000000000	sp	TAU_HUMAN	
		sp	P10636	TAU_HUMAN
		sp	Q6P0Q8	MAST2_HUMAN
		sp	P05534	1A24_HUMAN
		sp	P30447	1A23_HUMAN
4099	KGGSYSQAASSDSAQGS DVSLTACK2E+22	sp	P01891	1A68_HUMAN
		sp	P01892	1A02_HUMAN
		sp	Q09160	1A80_HUMAN
		sp	P10316	1A69_HUMAN
4100	EEDEEPESPPEK200000	sp	Q9H1E3	NUCKS_HUMAN
4101	SQSSEGVSSLSSSPNSLETQSQSLSR2. 0002E+24	sp	095155	UBE4B_HUMAN
4102	RKPEDVLDDDDAGSAPLK200000	sp	P35613	BASI_HUMAN
4103	SPSVSPEPAEK2020220000000	sp	Q9UQ35	SRRM2_HUMAN
4104	DPLTHTSNLPR20000	sp	Q9UBP0	SPAST_HUMAN
4105	TPILEDFELEGVCQLPDQSPPR20000	sp	Q7Z2Z1	TICRR_HUMAN
4106	SPAVKPAAPK2000000000000	sp	Q14978	NOLC1_HUMAN
4107	EEPSQNDISPK2000	sp	Q9NR30	DDX21_HUMAN
4108	DGLNQTTIPVSPSTTKPSR20000000000	sp	Q71RC2	LARP4_HUMAN
4109	AQQGADESSENEEEEEIELIPEK2200000000000000	sp	Q9Y4A0	JERKL_HUMAN
4110	SVSTPSEAGSQDSGDGAVGSR2000000000000000000	sp	Q13409	DC1I2_HUMAN
4111	SGLELTELQNMTPEDDNISNDSNDFTEVENGQINSK1. 200000002E+27	sp	Q9H2H9	S38A1_HUMAN
4112	VLSTSTDLEAAVADALK200000000000000	sp	Q7Z460	CLAP1_HUMAN
4113	SLSNSNPDISGTPTSPDDEV2. 02E+21	sp	Q96N67	DOCK7_HUMAN
4114	SPSKPLPEVTDEYK2000000000000	sp	P05455	LA_HUMAN
4115	TMFAQVESDDEEAKNEPEWK200000000000000	sp	Q9NW82	WDR70_HUMAN
4116	KELSESEDEELQLEEFPLMK2002000000000000000	sp	Q99543	DNJC2_HUMAN
4117	TNSSDSERSPDLGHSTQIPR2020000000000000000	sp	Q6VY07	PACS1_HUMAN

4118	LQTPNTFPK20020000	sp Q14978 NOLC1_HUMAN
4119	KVELSESEEDK20200000	sp Q8IYB3 SRRM1_HUMAN
4120	AAPEASSPPASPLQHLLPGK2000020000000000	sp Q96TA1 NIBL1_HUMAN
4121	ISASSATR20000000	sp P49023 PAXI_HUMAN
4122	SPSAIPEQNHSLNDQAK200000000000000000	sp Q13129 RLF_HUMAN
4123	AGSISTLDSLDFAR20020020000000	sp Q9P260 K1468_HUMAN
4124	TQLWASEPGTPPLPTSLPSQNPILK2E+20	sp Q9BXP5 SRRT_HUMAN
4125	SSEGGVGVGPGGGDEPPTSPPR2E+20	sp Q9Y4H2 IRS2_HUMAN
4126	FNGGHSPHSPK200020000	sp Q9Y5K6 CD2AP_HUMAN
4127	IKEEPEVEVDSSPPDSPASSPCSPPLK220000002200000000	sp P17544 ATF7_HUMAN
4128	EYIPGQPPLSQSSDSSPTR20000	sp P07814 SYEP_HUMAN
4129	SGSPAPETTNESVPFAQHSSLSR2. 00002E+22	sp O15047 SET1A_HUMAN
4130	WTISSSESEEEKPKPKPSTSSLLCAR2. 22002E+25	sp Q9NUW8 TYDP1_HUMAN
4131	MSDSESEELPKPVSDSESEEPPI. 20200000000002E+24	sp Q96ST2 IWS1_HUMAN
4132	YETPSDAIEVISPASSAPPQEK20000000000000	sp O75376 NCOR1_HUMAN
4133	QSHSGSISPYPK20200000	sp Q9UQ35 SRRM2_HUMAN
4134	KPHDCESSTVSEEDYFSSHR2220200000000000	sp Q9H4H8 FA83D_HUMAN
4135	FSEMMNMGDEDVDLPEVDGADDDSDSDEK1. 0001E+30	sp Q15185 TEBP_HUMAN
4136	SAGGSSPEGGEDSDREDGNYCPPVKR2. 20000002E+22	sp Q9H6Z4 RANB3_HUMAN
4137	QFLISPPASPPVGWK200020000000	sp P53805 RCAN1_HUMAN
4138	SEAEEGEVTRPTK20000000020000	sp Q9UKA8 RCAN3_HUMAN
4139	AGMSSNQSISSPVLDVPR200000020000000000	sp Q9H4G0 E41L1_HUMAN
4140	EKGSTLDSLDAEK200000000000	sp Q9UQ35 SRRM2_HUMAN
4141	AVATAAAQAQTGPEEDSGSSEEEESDSEEEAETLAQVKPSGK2. 000002022000 2E+31	sp O00767 ACOD_HUMAN
4142	TSPLNSSGSSQGR20000000000000	sp Q13428 TCOF_HUMAN
4143	AATAARPPAPPAPQPPSPTSPRPTLAR20002000000000	sp P51003 PAPOA_HUMAN
4144	GGPGAAPGGASPASSSAASSPSSGR2000000	sp Q96B36 AKTS1_HUMAN
4145	VHPYEDSDSEDEVDWQDTR300002200000000000	sp Q9BXX1 KLF16_HUMAN
4146	KAPAGQEEPGTPSSPLSAEQ LDR2000000000000000	sp O14730 RIOK3_HUMAN
4147	SLSESSVIMDR202002000000	sp P13051 UNG_HUMAN
4148	AQSPGAVEEILDR200000000000	sp Q9H2Y7 ZN106_HUMAN
4149	LEVTEIVKPSPK2000	sp Q9NYM9 BET1L_HUMAN
4150	SRSPLLEPEAK200000000000	sp Q7Z6E9 RBBP6_HUMAN
4151	HEAPSSPISGQPCGDDQNASPSK2E+19	sp Q92466 DDB2_HUMAN
4152	KDDSHAEDSEDEKEDHK2000002000000000	sp O75376 NCOR1_HUMAN
4153	TPQSPTLPPAK200000000	sp Q9H1E3 NUCKS_HUMAN
4154	SRSSPVTELASR20002000000000	sp P78310 CXAR_HUMAN
4155	KQHLYSSEDEPDNDVLDLDR2200000000000000	sp Q9UQ35 SRRM2_HUMAN
4156	LPSSGLPAADVSPATAEEPLSPSTPTR20200000	sp Q8NFC6 BD1L1_HUMAN
4157	FSISPDESSSYSSNSDFNYSYPTK20200000000000	sp O43149 ZZEF1_HUMAN
4158	VISDESIDIGGSDVEFKPDK2. 0002E+19	sp Q96QD8 S38A2_HUMAN
4159	SPAGSPELR2000000000	sp P52701 MSH6_HUMAN
4160	GPPDFSSDEEREPTPVLGSGAAAAGR2. 0000002E+20	sp Q96JM3 CHAP1_HUMAN
4161	DGTAPPQSPGSPGTGQDEEWSDEESPR2. 0000000000002E+20	sp P42166 LAP2A_HUMAN
4162	LGPGRPLPTFPTSECTSDVEPDTR2000000000	sp P42167 LAP2B_HUMAN
4163	SPSAEFSPAAPPGISSIHSPSLR2E+23	sp Q9UKS6 PACN3_HUMAN
4164	ELEEVSPETPVVATTQR20000000000000	sp Q8TDD1 DDX54_HUMAN
4165	KPPAPPSPVQSQSPSTNWSPAVPVKK2. 000000002002E+20	sp Q68DK2 ZFY26_HUMAN
4166	ISMQDVDSLGLSPK2000	sp Q15149 PLEC_HUMAN
4167	GSFSSSENYWR2000000000	sp Q8IYB3 SRRM1_HUMAN
4168	STGSATSLASQGER200000200000	sp Q09666 AHNK_HUMAN
		sp Q6P6C2 ALKB5_HUMAN
		sp Q5SW79 CE170_HUMAN



4169	DSHSSEEDEASSQTDLSQTISK2. 022E+21	sp Q5JTV8 TOIP1_HUMAN
4170	AAALQALQAQAPTSPPPPPPLK20000000000	sp Q8NAF0 ZN579_HUMAN
4171	KRESESEDETTPAAPQLIK202000020000000000	sp 060832 DKC1_HUMAN
4172	EVSPSDVR2000000	sp Q8WYP5 ELYS_HUMAN
4173	VTLYNTDQDGS DSPR2000	sp Q9UPW0 FOXJ3_HUMAN
4174	SPPGSGASTPVGPWDQAVQR20000000000000	sp Q8TE77 SSH3_HUMAN
4175	DGETLEGSDAEESL DK2000000000	sp Q9P2D1 CHD7_HUMAN
4176	TNSSDSERSPDLGHSTQIPR220200000000000000	sp Q6VY07 PACS1_HUMAN
4177	SVCPGSSSPGSSSGGR20000000	sp Q53LP3 SWAHC_HUMAN
4178	GIPHSASPVSPDGVIPLK2002000000000000	sp Q6PJF5 RHDF2_HUMAN
4179	KETPPPLVPPAAR200000000000	sp Q9BQA1 MEP50_HUMAN
4180	AKPAMPQDSVPSPR10000002000	sp P53396 ACLY_HUMAN
4181	LSSWDQAETPGHTPSLR2000200000	sp 075533 SF3B1_HUMAN
4182	NAEAVLQSPGLSGK20000000	sp Q13045 FLII_HUMAN
4183	CPFPAGAALACSEDEEDDEEHEGGSR2000000000000000	sp Q147X3 NAA30_HUMAN
4184	VESLESLYFTPIPAR20000000000000	sp Q14980 NUMA1_HUMAN
4185	AAHTEDINACTLTTSPR20000	sp P04049 RAF1_HUMAN
4186	SSGSPYGGYGGSGGGSGYGSR300000000000000000	sp P51991 ROA3_HUMAN
4187	SQSSEGVSSLSSSPNSLETQSQSLSR2E+19	sp 095155 UBE4B_HUMAN
4188	TETLVEETAAESSEDDVAPSQSLLSGYMTPGLTPANSQASK2. 2E+31	sp Q13428 TCOF_HUMAN
4189	TDASSASSFLDSELER220000000000	sp Q14498 RBM39_HUMAN
4190	SRSSSPVTELASR20200000000000	sp Q9UQ35 SRRM2_HUMAN
4191	CPEILSDESSSDEDEKK20002200000000	sp P35659 DEK_HUMAN
4192	SPTGPSNSFLANMGTVAHK2. 000000000001E+20	sp Q96I25 SPF45_HUMAN
4193	VASSPVMVSNPATR200000000000	sp P51610 HCFC1_HUMAN
4194	SPGPPPVGVK200000000000	sp Q9H1B7 I2BPL_HUMAN
4195	QVSSPSFTNR20000000	sp Q86V48 LUZP1_HUMAN
4196	LKSDERPVIHK2000000000	sp Q9UPN9 TRI33_HUMAN
4197	SQSESSDEVTELDLSHGK200200000000000000	sp Q86YS7 C2CD5_HUMAN
4198	NYAGEEEEEEGSGSSEGFDPATDR2020000000000000	sp P16989 YBOX3_HUMAN
4199	SQSPVHLR202000000	sp Q96T58 MINT_HUMAN
4200	GDSESEDEDELEVVPVPSR200000000000000000	sp P13861 KAP2_HUMAN
4201	ANTLSPVR20000	sp Q9H792 PEAK1_HUMAN
4202	TSAACA VTDLSDDSD FDEK2000000	sp Q9NQZ2 SAS10_HUMAN
4203	IASP VSR200000	sp P09884 DPOLA_HUMAN
4204	STSAPQMSPGSSDNQSSSPQPAQK1. 0002E+19	sp Q14157 UBP2L_HUMAN
4205	STDSPGADAELPES AAR2000000000000000	sp Q9Y4K1 AIM1_HUMAN
4206	KEPDDSRDEDEDEDESSEEDSEDEEPPPK2200020000000000	sp Q5SSJ5 HP1B3_HUMAN
4207	EMEHTVCAAGTSPVGEIGEEK1. 00000000002E+21	sp P18583 SON_HUMAN
4208	GDSLKEPTSIAESSR20000000000000	sp Q9C0B5 ZDHC5_HUMAN
4209	STSPIIGSPVVR200000000000	sp Q86TB9 PATL1_HUMAN
4210	SGSSSPDSEITELKFPSINHD200000000000000000	sp P17812 PYRG1_HUMAN
4211	ASPPASPIAFPPAFEAQVEAKPDELK2. 0002E+27	sp Q8NEZ4 KMT2C_HUMAN
4212	NSSPAPPQPAPGK20000000000000	sp Q9NYV4 CDK12_HUMAN
4213	NSSDSAIDNPKPNK20000000000000	sp P46100 ATRX_HUMAN
4214	GHTDTEGRPPSPPTSTPEK220000	sp Q00613 HSF1_HUMAN
4215	LKSVEDEMDS PGEEPFTGQGR2000000000000000	sp Q12857 NFIA_HUMAN
4216	TASRPDDIPDPS SPSK2002000	sp Q5VZK9 LR16A_HUMAN
4217	DHAPSIPTSPFGDSLESSTELR2000000000000000	sp Q9Y4B5 MTCL1_HUMAN
4218	SLSSSPVKK2000200000	sp Q9H3Q1 BORG4_HUMAN
4219	SNSDES NFSEK2000000000	sp 095999 BCL10_HUMAN
4220	ELSPAGSISK200000000	sp 095453 PARN_HUMAN
4221	SPTTASVSATAAPR20000000000000	sp Q15723 ELF2_HUMAN

4222	LASPELER2000000	sp P17535 JUND_HUMAN
		sp P05412 JUN_HUMAN
4223	EDNSSEEEEEIEPFPEER2200000000000000	sp Q4LE39 ARI4B_HUMAN
4224	SVDAIGGESMPIPTIDTSR2E+19	sp Q9BWT3 PAPOG_HUMAN
4225	LGHPEALSAGTGSPQPPSFTYAQQR20000000000000	sp Q15942 ZYG_HUMAN
4226	CPEILSDESSSDEDEKK2220000000	sp P35659 DEK_HUMAN
4227	ASESSKPWPDATYGTGSASR20000	sp Q9Y2W1 TR150_HUMAN
4228	DCSYGAVTSPTSTLESR2000000000	sp Q9HCD6 TANC2_HUMAN
4229	LCDPDGLSDEEDQKPVR20000000000	sp Q96RY5 CRML_HUMAN
4230	GYSPYSVSGSGSTAGSR2200000	sp Q15149 PLEC_HUMAN
4231	KASATCSSATAAASSGLEEWSR2000000000000000	sp Q96G74 OTUD5_HUMAN
4232	MQNTDDEERPQLSDDER200000000200000	sp Q8WVC0 LEO1_HUMAN
4233	AASDLLSSVSPAVSK200000200	sp P19532 TFE3_HUMAN
4234	SESSDFEVVVK200200000000	sp Q9Y520 PRC2C_HUMAN
4235	EEGPPPPSPDGASSDAEPEPPSGR2200000000000	sp Q9NTJ3 SMC4_HUMAN
4236	ENASPAPGTTAEEMSR200000000001000	sp P46379 BAG6_HUMAN
4237	HQQQLLASPGSSTVDNK2000000	sp Q9C0E2 XPO4_HUMAN
4238	TLSDPPSPLPHGPPNK2000200000000000	sp Q9ULH1 ASAP1_HUMAN
4239	HTGMASIDSSAPETTSDSPTLSR2000000	sp A7E2V4 ZSWM8_HUMAN
4240	SLSGSSPCPK200200000	sp Q9UQ35 SRRM2_HUMAN
4241	SRSGEGEVSGLMR200000000100	sp Q13263 TIF1B_HUMAN
4242	CVSMPGDISGLQGGPR2000000000000000	sp Q8IY17 PLPL6_HUMAN
4243	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2.002E+32	sp Q8WUZ0 BCL7C_HUMAN
4244	AEEKSPISINVK200000000	sp Q9UK58 CCNL1_HUMAN
4245	SVSGSPEPAAK202000000000	sp Q8IYB3 SRRM1_HUMAN
4246	ASSFSGISVLTR200000000000	sp Q15032 R3HD1_HUMAN
4247	HKSDNETNLQQQVVWGNR2000000000000000	sp Q7Z401 MYCPP_HUMAN
4248	IKEEVLESEEAENQQAGAAALAPEIVIK2E+22	sp Q9UJX5 APC4_HUMAN
4249	HSDNPSEEGEVKDDGLEK200020000000000000	sp O14647 CHD2_HUMAN
4250	ESVSTEDLSPPSPPLPK2000000	sp Q13127 REST_HUMAN
4251	SQASMVINSDDTDEEEVSAALTLAHLK2.1000002E+24	sp Q14676 MDC1_HUMAN
4252	LVGATATSSPPPK200000	sp Q96QD9 UIF_HUMAN
4253	GQSQLSNPTDDSWK2000000000	sp Q9Y6D5 BIG2_HUMAN
4254	DLHSDNNEESDEVEDNDNDEMERPVNR2.0000002E+26	sp P42568 AF9_HUMAN
4255	RGSVSSVSPKPPSSFK2000020000000000	sp Q9NZJ0 DTL_HUMAN
4256	SPEPEVLSTQEDLFDQSNK200000000000	sp Q12888 TP53B_HUMAN
4257	RESESEDETPPAAPQLIK202000000000000000	sp O60832 DKC1_HUMAN
4258	SIKSDVPVYLK200200000000	sp Q5SW79 CE170_HUMAN
4259	ELVGDGTSQEGDHEPSGSETEEDTSSSPHR2020000000000	sp PODJ93 SIM13_HUMAN
4260	EENIKDSSGQEDETQSSNDPSQSVASQDAQEIRPR2.00000202E+30	sp Q8N108 MIER1_HUMAN
4261	VLDTSSLTQSAPASPTNK2000000000000000	sp Q8N122 RPTOR_HUMAN
4262	SHSPSPDPDTPSPVGDSR200000000000000000	sp Q13586 STIM1_HUMAN
4263	KSSEGGVGVGGGDEPPTS2E+21	sp Q9Y4H2 IRS2_HUMAN
4264	LSPEVAPPAHR20000000000	sp Q8TAD8 SNIP1_HUMAN
4265	DLSPQHMVVR200000000	sp P49590 SYHM_HUMAN
4266	GPSPAPAPAASPQPSR2000000000000000	sp Q8IX07 FOG1_HUMAN
4267	TPVDESDEIQHDEIPTGK2E+19	sp Q86TC9 MYPN_HUMAN
4268	GPSSPPTTAPMR20000000000	sp Q9NXL9 MCM9_HUMAN
4269	GLLSDSMTDVPVDTGVAAR2100000000000000	sp O43399 TPD54_HUMAN
4270	HQDGLPYIDDSPPSSPHLSSK220000000	sp P11274 BCR_HUMAN
4271	SASWGSTDLK2000000000	sp Q6P1L5 F117B_HUMAN
4272	SWDSSSPVDRPEPEAASPTTR2E+21	sp Q86WB0 NIPA_HUMAN

4273	IGEGTYGVVYK3000000	sp Q00526 CDK3_HUMAN
		sp P06493 CDK1_HUMAN
		sp P24941 CDK2_HUMAN
4274	SVSVDSGEQR200000000	sp Q86UU0 BCL9L_HUMAN
4275	EVSDDEAEEK200000000	sp P07900 HS90A_HUMAN
4276	ENQEPLRSPEVGDDEALRPLTK20000000000000000000	sp P48681 NEST_HUMAN
4277	SSSVTTSETQPCTPSSSDYSDLQR2E+22	sp P50552 VASP_HUMAN
4278	GSPATSPHLGR20000000000	sp Q13671 RIN1_HUMAN
4279	TRLSTASEETVQNR220000000000	sp O43379 WDR62_HUMAN
4280	MAEESSSSSSSSPTAATSQQQLK2E+20	sp Q13620 CUL4B_HUMAN
4281	GFGFGQGAGALVHSE200	sp P21291 CSRP1_HUMAN
4282	AEQSPPNDSGQDAHPDPDANPDAAR2.0002E+23	sp Q9BUA3 CK084_HUMAN
4283	KA AVLSDSEDEEK202000000	sp Q96ST2 IWS1_HUMAN
4284	VSSPAPMEGEEEEELLGPK20000000000000000000	sp O14497 ARI1A_HUMAN
4285	DVDDGSGSPHSPHLSSK202002000000000	sp Q9H4A3 WNK1_HUMAN
4286	SPTPPSSAGLGSNSAPPIDSR200002020000000000	sp Q8IWX8 CHERP_HUMAN
4287	HVTLPSSPR20002000	sp Q5T4S7 UBR4_HUMAN
		sp P24941 CDK2_HUMAN
4288	IGEGTYGVVYK3000000	sp P06493 CDK1_HUMAN
		sp Q00526 CDK3_HUMAN
4289	LSTTPSPTSSLHEDGVEDFR2.00000022E+19	sp O94842 TOX4_HUMAN
4290	DPHSPEDDEEQPQGLSDDDILR20000000000020000000	sp Q86VM9 ZCH18_HUMAN
4291	KRPSLPSSPSPGLPK22000000000	sp O43294 TGF11_HUMAN
4292	HIVSNDSDDDESHEPK202000000000	sp P47712 PA24A_HUMAN
4293	ESLSTHTSPSQSPNSVHGSR202000000000000	sp Q9H1K0 RBNS5_HUMAN
4294	SSSLLASPGHISVK220000000000000	sp A0FGR8 ESYT2_HUMAN
4295	GLFQDEDCSDCSYR200000000	sp Q8NEM2 SHCBP_HUMAN
4296	MLPHAPGVQMAIPEDAVHEDSGDEDGEDPDKR1.000000001E+33	sp Q92769 HDAC2_HUMAN
4297	LGSQHSPGR20000	sp Q8IZP0 ABI1_HUMAN
		sp Q9UQ88 CD11A_HUMAN
4298	GTSRPPPEGGLGYSQLGDDDLK2.2E+21	sp P21127 CD11B_HUMAN
		sp P35659 DEK_HUMAN
4299	KESEEDSSDDEPLIK2000022000000000	sp O60238 BNI3L_HUMAN
4300	DHSSQSEEEVVEGEK220200000000000	sp Q9NTI5 PDS5B_HUMAN
4301	LDSEMDSSENYDTMSSPLPGK20000300000000000	sp Q6PIJ6 FBX38_HUMAN
4302	TVNSGGSSEPSPTVDVSR20020000000000	sp Q9C0B5 ZDHC5_HUMAN
4303	SSSLKSAQGTGFEGLQLQSIR2000200000000000000	sp Q9Y4E1 FA21C_HUMAN
4304	GLFSDEEDSEDLFSSQSASNLK2000000000000000	sp P49756 RBM25_HUMAN
4305	LGASNSPGQPNVSVKR20000020000	sp Q3YEC7 RABL6_HUMAN
4306	GSPPLPAGVPVSDITLSSEEEAEVAAPTK2.00000000020002E+29	sp P00558 PGK1_HUMAN
4307	ALESPERPFLAILGGAK2000000000000000	sp Q9BRD0 BUD13_HUMAN
4308	YEYDPDISPPR20000	sp Q9P1Y6 PHRF1_HUMAN
4309	AEAPSSPDVAPAGK200000000000	sp Q92805 GOGA1_HUMAN
4310	SVKESVASMADSGDDFASDGSSSR2E+21	sp Q9UQ35 SRRM2_HUMAN
4311	SGSSQELDVKPSASPQER20020000000000000000	sp P24928 RPB1_HUMAN
4312	YSPTSPTYSPVYTPSPK3000000000002000	sp Q8WWI1 LMO7_HUMAN
4313	SHSPSASQSGQLR2000000000000000	sp Q9NR09 BIRC6_HUMAN
4314	SPATSPISSNSHR2000000000000000	sp Q9GZR1 SEN6_HUMAN
4315	ESISPQPADSACSSPAPSTGK20020000000000	sp Q6PKG0 LARP1_HUMAN
4316	TASISSPSEGTPVGSYGCTPQSLPK2.202E+22	sp Q13247 SRSF6_HUMAN
4317	SVSPPPKR202000000	sp Q14202 ZMYM3_HUMAN
4318	GWAPPGPSSSGALDLLDTPAGLEK20000000000000000000	sp Q6Y7W6 PERQ2_HUMAN
4319	ALSSGGSITSPPLSPALPK20000000002000000	sp P27816 MAP4_HUMAN
4320	DMSPLSETEMALGK20020000000000	sp O60832 DKC1_HUMAN
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4322	SQSSGSSATHPI SVPGAR200000000000	sp Q9Y4H2 IRS2_HUMAN
4323	IQSSLSVNSDISK200	sp Q9P1T7 MDFIC_HUMAN
4324	DGSDYDPYDFSDTEEMPQVHTPK2E+21	sp P25205 MCM3_HUMAN
4325	GFNGCDSPEPDGSDSLEQSPILLEDK2. 00000002E+19	sp Q14814 MEF2D_HUMAN
4326	SSTPLHSPSPIR20002000000	sp 095817 BAG3_HUMAN
4327	ESSEPPAPASSPEAASPTEQGPAGTSK2200002000000000000	sp Q92766 RREB1_HUMAN
4328	KQDSSDEDDDDDEAPK22000000000000	sp 014647 CHD2_HUMAN
4329	SRSSSPVTELASR202000000000	sp Q9UQ35 SRRM2_HUMAN
4330	TVSSPPTSPRPGSAATVSASTSNIIPPR2. 2002E+26	sp 076094 SRP72_HUMAN
4331	YSDGDGKSDDLVVDVSNEDPATPR200000000000002000	sp Q04726 TLE3_HUMAN
4332	SSSLGSYDDEQEDLTPAQLTR2E+19	sp 094988 FA13A_HUMAN
4333	SRDATPPVSPINMEDQER2000200020001000000	sp P17275 JUNB_HUMAN
4334	LDSQPQETSPELPR2000000	sp 014545 TRAD1_HUMAN
4335	GPSPAPASSPKR20000020000	sp Q9H7N4 SFR19_HUMAN
4336	GSDALSETSSVSHIEDLEK200220200000000	sp Q6P996 PDXD1_HUMAN
4337	SPPVQPHTPVTISLGTAPSLQNNQPVEFNHAINVYVK2E+37	sp Q96ST3 SIN3A_HUMAN
4338	NEIIQSPISQVPSVEK200000020000	sp Q7Z3T8 ZFY16_HUMAN
4339	RASPPPPKR202000000	sp Q8IYB3 SRRM1_HUMAN
4340	GDSVSDSGSDALR200000000000	sp Q53EL6 PDCD4_HUMAN
4341	AGDPGEMPQSPTGLGQPK1002000000000	sp Q9UIF9 BAZ2A_HUMAN
4342	FVVIQNEDLGPASPLDSTFYR2000000000	sp P04626 ERBB2_HUMAN
4343	SPLLAGGSPQPVVPAHK2000000200000000000	sp Q8NFH5 NUP53_HUMAN
4344	SPSSLSANI ISSPK200000000002000	sp Q8NFA0 UBP32_HUMAN
4345	SRPTSEGSDIESTEPQK20020000000000	sp Q4G0J3 LARP7_HUMAN
4346	AGSNEDPILAPSGTTPPTIPPDEFGGR200000000000000000	sp Q8IZL8 PELP1_HUMAN
4347	SPAVATSTAAPPPSSPLPSK2000000000000000	sp Q14157 UBP2L_HUMAN
4348	RQSPEPSPVTLGR200020000000	sp Q9NRL2 BAZ1A_HUMAN
4349	RESESEDETPPAAPQLIK202020000000000000	sp 060832 DKC1_HUMAN
4350	SGDEMIFDPTMSK20000000000000	sp P20042 IF2B_HUMAN
4351	TAQVSPPPR20000	sp Q9UKV3 ACINU_HUMAN
4352	AWLDEDSNLSPSPLR20000	sp Q9C086 IN80B_HUMAN
4353	STGSATSLASQGER20000000000000	sp Q5SW79 CE170_HUMAN
4354	SRDSGDENEPIQER2002000000000000	sp Q8WX93 PALLD_HUMAN
4355	AAGGAPSPPPPVR20000000	sp Q4KMP7 TB10B_HUMAN
4356	SDPDGGDSPLPASGGPLTCK2E+20	sp Q8N1G0 ZN687_HUMAN
4357	FGDSSSDNGFSSTGSTPAKPTVEK2. 202E+22	sp Q8IU60 DCP2_HUMAN
4358	STPSPVLQPR2020000000	sp Q8TEK3 DOT1L_HUMAN
4359	SGSSSPDSEITELK22000000000000	sp P17812 PYRG1_HUMAN
4360	SASWGSADQLK2000000000	sp Q86VQ1 GLC11_HUMAN
4361	SPSPYYSR202000000	sp P62995 TRA2B_HUMAN
4362	QLSSGVSEIR20000000	sp Q13595 TRA2A_HUMAN
4363	SEVQQPVHPKPLSPDSR200000	sp P04792 HSPB1_HUMAN
4364	AEEQQLPPPLSPPSPSTPNHR200200000000	sp Q9UHB6 LIMA1_HUMAN
4365	KLPSDSGDLEALEGK20000000000000	sp Q6WCQ1 MPRIP_HUMAN
4366	LGSVEAPK2000000	sp 060524 NEMF_HUMAN
4367	QQAAYYAQTSPQGMPPHPPAPQGQ2000000000000000	sp P46087 NOP2_HUMAN
4368	ENSFSGSPLEFR2002000000	sp Q96AE4 FUBP1_HUMAN
4369	RPDHSGGSPSPPTSEPAR2002000000000000	sp Q9UQ35 SRRM2_HUMAN
4370	GSSDGRGSDSEDLPHR2000000200000000	sp Q8TAD8 SNIP1_HUMAN
4371	DNSISPESDVTASTISLVTGETER2. 02002E+23	sp Q9UPQ0 LIMC1_HUMAN
4372	EESDDEDEKESEEEPPKK22000000200000000	sp Q5SW79 CE170_HUMAN
4373	ALSSGGSITSPPLSPALPK220000000000	sp P35659 DEK_HUMAN
4374	EVEEDSEDEEMSEDEEDDSSGEEVVIPQK2. 00000200000022E+24	sp Q6Y7W6 PERQ2_HUMAN
		sp P19338 NUCL_HUMAN

4375	STSSSDTMSTFKPMVAPR220000000000000000	sp Q7Z6B7 SRGP1_HUMAN
4376	LDSSEMDHSENYDTMSSPLPGK2.0010020000001E+21	sp Q9NTI5 PDS5B_HUMAN
4377	SPSSLSANI ISSPKGSPSSSR220002000000	sp Q8NFA0 UBP32_HUMAN
4378	KPSVGVPPPASPSYPR2000000	sp Q9P206 K1522_HUMAN
4379	CQTAEADSESDHEVPEPESEM2020000000000000	sp Q9BPX3 CND3_HUMAN
4380	TAVQYIESSDSEEIETSELPQK30002020000000000000	sp Q14527 HLTF_HUMAN
4381	SSSVGSSSYPI SPAVSR22200020000000	sp Q15149 PLEC_HUMAN
4382	TDALTSSPGR2020000	sp P49736 MCM2_HUMAN
4383	LLEDSEDEEAAPSPLQPALRPNPTAILDEAPKPK2.02E+31	sp Q9BW85 CCD94_HUMAN
4384	APSTSPSFEGTQETYTVAHEENV20000000000000	sp Q9BUT9 F195A_HUMAN
4385	EKEEEEEPEGKPSDAESSESDDEKAWVEEVR2.00022022E+20	sp Q9BSC4 NOL10_HUMAN
4386	DATPPVSPINMEDQER200020001000000	sp P17275 JUNB_HUMAN
4387	HSVTGYGDCAVGAR2000000000000000	sp Q96IF1 AJUBA_HUMAN
4388	GSPPLPAGVPVSDITLSSEEEAEVAAPTK2.0000000002E+29	sp Q3YEC7 RABL6_HUMAN
4389	ISYTPPEPVPVSYASSTPLHVPVPR2.0002E+22	sp P41212 ETV6_HUMAN
4390	AGGSPAPGPETPAISPSK20000	sp P33316 DUT_HUMAN
4391	GRDDPGQQTDSSEDEDIIGMPAK20220000000010000	sp Q8IXQ4 GPAM1_HUMAN
4392	SSSPVTELASR200000000000	sp Q9UQ35 SRRM2_HUMAN
4393	RYPSSISSPQK20000	sp Q14157 UBP2L_HUMAN
4394	SGPKPFSAPKQTSPPSK202000	sp Q01518 CAP1_HUMAN
4395	DSPSKSSAEAQTPEDTPNK200000000000000000	sp 043493 TGON2_HUMAN
4396	TLSPTPSAEGYQDVR200000000000000000	sp 014639 ABLM1_HUMAN
4397	AASPESASSTPESLQAR200202000000000000	sp Q96BT3 CENPT_HUMAN
4398	TASGSSVTSLDGTR2022000000	sp Q92597 NDRG1_HUMAN
4399	SLVGTPTYWMAPELISR200000000000000000	sp 096013 PAK4_HUMAN
4400	TPLALAGSPTPK2000	sp 095785 WIZ_HUMAN
4401	HPASDSEIEELQK20200000000	sp Q96ST2 IWS1_HUMAN
4402	DTSPDKGELVSDEEEDT20000000	sp Q9UPU7 TBD2B_HUMAN
4403	AVPTWASVQVDDPGSLASVESPGTPK2000000	sp Q8IVT2 MISP_HUMAN
4404	LPQLVGVSTPLQGGNSAAAIGQSSGELR200000	sp P15291 B4GT1_HUMAN
4405	LGGLRPESPELTSVSR200200000000	sp Q9H6F5 CCD86_HUMAN
4406	GRLTPSPDIIVLSDNEASSPR2020000000000020000	sp Q8WXI9 P66B_HUMAN
4407	GLAGPPASPGK20000	sp Q9P206 K1522_HUMAN
4408	TSDIFGSPVTATSR200000000	sp Q9H910 HN1L_HUMAN
4409	SPAVATSTAAPPPSSPLPSK20000000	sp Q14157 UBP2L_HUMAN
4410	GPPSPAPVMHSPSR20000000000000	sp Q13573 SNW1_HUMAN
4411	ATSPSSVSGDFDDGHHSVSTPGPSR2E+24	sp Q86U86 PB1_HUMAN
4412	SSGTASSVAFTPLQGLEIVNPQAAEK2.02E+25	sp Q8WWY3 PRP31_HUMAN
4413	RGSLPDTQPSQGPSTPK2000000000000000	sp Q86Y91 KI18B_HUMAN
4414	LSSPVLHR2000000	sp Q16643 DREB_HUMAN
4415	IYQYIQR300000	sp Q9Y463 DYR1B_HUMAN
4416	QSLSSADNLESDAQGHQVAAR200000000000000000	sp Q13627 DYR1A_HUMAN
4417	KPSGSPDLWK2000000	sp Q9Y6D5 BIG2_HUMAN
4418	AAEMHQLKDSSR2000	sp Q96JM3 CHAP1_HUMAN
4419	KGGSYSQAASSDSAQGSVSLTACK2002000000	sp Q96BD0 SO4A1_HUMAN
		sp P05534 1A24_HUMAN
		sp Q09160 1A80_HUMAN
		sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
4420	DSSSTDSANEKPALIPR200200000000000000	sp Q9NUQ6 SPS2L_HUMAN
4421	GLPSPYNMSSAPGSR20000200	sp P15924 DESP_HUMAN
4422	LSPQDPRPLSPGALQLAGEK2.00000002E+19	sp Q8TEK3 DOT1L_HUMAN

4423	SLPGPAPCLK20000000000	sp Q01433 AMPD2_HUMAN
4424	LIPGPLSPVAR200000	sp P48634 PRC2A_HUMAN
4425	SPGSPVCR200200000	sp Q86VQ1 GLC11_HUMAN
4426	SPSEESAPTTSPESVSGVPSSGSSGR2000000000000	sp P54725 RD23A_HUMAN
4427	HASSSPESPKPAPAPGSHR22002000000000000	sp Q9UQ35 SRRM2_HUMAN
4428	HSVTAATPPPSPTSGESGDLLSNLQSPSSAK2. 2E+20	sp Q96QU8 XPO6_HUMAN
4429	VETVSQPSESPKDTIDK20000200000000000	sp Q99590 SCAFB_HUMAN
4430	FNSYDISR2000000	sp Q6IQ26 DEN5A_HUMAN
4431	QASPETSASPDGSQNLVYETELLR20000000000000000	sp Q14789 G0GB1_HUMAN
4432	QRGSETGSETHESDLAPSDK20200000000000000	sp P48634 PRC2A_HUMAN
4433	AAAPPQSPPPPTK2000000	sp P49848 TAF6_HUMAN
4434	GAASPVLQEDHCDLPSVLQVEEK2E+21	sp Q8N9B5 JMY_HUMAN
4435	GTEPSPGGTPQPSRPVPAGPPEGVPEEAQPPR2. 00000002E+25	sp Q8WUZ0 BCL7C_HUMAN
4436	NLVSPAYCTQESR200	sp Q9NUQ3 TXLNG_HUMAN
4437	TEQELPRPQSPSDLSDLGR200000200000	sp P54259 ATN1_HUMAN
4438	TGVTSTSDSEEGDDQEGEKK20200000000000000	sp 075475 PSIP1_HUMAN
4439	QEPIESDEEEAEPPR20000000000	sp P56524 HDAC4_HUMAN
4440	RGSPVPPVPER2000000000	sp Q6ZRV2 FA83H_HUMAN
4441	QNSQLPAQVQNGPSQEELEIQR2E+20	sp Q8N8S7 ENAH_HUMAN
4442	VTETEDSDSDSDDEDDVHVTIGDIK2. 0202E+20	sp Q6UN15 FIP1_HUMAN
4443	DDEKEAEEGEDDRDSANGEDDS200000000	sp P07910 HNRPC_HUMAN
4444	APSDSSLGTPSDGRPELR20000000000000000	sp Q15642 CIP4_HUMAN
4445	DLFDLNSSEEDDTEGFSEK200000000000000	sp Q9Y3T9 NOC2L_HUMAN
4446	SPTLSQVHSPLVTSANLK20002000000000	sp Q86UU0 BCL9L_HUMAN
4447	SLHGFEFSASDSSLTDIQETR2022000000000	sp 043166 SIL1_HUMAN
4448	SSTPLPTISSAENTR220000000	sp P42166 LAP2A_HUMAN
4449	LKEDIENEDEQNSPPK20000	sp P42167 LAP2B_HUMAN
4450	SPGSTPTTSSQAPQK202200000000000	sp Q9NTI5 PDS5B_HUMAN
4451	VPVQEIDDDFFPSSGEEAEASVGEVGGGGGRK2. 2E+20	sp P35658 NU214_HUMAN
4452	EGEETVYSDEEEKDESAR20000000000000000	sp Q03468 ERCC6_HUMAN
4453	EPAPLLSAQYETLSDSDD202000	sp 000264 PGRC1_HUMAN
4454	NAIASDSEADSDTEVVK200000200000000	sp 075376 NCOR1_HUMAN
4455	LEEPINEETPSQDEEEGVSDVESER2000000020002000	sp Q8WVC0 LEO1_HUMAN
4456	SSSPILTKPTLK220000000000000	sp Q9BYW2 SETD2_HUMAN
4457	LTSVSSVDFDQR22200000000	sp Q92794 KAT6A_HUMAN
4458	SAASPVSSMPER200000000000	sp Q5T3F8 CSCL2_HUMAN
4459	GPPSPAPVMHSPSR2000001020000	sp P18583 SON_HUMAN
4460	ESSTESSQSAKPVSGQDTSGNTEGSPAEEK2000000	sp Q13573 SNW1_HUMAN
4461	ESKDKPEIEDVGSDEEEK20000000	sp Q96S66 CLCC1_HUMAN
4462	FLETDSEEEQEEVNEKK20200000000000000	sp Q14568 HS902_HUMAN
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4464	GRLTPSPDIVLSDNEASSPR2000000000000000000	sp Q5VT52 RPRD2_HUMAN
4465	SRSPLDKDTYPPSASVVGASVGGHR2. 0000000002E+23	sp Q8WXI9 P66B_HUMAN
4466	YTPTSPSYSPSPEYTPSPK2. 000003E+20	sp Q96T37 RBM15_HUMAN
4467	GAAEEEQQDSGSEPRGEDDIPLGPQSNVSLLDQHHLK2. 02E+30	sp P24928 RPB1_HUMAN
4468	DAPAGSPASPQGPSTK20002000000000000	sp Q9UJV9 DDX41_HUMAN
4469	VVEAVNSDSDSEFGIPKK20202000000000	sp Q3MIN7 RGL3_HUMAN
4470	AADSSPDTSDNEAKEEIK22002200000000000	sp Q02880 TOP2B_HUMAN
4471	AASPSQSVR202000000	sp Q9NYD6 HXC10_HUMAN
4472	SVENLPECGITHEQR20000000000000000	sp Q8IYB3 SRRM1_HUMAN
4473	SGFGEISSPVIR200000	sp Q9UBF8 PI4KB_HUMAN
4474	EPVNLEGDDTSLIHGK2000000	sp P57740 NU107_HUMAN
4475	AGSNEDPILAPSGTTPPTIPPDETFGGR20000000000000000	sp Q9H2J7 S6A15_HUMAN
		sp Q8IZL8 PELP1_HUMAN

4476	RGEGDAPFSEPGTTSTQRPSSPETATK20000000	sp Q9UQ35 SRRM2_HUMAN
4477	DAAAHLQTSBKPSAEDAEGQSPLSQK2000000	sp Q96F24 NRBF2_HUMAN
4478	SRSRSTPPSSAGLGSNSAPPIDSR2. 0202E+26	sp Q8IWX8 CHERP_HUMAN
4479	KRSPSPSPTPEAK2000000000	sp Q8TAQ2 SMRC2_HUMAN
4480	EAARSPDKPGGSPSASR20000000000000	sp Q9Y3T9 NOC2L_HUMAN
4481	TPQGPSPTNTPLPEAAVSQPGSAVASDWR2E+26	sp Q9Y4R8 TELO2_HUMAN
4482	SAPASPTHPLMSPR2000000000002000	sp P85037 FOXK1_HUMAN
4483	LNQSSPDNVTDTK2000000000	sp Q08AD1 CAMP2_HUMAN
4484	SGDETPGSEVPGDK200200000000	sp Q1KMD3 HNRL2_HUMAN
4485	EFITGDVEPTDAESEWHSENEEEEEK200000000	sp Q99733 NP1L4_HUMAN
4486	KPGDGEVSPSTEDAPFQHSPLGK20000000000000	sp 060318 GANP_HUMAN
4487	SEPVINNDNPLESNDEK200000	sp P23497 SP100_HUMAN
4488	TTAPTEGKSPLKK200000	sp Q96T37 RBM15_HUMAN
4489	VSPAPGSAPEPTPEDK2000200000	sp Q13459 MYO9B_HUMAN
4490	AKSPTPDGSER2020000000	sp Q96MU7 YTDC1_HUMAN
4491	GEAERGGSSGDELREDDEPVK200220000000000000	sp Q7Z4V5 HDGR2_HUMAN
4492	SMSPTSTPK2020000000	sp 094913 PCF11_HUMAN
4493	GSFSDTGLGDGK2000000000	sp Q9Y2W1 TR150_HUMAN
4494	HGSVSADEAAR2000000000	sp Q9NWW5 CLN6_HUMAN
4495	HSHSHSPMSTR20202000000	sp P62995 TRA2B_HUMAN
4496	SSTPLPTISSAENTR20000000000	sp P42167 LAP2B_HUMAN
4497	TLSNAEDYLDDESD200	sp P42166 LAP2A_HUMAN
4498	KRPSLPSSPSPGLPK2000200000000	sp Q92882 OSTF1_HUMAN
4499	YLVDTGKPNAGSEEISSEDELVEEKK200022000000000000	sp 043294 TGF11_HUMAN
4500	AASSSPGSPVASSPSR200	sp Q9NY61 AATF_HUMAN
4501	SSSFTPFIDPR22000000000	sp Q92538 GBF1_HUMAN
4502	LPPTFFCGVCSDTDEDNGNGEDFQSELQK200000000000000000	sp 095819 M4K4_HUMAN
4503	LGFVDVQNCISR200	sp P49792 RBP2_HUMAN
4504	SRSSSPVTELASR22000000000	sp Q14573 ITPR3_HUMAN
4505	SGLELTELQNMTPEDDNISNDSNDFTEVENGQINSK2. 00000002E+26	sp Q9UQ35 SRRM2_HUMAN
4506	SEPSGEINIDSSGETVGSGER220000000000	sp Q9H2H9 S38A1_HUMAN
4507	GGSYVFQSDGPEPEAEESDLDSGSVHSASGRPDGPVR2E+20	sp P42695 CNDD3_HUMAN
4508	SALSPSKSPAK200020000	sp Q12873 CHD3_HUMAN
4509	SFTLDESLSK200000000	sp P49792 RBP2_HUMAN
4510	AGMSSNQSISSPVLDVAVPR2002022000000000	sp Q7Z3J3 RGPD4_HUMAN
4511	TIASDSEEEAGK2020000000	sp Q86WR7 PRSR2_HUMAN
4512	SPPAPGLQPMR200000000000	sp Q9UQ35 SRRM2_HUMAN
4513	SNSPLPSIQLQPQSPSASK2000000	sp Q96ST2 IWS1_HUMAN
4514	EAASSSGTQPAPPAPASPWDK2000000	sp P15408 FOSL2_HUMAN
4515	GFEEHKDSDDDSSDDEQEKKPEAPK20000200000000000000	sp Q08174 PCDH1_HUMAN
4516	ISSSFSR20000	sp Q9H7N4 SFR19_HUMAN
4517	GSVSPSTPSPPKPTGIFQTSANSSFEVPK2. 002E+27	sp Q13435 SF3B2_HUMAN
4518	FPTYASTSEDEFSGSNR222000000000	sp P05787 K2C8_HUMAN
4519	VGVSSKPDSSPVLSPGNK20000200000	sp 015027 SC16A_HUMAN
4520	SNSAPLIHGLSDTSPVFQAEAPSAR20000000000000	sp Q5SW79 CE170_HUMAN
4521	KRPACPEPGESESSSDEGCTVVRPEK202220000000000000	sp Q8N4C8 MINK1_HUMAN
4522	RPDHSGGSPSPPTSEPAR2000020000000000	sp Q96E09 F122A_HUMAN
4523	QPSSPSHNTNLR2000000000	sp 015541 R113A_HUMAN
4524	SPAVATSTAAPPPSSPLPSK2E+21	sp Q8TAD8 SNIP1_HUMAN
4525	SLTNSHLEK20000000	sp Q9UIF9 BAZ2A_HUMAN
4526	DAEPQGPSAAESLEEDAAAGLSSTK2E+20	sp Q14157 UBP2L_HUMAN
4527	TPTAPAVNLAGAR200000000000	sp Q9H2H9 S38A1_HUMAN
		sp Q9H9J4 UBP42_HUMAN
		sp Q9UQ35 SRRM2_HUMAN

4528	HKSDNETNLQQVVGWNR200000000000	sp Q7Z401 MYCPP_HUMAN
4529	ASPPQGPLPGPPGALHR2000000000000000	sp Q96G74 OTUD5_HUMAN
4530	QDPVTYISETDEEDDFMCK2020000001000	sp P35251 RFC1_HUMAN
4531	TRSVEDDEEGHLICQSGDVLSAR2.02E+23	sp P49759 CLK1_HUMAN
4532	SIVECQPDASATPGVGTADSPAAPTDSR2000000000	sp Q7Z2Z1 TICRR_HUMAN
4533	SSSQPSSCCSDPSKPGNVEGATQSLAEQMR2E+29	sp Q13501 SQSTM_HUMAN
4534	SIDTQTPSVQER20000000000000	sp Q86VQ1 GLC11_HUMAN
4535	SKSDGEAKPEPSPSPR2000000000000000	sp P26358 DNMT1_HUMAN
4536	ETSAATLSPGASSR20000000	sp Q9Y6I9 TX264_HUMAN
4537	AKTQTPPVSPAPQPTTEER2000200000000000	sp Q14247 SRC8_HUMAN
4538	KPSPQPSSPR200002000	sp 075909 CCNK_HUMAN
4539	SGMSPEQSR2000000	sp Q9UQ35 SRRM2_HUMAN
4540	NSPNNISGISNPPGTPR2000	sp Q9BWW4 SSBP3_HUMAN
4541	LRLSPSPTSQR200000000	sp P02545 LMNA_HUMAN
4542	AWLDEDSNLSPSPLR2002020000	sp Q9C086 IN80B_HUMAN
4543	SHSPSASQSGSQLR200000000	sp Q8WWI1 LM07_HUMAN
4544	EVDDILGEGSDSDSEK200200000	sp Q9Y5B0 CTDP1_HUMAN
4545	NSSTETDQQPHSPDSSSVHSIR20000000000000	sp Q92613 JADE3_HUMAN
4546	EIQGNLHESDSESVPR202000000	sp Q86UP2 KTN1_HUMAN
4547	LDEDDSCSLTK20200000	sp Q9HAW4 CLSPN_HUMAN
4548	QPEYSPESPR2000	sp Q9H6F5 CCD86_HUMAN
4549	GKGSRPQAPK200000000	sp Q14978 NOLC1_HUMAN
4550	ATSSSSGSLSATGR20000000000000	sp Q03252 LMNB2_HUMAN
4551	TDSVIIADQTPPTR20000000020000	sp P17544 ATF7_HUMAN
4552	ETIELSPTGRPK20000000	sp Q9NRZ9 HELLS_HUMAN
4553	NHSGSRTPPVALNSSR2000200000000000	sp Q9UQ35 SRRM2_HUMAN
4554	ALSSGGSITSPPLSPALPK200000020002000000	sp Q6Y7W6 PERQ2_HUMAN
4555	TVIIIEQSWGSPK2000	sp P10809 CH60_HUMAN
4556	EAEDSDSDDNIKR20000000000	sp Q96ST2 IWS1_HUMAN
4557	RPASPSSPEHLPATPAESPAQR2.02E+19	sp Q9H7L9 SDS3_HUMAN
4558	SDSGSSSEPFDR20000200000000	sp P48634 PRC2A_HUMAN
4559	GPRTPSPPPPYPEDIALGK2020000000000000	sp Q5T200 ZC3HD_HUMAN
4560	KRSSSEDAESLAPR220000000000	sp Q4G0J3 LARP7_HUMAN
4561	LEKPETQSSPITVQSSK2000000000	sp Q96CB8 INT12_HUMAN
4562	KSSADTEFSDECTTAER2000000000000000	sp Q9H6S0 YTDC2_HUMAN
4563	ECELSPGVNR2000000	sp Q9BXP5 SRRT_HUMAN
4564	VGSPLTLTDAQTR2000000000000	sp Q99569 PKP4_HUMAN
4565	DPSSGQEVATPPVQLQVCEPK20000000000000	sp Q9UKV3 ACINU_HUMAN
4566	MNSTGGQTPR200000000	sp Q96ST2 IWS1_HUMAN
4567	GYSPYSVSGSGSTAGSR30000000002000000	sp Q15149 PLEC_HUMAN
4568	LASPMKVPPTSSK201000002000000	sp Q15648 MED1_HUMAN
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4570	SPISVPGGSALISNLGK200000000000000000	sp P51610 HCFC1_HUMAN
4571	RPAEATSSPTSPERPR2000000	sp Q9BU76 MMTA2_HUMAN
4572	RNSVERPAEPVAGAATPSLVEQQK2E+22	sp Q9BTC0 DIDO1_HUMAN
4573	SPLELEPEAK20000000000	sp Q92466 DDB2_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
4574	GGSYSQAASSDSAQGSVSLTACK2000000000000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
4575	ENSGSPLEFR2000000000	sp Q9UQ35 SRRM2_HUMAN
4576	QSVVYGGKSTIR22000	sp Q8WWY3 PRP31_HUMAN



4577	SGGVTEASSLLGGSPR2000	sp Q92729 PTPRU_HUMAN
4578	SAYQEYSDSDVPEELKR2000000000	sp Q96JG6 CC132_HUMAN
4579	EVEEDSEDEEMSEDEEDDSSGEEVVIPQK2.00001000000022E+24	sp P19338 NUCL_HUMAN
4580	EVYELLDSPGK20000	sp P22234 PUR6_HUMAN
4581	VGEDEDGSDEEVVHNEDIHFEPVIVSLPEVEVK2E+25	sp P49792 RBP2_HUMAN
4582	SSVGPSPKPVSPQR2000000000000	sp Q9Y657 SPIN1_HUMAN
4583	SSTPSPSPLNLSSSR22000000000000	sp P37275 ZEB1_HUMAN
4584	NSSSPVPASVPGQR2020020000000000	sp Q9H1B7 I2BPL_HUMAN
4585	GSSPGIQDTLEAEDGAFETDEAPEDR2E+24	sp Q9UNX4 WDR3_HUMAN
4586	VENMDISDEEEGGAPPPGSPQVLEEDEAELELQK1.002E+31	sp O43290 SNUT1_HUMAN
4587	HEHPNPPVSPGK20000	sp Q6VMQ6 MCAF1_HUMAN
4588	ESKEETSIDVAGKPNEVTK20000000000000	sp P53985 MOT1_HUMAN
4589	STPLASPSPPGR200020000	sp Q8WYP5 ELYS_HUMAN
4590	FTDKDQPSGSEGEDDDAEALKK2000000000000000	sp Q9NXG2 THUM1_HUMAN
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4593	LAAQLGAPTSPIPDSAIVNTR2000000000000000	sp Q9H8Y5 ANKZ1_HUMAN
4594	DHSPTPSVFNSDEER20002000200000	sp Q6UN15 FIP1_HUMAN
4595	DSSTSYTETKDPSSGQEVATPPVPQLQVCEPK2.2E+20	sp Q9UKV3 ACINU_HUMAN
4596	KVDYDYSQFQESDDAEDYGR2000000000000000	sp Q9H1E3 NUCKS_HUMAN
4597	IEEQELSENTSLPAEEANGSLSEEEANGPELGSGK2.00000000202E+25	sp Q9BUR4 WAP53_HUMAN
4598	TSPADHGGVSGSESGSAVDSVAGEHSVSGR2000000000000000	sp Q5T4S7 UBR4_HUMAN
4599	HLFSSTENLAAGSWK20000000000000	sp Q6WKZ4 RFIP1_HUMAN
4600	HSTPSNNSNSGPPSPNSPHR200020000000	sp Q9Y5S2 MRCKB_HUMAN
4601	GADNDGSGSESGYTPK200000000000	sp Q7Z417 NUFP2_HUMAN
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4603	SPGTSPLLSR20002000000	sp Q9BRS8 LARP6_HUMAN
4604	TPPAPSPFDLPELK2000000000000000	sp O95239 KIF4A_HUMAN
4605	ESLSTHTSPSQSPNSVHGSR202000000000	sp Q9H1K0 RBNS5_HUMAN
4606	DDSHAEDSEDEKEDHK20002000000000	sp Q9H1E3 NUCKS_HUMAN
4607	QLSALSSDGEQGGRPVPR200200000000000000	sp O15438 MRP3_HUMAN
4608	HQSDKNPGSSNLQK20000000000000	sp Q12986 NFX1_HUMAN
4609	TASRPDDIPDSPSSPK2000	sp Q5VZK9 LR16A_HUMAN
4610	EALLQASR200	sp P26038 MOES_HUMAN
4611	ESISQPADSACSSPAPSTGK20000020000000000000	sp Q9GZR1 SENP6_HUMAN
4612	SAADSISESVPVGPK200000000000	sp P45974 UBP5_HUMAN
4613	NTPSQHSHSIQHSPER200020000	sp Q9NYF8 BCLF1_HUMAN
4614	SRSTTAHSWQR202000000000	sp Q9UKJ3 GPTC8_HUMAN
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4616	RLSPSASPPR202020000	sp Q8IYB3 SRRM1_HUMAN
4617	ELSNSPLRENSFGSPLEFR2000000002000000	sp Q9UQ35 SRRM2_HUMAN
4618	GSSLSGTDGAQEVVK2000000000000000	sp Q9Y6D5 BIG2_HUMAN
4619	SQDQDSEVNELSR200000200	sp Q86VM9 ZCH18_HUMAN
4620	RLSSASTGKPPLSVEDDFEK20002000000000000000	sp O75152 ZC11A_HUMAN
4621	RGFSDSGGPPAK20000000000	sp Q9H307 PININ_HUMAN
4622	RSSLSSHSHQSQIYR2200000000000000	sp O15027 SC16A_HUMAN
4623	NHLLQFALESPAKSPASSSSK20002000000000	sp Q9NS91 RAD18_HUMAN
4624	AKSPQPPVEEEDHFDDTVVCLDTYNCDLHFK2000000000000000	sp Q00839 HNRPU_HUMAN
4625	SRTPPSAPSQR20200000000000	sp Q9UQ35 SRRM2_HUMAN
4626	SPAVATSTAAPPPSSPLPSK2000000000000000	sp Q14157 UBP2L_HUMAN
4627	GFEEEHKSDDDSSDDEQEKKPEAPK20002200000000000000	sp Q13435 SF3B2_HUMAN
4628	SSSQRSPPGNHTSNSSNASNATVVPQNSSAR2.2002E+32	sp Q9BTA9 WAC_HUMAN
4629	SAAGSEKEEPEDEEEEEEEEEEEYDEEEEEEDDRPPK2.0002E+37	sp O00267 SPT5H_HUMAN

4630	IIIEELSLVAASPTLSR2000	sp Q16513 PKN2_HUMAN
4631	SLSQGSTNSNMLDVQGAHK20000000000000000000	sp Q8TEU7 RPGF6_HUMAN
4632	LQEDPNYSPQR20000	sp P46379 BAG6_HUMAN
4633	DLQEADAGSER2000	sp Q6P158 DHX57_HUMAN
4634	SSLGSLQTPAVTTR2000002000000000	sp Q7Z2W4 ZCCHV_HUMAN
4635	HQEVQDQDPVFQGSDDSSGYQSDHKK200000	sp Q3B726 RPA43_HUMAN
4636	SRPTSFADELAAR22000000000	sp Q641Q2 FA21A_HUMAN
		sp Q9Y4E1 FA21C_HUMAN
		sp Q5SNT6 FA21B_HUMAN
4637	TEIKKEEDQPSTSATQSSPAPGQSK200000000	sp Q09472 EP300_HUMAN
4638	TASRPDDIPDSSPK200000000002000	sp Q5VZK9 LR16A_HUMAN
4639	QQFYHSVQDLGGSR200	sp P10398 ARAF_HUMAN
4640	SGGVTEASSLLGGSPR200002000	sp Q92729 PTPRU_HUMAN
4641	LSAITMALQYGESEEDAALAAAR20200000000000	sp Q86U86 PB1_HUMAN
4642	TPTPSTPGDTQPNTAPVPPAEDGIK2.00000000002E+24	sp Q14839 CHD4_HUMAN
4643	ALVHQLSNESR200000	sp Q9H4L5 OSBL3_HUMAN
4644	LSLTSDPEEGDPLALGPESPGEPQPPQLK2E+28	sp Q5T1M5 FKB15_HUMAN
4645	ALSSAVQASPTSPGGSPSSSGQR200020000000000	sp Q03164 KMT2A_HUMAN
4646	LVEPHSPSPSSK20200000	sp O15014 ZN609_HUMAN
4647	GSVPLSSPPATHFPDETEITNPVK2E+21	sp P11388 TOP2A_HUMAN
4648	SPTPPSSAGLGSNSAPPIDSR2.002E+20	sp Q8IWX8 CHERP_HUMAN
4649	CSASPPLR200000	sp P46013 KI67_HUMAN
4650	WAHDKFSGEEGEIEDDESGTENR200000000000020000	sp Q9Y2W1 TR150_HUMAN
4651	HSPIAPSSPSPQVLAQK20000020200000000	sp Q9NQS7 INCE_HUMAN
4652	EPLENSASESSDELPEK202002000000000	sp P78337 PITX1_HUMAN
4653	SLPAEEGSSGK200000000000	sp Q9UKJ3 GPTC8_HUMAN
4654	VETVSPSESPKDTIDK200000000	sp Q99590 SCAFB_HUMAN
4655	EGEDGDQPTTPKPLK20000000	sp P18858 DNLI1_HUMAN
4656	NYAGEEEEEEGSGSSEGFDPATDR2200000000000	sp P16989 YBOX3_HUMAN
4657	GIAPASPMLGNASPNK2010000000000	sp P27448 MARK3_HUMAN
4658	SESPPAELPSLR2020000000000	sp Q14669 TRIPC_HUMAN
4659	ATPSENLPSSAR2000000000000	sp Q8N684 CPSF7_HUMAN
4660	RKSLDSESDSK2020200000000	sp Q13185 CBX3_HUMAN
4661	KDDSHAEDSEDEKEDHK20200020000000000	sp Q9H1E3 NUCKS_HUMAN
4662	SATRPSPPERSSTGPEPPAPTLLAER2.00022E+21	sp Q9UQ35 SRRM2_HUMAN
4663	DKDQPPSPSPPPQSEALSSTR20000000000000000	sp Q7L1V2 MON1B_HUMAN
4664	EEGSLDTEADAVSGQLPDPTNPSAGK2E+21	sp Q8N1G4 LRC47_HUMAN
4665	DDISEIQLASDHSGR20002002002000	sp Q07157 ZO1_HUMAN
4666	RSPSPAPPPR20000000000	sp Q8IYB3 SRRM1_HUMAN
4667	SPLLAGGSPQPVVPAHK200000000000	sp Q8NFH5 NUP53_HUMAN
4668	TSSLDTITGPYLTGQWPR20000000000000000	sp Q86VQ1 GLCI1_HUMAN
4669	EVSPEVVR2000000	sp Q5T200 ZC3HD_HUMAN
4670	FHSPSTWSPNKDTPQEK20000020000000000	sp Q99590 SCAFB_HUMAN
4671	MSSPPSPQK200000000	sp Q9Y2H6 FND3A_HUMAN
4672	ASSPSLIER200000000	sp A1L170 CA226_HUMAN
4673	RNSSDSAIDNPKPNK2000000000000	sp P46100 ATRX_HUMAN
4674	GESAPTLSTSPSPSSPSTSPSPTLGR2E+25	sp Q9H330 TM245_HUMAN
4675	RLSSASTGKPLSVEDDFEK2002000000000000000	sp O75152 ZC11A_HUMAN
4676	HDYDSSSEEQAEDSYEASPGSETQR3E+24	sp P98175 RBM10_HUMAN
4677	IGELGAPEVWGLSPK2000	sp Q8N5F7 NKAP_HUMAN
4678	LTGSTSSLNK20000	sp Q9NV70 EXOC1_HUMAN
4679	GTLSDSEIETNSATSTIFGK200000000000	sp Q8WXG6 MADD_HUMAN
4680	SNSPLPSIQLQPQSPSASK200000000002000000	sp Q08174 PCDH1_HUMAN
4681	SPTPPSSAGLGSNSAPPIDSR2.02002E+22	sp Q8IWX8 CHERP_HUMAN

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4683	QLSALSSDGEQGGRPVPR200220000000000000	sp 015438 MRP3_HUMAN
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4685	SIDQDYER200000000	sp Q9UDY2 ZO2_HUMAN
4686	SPDKPGGSPSASR2000000	sp Q9Y3T9 NOC2L_HUMAN
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4688	GLSPAMSPALQR20002000000	sp Q96DF8 DGC14_HUMAN
4689	SASPEVSEGHENQHGESEAK2.02E+21	sp 060930 RNH1_HUMAN
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4691	LISPLASPADGVK200020000000	sp Q8WYP5 ELYS_HUMAN
4692	SSEGGVGVGPGGGDEPPTS20000	sp Q9Y4H2 IRS2_HUMAN
4693	YPSSISSSPQK20000	sp Q14157 UBP2L_HUMAN
4694	SPSPGNHTSNSSNASNATVVPQNSSAR220000000000000000	sp Q9BTA9 WAC_HUMAN
4695	AGMSSNQSISSPVLDAVPR120000002000000000	sp Q9UQ35 SRRM2_HUMAN
4696	SVQSAAVPAPTSQLLSSLEK2.002E+20	sp Q96T58 MINT_HUMAN
4697	TKPTQAAGPSSPQKPPTPEETK20000002000000	sp P18887 XRCC1_HUMAN
4698	HHFETLTESPCR20000	sp 043379 WDR62_HUMAN
4699	LCSSDTLVSEGEENQKPK200000000000000000	sp Q5T0W9 FA83B_HUMAN
4700	LGSTSGEESDLER22000200000	sp Q9BW71 HIRP3_HUMAN
4701	ASESSSEKDDYEIFVK200200000000000000	sp P18583 SON_HUMAN
4702	FDDMSSPGLLEPSCELSR2000000000000000	sp Q6IN85 P4R3A_HUMAN
4703	SLAFDSEHSADEK200200000	sp Q96L73 NSD1_HUMAN
4704	YPSSISSSPQK200000	sp Q14157 UBP2L_HUMAN
4705	REEQTDTSDGESVTHHIR2020000000000000	sp P36915 GNL1_HUMAN
4706	SESSDFEVVPK200000000000	sp Q9Y520 PRC2C_HUMAN
4707	CGGVEQASSSPR2000	sp Q5VZL5 ZMYM4_HUMAN
4708	RNSSSPVSPASVPGQR2200020000000000	sp Q9H1B7 I2BPL_HUMAN
4709	NTGIICTIGPASR200	sp P14618 KPYM_HUMAN
4710	SSPELVTHLK22000000000	sp Q07960 RHG01_HUMAN
4711	VLKEPVNLEGDDTSLIHGK2000000	sp Q9H2J7 S6A15_HUMAN
4712	SVSEHTSCPAESASDAAPLQR2E+20	sp Q8WUA7 TB22A_HUMAN
4713	VDSPSHGLVTSSLCIPSPAR20000000000020000	sp Q9UER7 DAXX_HUMAN
4714	EGHETPMDIDSDDSK200000	sp Q8TDB6 DTX3L_HUMAN
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4718	SSPGSVAASPSGGGR2000000000000000	sp Q9BXB5 OSB10_HUMAN
4719	RNSSEASSGDFLDLK2000000000	sp Q9UK76 HN1_HUMAN
4720	SPFLKPDER2000000000	sp Q9H7N4 SFR19_HUMAN
4721	RKTSDANETEDHLES LICK220000000000000000	sp Q09161 NCBP1_HUMAN
4722	TASRPDDIPDPSPPK2000000	sp Q5VZK9 LR16A_HUMAN
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4724	KDELSDWSLAGEDDRDSR2002000000000000	sp A8CG34 P121C_HUMAN
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4728	NSSDSAIDNPKPNK20020000000000	sp Q9UQ35 SRRM2_HUMAN
4729	DRSSPPPGYIPDELHQVAR200000000000000000	sp P46100 ATRX_HUMAN
4730	LQQHSEQPPLQPSVMTR2000000	sp Q9Y2U5 M3K2_HUMAN
4731	YPSSISSSPQK220000	sp Q5JTV8 TOIP1_HUMAN
4732	SKSDGEAKPEPSPSPR200000000002000	sp Q14157 UBP2L_HUMAN
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4734	SYTSGPGR2000000000	sp Q8WVC0 LEO1_HUMAN
		sp P05787 K2C8_HUMAN

4735	ASPGTPLSPGSLR2000000	sp Q96BD0 S04A1_HUMAN
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4737	RNSVTPLASPEPTK2000002000000	sp Q16875 F263_HUMAN
4738	HASSSPESPAPAPGSHR202002000000000000	sp Q9UQ35 SRRM2_HUMAN
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4740	TPASTPVSQTPQASPMIER2000000	sp Q9UHR4 BI2L1_HUMAN
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4745	KPPAPSPVQSQSPSTNWSPAVPVKK20202000000000000	sp Q8IYB3 SRRM1_HUMAN
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4747	TPQAPASANLVGPR2000000000000000	sp Q9UQ35 SRRM2_HUMAN
4748	ADAPDAGAQSDSELPSYHQNDVSLDR200000000000020000	sp Q92538 GBF1_HUMAN
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4750	GTGSGGQLQDLDCSSDDEGAAQNSTKPSATK2E+29	sp P08240 SRPR_HUMAN
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4752	AKNSPPQAPSTR2000000000	sp Q8N1F8 S11IP_HUMAN
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		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
4753	GGSCSQAASSNSAQGSDESLIACK2E+20	sp Q29963 1C06_HUMAN
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		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
4754	VQSGMGLILQGYESGSEEEGEIHEK2020000000000	sp Q13523 PRP4B_HUMAN
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4757	LTSVSSVDFDQR220000000	sp Q5T3F8 CSCL2_HUMAN
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4759	SFPLHSPVAGVAHR2000020000000000	sp Q96JN8 NEUL4_HUMAN
4760	KIPDPSDDVSEVDAR20002000000	sp P51532 SMCA4_HUMAN
4761	ASPAPGSGHPEGGAHLDMNSLDR2E+23	sp O94826 TOM70_HUMAN
4762	SVSPCSNVESR202002000000	sp Q9UQ35 SRRM2_HUMAN
4763	VGDTKEPEPERSPPNR200000	sp Q9NZ63 CI078_HUMAN
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4767	SLPTTVPESPNYR20000000200000	sp Q6PKG0 LARP1_HUMAN
4768	SGTSSPQSPVFR2200200000	sp Q8N556 AFAP1_HUMAN
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4770	LIDLESPTPESQK200000000	sp O43166 SI1L1_HUMAN
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4773	KLSSSEPYEEDEFNDDQS1KK2.02E+20	sp Q9H4L7 SMRCD_HUMAN
4774	SPPADAIK2000000000	sp P18754 RCC1_HUMAN
4775	SSSTDILEPFTVER2000000000000000	sp Q6GYQ0 RGPA1_HUMAN
4776	NSCNVGGGGGFK2000000000000	sp Q7L2J0 MEPCE_HUMAN
4777	SPQVKPASTMGMPGLGK2000000000000000000	sp Q13428 TCOF_HUMAN
4778	IEDSEPHIPLIDDTDAEDDAPTK2E+20	sp P20020 AT2B1_HUMAN

4779	SLHSAHSLASR200200000	sp 015027 SC16A_HUMAN
4780	KLSSERPSSDGEVVENGGITTCNGK2.000002E+23	sp Q8N556 AFAP1_HUMAN
4781	KSDGACDPSDDKENSQIAQDHQK220000000000000000	sp Q9UIG0 BAZ1B_HUMAN
4782	SQDQDSEVNELSR20000200000200	sp Q86VM9 ZCH18_HUMAN
4783	DEALQNISADDLPDSASQAHPQDSAFSYR2E+23	sp Q14C86 GAPD1_HUMAN
4784	SSPEQPRPSTSK200000020000	sp P29590 PML_HUMAN
4785	KESESEDSSDDEPLIKK2020022000000000	sp P35659 DEK_HUMAN
4786	SLDSESEDEEDDYQQKR2002000000000000	sp Q13442 HAP28_HUMAN
4787	HLLTDLPLPELPGDLSPPDSPEPK2000200000	sp Q9NYV4 CDK12_HUMAN
4788	AFLSPPTLLEGPLR200000000000	sp 015320 CTGE5_HUMAN
4789	AASSAAQGAFQGN200000000000	sp 015127 SCAM2_HUMAN
4790	VELSESEEDK20200000	sp Q8IYB3 SRRM1_HUMAN
4791	RSPSPYYSR202000000	sp Q13595 TRA2A_HUMAN
4792	LLELTSSYSPDVSDYK20000000000	sp P62995 TRA2B_HUMAN
4793	WTKPSSFSDSER200000	sp P38432 COIL_HUMAN
4794	SNSTTQVSQPR2000000000	sp Q8NC56 LEMD2_HUMAN
4795	SASASHQADIK2000000000	sp Q9UPN4 CP131_HUMAN
4796	GPLAGLPGSTPR22000	sp Q14739 LBR_HUMAN
4797	SGSALLQSQSTEDPKDEPAELKPDSEDLSSQSSASK2.2E+28	sp Q8WUM0 NU133_HUMAN
4798	SPSSKPSLAVAVDSAPEVAVK2E+21	sp Q5VTR2 BRE1A_HUMAN
4799	QISYNYSDLQSNVTEETPEGEEHHPVADTENK2E+31	sp 060508 PRP17_HUMAN
4800	YEPQSPGYEPR20000000	sp Q8NC51 PAIRB_HUMAN
4801	HKSDSPESDAER20200200000	sp Q9H5H4 ZN768_HUMAN
4802	ASPSPQSSQLQIHR2000000000000000	sp 075400 PR40A_HUMAN
4803	SNTPESIAETPPAR2000000000000000	sp Q9HC35 EMAL4_HUMAN
4804	GNDPLTSSPGR220000	sp Q9Y4H2 IRS2_HUMAN
4805	GSGGLFSPSTAHPDVGALGQR2.00002E+20	sp P49736 MCM2_HUMAN
4806	AGSPLCNSNLQDSEEDIEPSYSLFR2.0000000002E+23	sp Q9C0C2 TB182_HUMAN
4807	ENSPVQSIR20000000	sp Q9NRY4 RHG35_HUMAN
4808	HKLTVDGEGEGEKK20002000000	sp Q7Z2Z1 TICRR_HUMAN
4809	KPVTVSPTTPTSPTTEGEAS22000000000000	sp P46100 ATRX_HUMAN
4810	TASPPPPPKR20200000000	sp Q9Y6G9 DC1L1_HUMAN
4811	ENSGPVENGVDQEGEEQAREPELPCGLAPAVSR2E+24	sp Q8IYB3 SRRM1_HUMAN
4812	YGGSHYSSGYSNSR20000	sp Q8TEQ6 GEMI5_HUMAN
4813	AGGSAALSPSK200	sp Q9BRL6 SRSF8_HUMAN
4814	KSSSEAKPTSLGLAGGHK202000000000000000	sp Q92522 H1X_HUMAN
4815	KGGSYSQAASSDSAQGSVDVSLTACK2000000000	sp Q96RK0 CIC_HUMAN
4816	KLSSSSEPYEDEFNDDQSIKK2.002E+20	sp P01891 1A68_HUMAN
4817	TASISSPSEGTPVGSYGCTPQSLPK2E+27	sp P01892 1A02_HUMAN
4818	TSSAFVGTPEASPEPK2000000000200000	sp P05534 1A24_HUMAN
4819	KYSISSDNDTDDSHATSTSASR2.02E+21	sp Q09160 1A80_HUMAN
4820	SSSLQGMASLPPR2000000000000000	sp P10316 1A69_HUMAN
		sp P30447 1A23_HUMAN
		sp Q9H4L7 SMRCD_HUMAN
		sp Q6PKG0 LARP1_HUMAN
		sp Q99460 PSMD1_HUMAN
		sp Q9NQC1 JADE2_HUMAN
		sp Q9C0J8 WDR33_HUMAN

		sp P30510 1C14_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
4821	GGSCSQAASSNSAQGSDESLIACK20002000000000	sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
4822	LSSSDRYSDASDDSFSEPR20020000000000	sp Q9NRF8 PYRG2_HUMAN
4823	VADNSFDAK200000	sp Q14978 NOLC1_HUMAN
4824	TQDPSSPGTTPPQAR20000000000000	sp Q13112 CAF1B_HUMAN
4825	GAAAAATGQPGTAPAGTPGAPPLAGMAIVK2E+24	sp Q13263 TIF1B_HUMAN
4826	SKTFNPGAGLPTDK2000000000000000	sp P09661 RU2A_HUMAN
4827	SSTPLHSPSPIR200002000000	sp O95817 BAG3_HUMAN
4828	NDFEGLSPVVIAPK2020000000	sp Q9H1A4 APC1_HUMAN
4829	AAPPPPALTPDSQTVDSCK20000000000000	sp Q9H7N4 SFR19_HUMAN
4830	QLPALDGLMGPESPPAQEEEAPVSPHK2E+21	sp Q8N9T8 KRI1_HUMAN
4831	SFSMQDLR2000000	sp Q9H6H4 REEP4_HUMAN
4832	ATSSSNPSSPAPDWYK2000000200000000	sp P46100 ATRX_HUMAN
4833	SKSETGDSSIFR200000000000	sp O43166 SI1L1_HUMAN
4834	GSVILDSGHLSTASSSDDLKGEESFR2200000000000000	sp Q92614 MY18A_HUMAN
4835	CSGPGLSPGMVR2001000	sp P21333 FLNA_HUMAN
4836	SGDETPGSEVPGDKAAEEQGDDQDSEK2E+27	sp Q1KMD3 HNRL2_HUMAN
4837	GPPDFSSDEER2200000	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
		sp P30510 1C14_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
4838	GGSCSQAASSNSAQGSDESLIACKA200200000000	sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
4839	VISHDRDSPPPPPPPPPPPPPQPPSLK2E+19	sp Q69YN4 VIR_HUMAN
4840	HSASPVVFTSAR20200000000000	sp O43166 SI1L1_HUMAN
4841	ESKEETSIDVAGKPNEVTK2E+19	sp P53985 MOT1_HUMAN
4842	SGSPSDNSGAEEMEVSLAKPK2.00002E+19	sp P31749 AKT1_HUMAN
4843	MAESPCSPSGQQPPSPSPDEL PANVK1.00000002000002E+27	sp Q14160 SCRIB_HUMAN
4844	ERSDSGGSSEPFDR200002000000	sp P48634 PRC2A_HUMAN
4845	EVEEDESEEMSEDEEDDSSGEEVVIPQK2.00001200000002E+24	sp P19338 NUCL_HUMAN
4846	MSPSGGAPQTPTLR2000000000000000	sp Q9BY89 K1671_HUMAN
4847	SINEENGEVSEDQSQNK200000000200000000	sp Q13523 PRP4B_HUMAN
4848	DTLSNPQSPQSPYNSPKPQHK2.0000000003E+19	sp Q96PU5 NED4L_HUMAN
4849	SPSGPVKSPPLSPVGTTPVK2000000002000000000	sp Q9BVC5 ASHWN_HUMAN
4850	YSPTSPTYSPTSPVYTPTS PK3.00000002000003E+21	sp P24928 RPB1_HUMAN
4851	TPSPPPR20200000	sp Q8IYB3 SRRM1_HUMAN
4852	QFIAAQNGLPASGHGTPASSPSSSLPSPMSPTPR2E+24	sp Q8WX93 PALLD_HUMAN
4853	SVSPCSNVESR2000000000	sp Q9UQ35 SRRM2_HUMAN

4854	KKSSSEAKPTSLGLAGGHK220000000000000000	sp Q96RK0 CIC_HUMAN
4855	SNSSEAVLQGEELSAQAK200000000000000000	sp Q9BXF6 RFIP5_HUMAN
4856	SVCGHLENTSVGNPNPSSAENSFR220002000000000000	sp Q96HH9 GRAM3_HUMAN
4857	HTPNTSDNEGSDTEVCGPNSPSK2. 2E+19	sp Q8NEN9 PDZD8_HUMAN
4858	SPSDLHISPLAK2000000200000	sp 095785 WIZ_HUMAN
4859	TPTPSTPGDTQPNTAPVPPAEDGIK200020000000000000	sp Q14839 CHD4_HUMAN
4860	GPASTPTYSPAPTQPAPR200000000000000000	sp Q6P1N0 C2D1A_HUMAN
4861	RHSTSSLSQR2000020000	sp Q03164 KMT2A_HUMAN
4862	TPASTPVSQTPQASPMIER2000002000000	sp Q9UHR4 BI2L1_HUMAN
4863	DGQSSPQPSPR20000000	sp Q5VZK9 LR16A_HUMAN
4864	KPISDNSFSSDEEQSTGPIK20000000000000	sp 060293 ZC3H1_HUMAN
4865	RVSVCAETYNPDEEEEDTDPR2E+19	sp P13861 KAP2_HUMAN
4866	KDEETESEYDSEHENSEPVTNIR2. 000030200002E+20	sp Q15154 PCM1_HUMAN
4867	KELSPAGSISK200000000	sp 095453 PARN_HUMAN
4868	QEKPAEKPAETPVATSPTATDSTSGDSSR2. 0002E+19	sp P54727 RD23B_HUMAN
4869	EDAATPKAPPAPPPQNLQPESDAPQPGSSPR2000	sp Q9Y618 NCOR2_HUMAN
4870	YGMGTSVER20000	sp P08559 ODPA_HUMAN
4871	VVPSFLPVDQGGSLVGR200000	sp Q14126 DSG2_HUMAN
4872	GPSQATSPIR200020000	sp Q7RTP6 MICA3_HUMAN
4873	SESSDFEVVPK2200000000	sp Q9Y520 PRC2C_HUMAN
4874	GISPIVFDR20000000	sp Q96MU7 YTDC1_HUMAN
4875	GSISSTSEVHSPNVGLR200200000000000000	sp Q8WWQ0 PHIP_HUMAN
4876	SYLEGSSDNQLKDESETPVDDR2. 0000220000002E+22	sp Q9UHI6 DDX20_HUMAN
4877	KEDSDEEEDDDSEDEEDEDDEDEDEDEIEPAAMK2. 00000002E+32	sp P19338 NUCL_HUMAN
4878	EFTGSPSSATK200000000	sp Q92609 TBCD5_HUMAN
4879	DSAPSTPTSPTFLTPGLR200000000000	sp Q9NZJ4 SACS_HUMAN
4880	TPTSSPASSPLVAK20002000000	sp Q14684 RRP1B_HUMAN
4881	RPTPNDDLDEGVGLVHSNIATEHIPSPAK20000	sp Q01581 HMCS1_HUMAN
4882	TPASTPVSQTPQASPMIER2000000002000000	sp Q9UHR4 BI2L1_HUMAN
4883	AGGSAALSPSKK200000	sp Q92522 H1X_HUMAN
4884	TEMDKSPFNPSQDSR200000000000000000	sp P08651 NFIC_HUMAN
4885	MSGFIYQ GK200000000	sp Q14155 ARHG7_HUMAN
4886	DTHEDHDTSTENTDESNHDPQFEPVSLPEQEIK2. 2E+27	sp Q15052 ARHG6_HUMAN
4887	SPPPGMGLNQR200000000000	sp P43487 RANG_HUMAN
4888	TLNAETPKSSPLPAK20020000000	sp P23246 SFPQ_HUMAN
4889	GGQGDPAVPAQQPADPSTPER20000	sp Q7L2J0 MEPCE_HUMAN
4890	TLLPWDSDESPEASPGPPGPR2002000200000000	sp 015231 ZN185_HUMAN
4891	ENESSDSEEEVSRPR220200000000	sp Q9UMN6 KMT2B_HUMAN
4892	KESEEDSSDDEPLIKK2020002000000000	sp Q6KC79 NIPBL_HUMAN
4893	TDCEPVGLEPAVEQSSAASELAEASSEELAEAPTEAPSPPEAR20000000200000000	sp P35659 DEK_HUMAN
4894	LNHSPPQSSSR200000000	sp Q15424 SAFB1_HUMAN
4895	LHQASASSTSSLSTR2000000	sp Q14151 SAFB2_HUMAN
4896	GSSGSPAHAESYSSGGGQK2. 2E+20	sp Q5T8P6 RBM26_HUMAN
4897	SGSSQELDVKPSASPQER202200000000000000	sp 015085 ARHGB_HUMAN
4898	VAESAAAAPQGGPELPPSPASPPEQPPEER2002000000000000	sp P15408 FOSL2_HUMAN
4899	RVSHSPPPK20200000	sp Q9UQ35 SRRM2_HUMAN
4900	SQEPIDDDQKVSDDDKEK20000000	sp Q96N64 PWP2A_HUMAN
4901	DDGVFVQEVTVQNSPAAR200000	sp Q8IYB3 SRRM1_HUMAN
4902	SPSPPPTR202000000	sp Q86UE4 LYRIC_HUMAN
4903	VGVSCKPDSSPVLSPGNK2000200000	sp Q09666 AHNK_HUMAN
4904	FHALSSPQSPFPSTPTR20020000000000	sp Q8IYB3 SRRM1_HUMAN
4905	AVPSPTTGEEGSHSR20000000000000	sp Q8N4C8 MINK1_HUMAN
		sp 043166 SI1L1_HUMAN
		sp Q5SYE7 NHSL1_HUMAN

4906	SGSMEEDVDTSPGGDYITSPSSPTSSSR33000020000000	sp P08651 NFIC_HUMAN
4907	AEEKSPIINVK200200000	sp Q9UK58 CCNL1_HUMAN
4908	NHSGSRTPPVALNSSR2020200000000000	sp Q9UQ35 SRRM2_HUMAN
4909	TPPRDLPTIPGVTSPSSDEPPMEASQSHLR2.00002E+23	sp Q13541 4EBP1_HUMAN
4910	SHSESASPSALSSSPNLSPTGWSQPK2000020000000000	sp P04049 RAF1_HUMAN
4911	SPSPFTHTHLAQGYR2020000000000000	sp P38398 BRCA1_HUMAN
4912	LSAVSLR2002000	sp Q15036 SNX17_HUMAN
4913	HGGPGPGPEPELSPITEGSEAR200000000000	sp Q8WUF5 IASPP_HUMAN
4914	GTPSQSPVGR200020000000	sp P52594 AGFG1_HUMAN
4915	EKDKEPFTFSSPASGR2002000	sp Q6P1X5 TAF2_HUMAN
4916	SRSPIIHSPK200002000	sp O94913 PCF11_HUMAN
4917	NQGGGLSSSGAGEGQPK200000000000	sp P07814 SYEP_HUMAN
4918	ENSPAAFPDR2000000000	sp Q9Y5B0 CTDP1_HUMAN
4919	TLEAEFNPSPTPEPEGPR200000000000	sp A0MZ66 SHOT1_HUMAN
4920	KPPAPSPVQSQSPSTNWSPAVPVK2.000002002002E+19	sp Q8IYB3 SRRM1_HUMAN
4921	GHPAQRPEPSPPEPEQAPEGSQAGAEGPSSPEASR20000000	sp Q6ZW31 SYDE1_HUMAN
4922	SLSSGESGPGSPTHSHLSR20020000000000000000	sp Q6P0Q8 MAST2_HUMAN
4923	DMESPTKLDVTLAK1020000000000000	sp P27816 MAP4_HUMAN
4924	SPSPPTPEAK20200000000000	sp Q8TAQ2 SMRC2_HUMAN
4925	ATLGPTPTPPQPPDPSPPPQPMQH2E+21	sp O60927 PP1RB_HUMAN
4926	QGSITSPQANEQSVTPQR2200000000000000	sp Q9UQ35 SRRM2_HUMAN
4927	IESDTEETQDTSVDHNETGNTGESSVEENEK2E+29	sp Q6PL18 ATAD2_HUMAN
4928	KKSPNELVDDLK20000000000000	sp Q9UNZ2 NSF1C_HUMAN
4929	REEQTDSDGESVTHHIR2022000000000000	sp P36915 GNL1_HUMAN
4930	SCFESSPDELKSR2000000200	sp Q9UQ35 SRRM2_HUMAN
4931	GSVPLSSPPATHFPDETEITNPVPK2E+19	sp P11388 TOP2A_HUMAN
4932	TDALTSSPGR220000	sp P49736 MCM2_HUMAN
4933	VKASPIINDGEDEFVPSDGLDKDEYTFSPGK2.000000000002E+28	sp Q02880 TOP2B_HUMAN
4934	ALSSDSILSPAPDAR20200000000000	sp Q8IVT2 MISP_HUMAN
4935	MESEGGADDSAEGLDLLDDDNEDRGDDQLELIKDDEK1.02E+38	sp P07910 HNRPC_HUMAN
4936	GPTTGEGALDLSDVHSPKSPEGK2.000000002E+22	sp O95684 FR10P_HUMAN
4937	ISDSESDPPR20200000000000	sp Q96ST2 IWS1_HUMAN
4938	AGSSAAGASGWSAGSLNSVPTNSAQGHNSPDSPTSAAK2000000000	sp Q86XP3 DDX42_HUMAN
4939	AYSQEEITQGFEEETGDTLYAPYSTHFQLQNQPPQK2E+33	sp Q8NFJ5 RAI3_HUMAN
4940	GPPDFSSDEER2200000	sp P42166 LAP2A_HUMAN
4941	RTPTPPPR20200000	sp P42167 LAP2B_HUMAN
4942	GSPNTASAEATLPR2000000000000000	sp Q8IYB3 SRRM1_HUMAN
4943	DMDEPSPVNVVEVTLPK2000000000000000	sp Q6PJG6 BRAT1_HUMAN
4944	APSPTGPAALISGASPVHCAADGTVELK2.0000000002E+25	sp Q8TAQ2 SMRC2_HUMAN
4945	SPSPEPIYNSEGKR2020000000000000	sp Q76L83 ASXL2_HUMAN
4946	SVASNQSEMEFSSLQDMPK200201000000000000	sp Q15637 SF01_HUMAN
4947	SQEDEISSPVNK20000000000000	sp Q8N163 CCAR2_HUMAN
4948	RLSPSASPPR202000000	sp Q5UIP0 RIF1_HUMAN
4949	SASQSSLDKLDQELK20000000000000	sp Q8IYB3 SRRM1_HUMAN
4950	SAPASPTHPLMSR2000200000002000	sp O60271 JIP4_HUMAN
4951	AAKPGPAEAPSPTASPSGDASPPATAPYDPR200000200000000000	sp P85037 FOXK1_HUMAN
4952	SPSKLPEVTDEYK2000000000000000	sp Q9UPT8 ZC3H4_HUMAN
4953	SDSSGGYNLSDIIQSPSSTGLLK300000002000000000	sp P05455 LA_HUMAN
4954	ASHASSDWTTPRPR220000000000	sp Q9P2D0 IBTK_HUMAN
4955	RGPNYTSGYGTNSELSNPSETESER202000	sp Q96EB6 SIR1_HUMAN
4956	SPPEALVQGR200000000000	sp P51114 FXR1_HUMAN
4957	SSSISEEKGSDDEKPR2020000000000000	sp Q9UHB6 LIMA1_HUMAN
4958	LLPSAPQTLPGPLASPAR20000	sp Q9UKV3 ACINU_HUMAN
		sp O15027 SC16A_HUMAN



4959	HAASYSSDSENQGSYSGVIPPPGR300000000000	sp Q96D71 REPS1_HUMAN
4960	AGMSSNQSISSPVLDAVPR102002022000000000	sp Q9UQ35 SRRM2_HUMAN
4961	DEEQSEADAGSGPPTPGPTTLGPK2.000000002E+20	sp Q6ZRS2 SRCAP_HUMAN
4962	HGGSPQPLATTPLSQEPVNPPEASPTR2E+25	sp Q9UQ35 SRRM2_HUMAN
4963	ASPGTPLSPGSLR2000002000	sp Q96BD0 SO4A1_HUMAN
4964	AQTLPTSVVTTITSESSPGKR200000	sp Q01082 SPTB2_HUMAN
4965	AEDEILNRSR2000	sp P27824 CALX_HUMAN
4966	RGTSPRPPEGGLGYSQLGDDDLK2.2E+21	sp Q9UQ88 CD11A_HUMAN
4967	KPSPSEPEPWKPFPAVSPEPR2.02E+20	sp P21127 CD11B_HUMAN
4968	GDRSPEPGQWTR20000000000	sp Q96JM3 CHAP1_HUMAN
4969	GQEVETSVTYR2000000	sp Q09666 AHNK_HUMAN
4970	GGNFGFGDSR200	sp 043169 CYB5B_HUMAN
4971	SPSAEFSPAAPPGISSIHSPSLR2E+21	sp P22626 ROA2_HUMAN
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4973	SLNFQEDDTSR2000	sp P46013 KI67_HUMAN
4974	SPGSTPTTSSQAPQK2202000000000000	sp Q9H814 PHAX_HUMAN
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4976	DSSSTDSANEKPALIPR20020200000000000	sp Q92609 TBCD5_HUMAN
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4978	ILCKSPQSDPADTPTNTK200000000000000	sp Q12873 CHD3_HUMAN
4979	SCTPSPDQISHR2020200000000	sp P46013 KI67_HUMAN
4980	HQSDGNEIAHTR20000000000	sp Q7Z2W4 ZCCHV_HUMAN
4981	NSSDSAIDNPKPNKLPK2200000000000000	sp 043166 SIL1_HUMAN
4982	KRSPTESVNTVPVK20000000000	sp P46100 ATRX_HUMAN
4983	QSSGSATNVASTPDNR20000000000	sp Q8WXG6 MADD_HUMAN
4984	TLENSPISCDTSDTEGPVNVNSAAVLK2.0022002E+24	sp Q7Z460 CLAP1_HUMAN
4985	AKSPTPSPPR20202000000	sp Q8N556 AFAP1_HUMAN
4986	SRDATPPVSPINMEDQER200020002000000000	sp Q8IYB3 SRRM1_HUMAN
4987	TNLTQSSTTNLPGSPGSPGSPGSPGSPGSPK200200200000000	sp P17275 JUNB_HUMAN
4988	SQSIEQESQEK20000000000	sp Q6IN85 P4R3A_HUMAN
4989	LALAGEGEACPLLGSPGAGR2000000	sp Q5VTL8 PR38B_HUMAN
4990	EHYPVSSPSPSPAQPGGVSRR2000000000000000	sp Q96F46 I17RA_HUMAN
4991	NYAGEEEEEEGSGSSEGFDPATDR20020000000000	sp 075179 ANR17_HUMAN
4992	SLSPNHNTLQTLK2020000000000	sp P16989 YBOX3_HUMAN
4993	GQKPALDCSQVSPRPATSPENNASLSDTSPMDSSPSGIPK2.000002E+23	sp 075592 MYCB2_HUMAN
4994	YQPLASTASDNDFVTPEPR20000000000000	sp P18858 DNLI1_HUMAN
4995	SVEDVRPHHTDANNQSACFEAPDQK2000000000000000	sp Q15021 CND1_HUMAN
4996	SSDYQFPSSPFTDTLK2000000200000000	sp Q9Y520 PRC2C_HUMAN
4997	DLSTSPKPSPIPSPVLGR200020002000000	sp Q9UK61 F208A_HUMAN
4998	GSAPHSEDLPEQEEEILGSDDEQEDPNDYCK2.0002E+32	sp Q8NDI1 EHP1_HUMAN
4999	KLSSSEPYEDEFNDDQS IKK2.2E+20	sp Q96SB4 SRPK1_HUMAN
5000	SSSGLGSSGSLSQK20000000	sp Q9H4L7 SMRCD_HUMAN
5001	RPSGPDGAAR200000000	sp Q15648 MED1_HUMAN
5002	LPQSSSESSPPSPQPTK2002000000	sp Q8NDX5 PHC3_HUMAN
5003	SSGHSSSELSPDAVEK200020000000	sp Q9UQ35 SRRM2_HUMAN
5004	QLSALHR200000	sp Q9UQ35 SRRM2_HUMAN
5005	SPVGSQAPAAAPAAHVAGNPGDAAPAATGTAASLATAAGSEDAEKK2E+52	sp P61313 RL15_HUMAN
5006	GEPNVSYICSR300000	sp P16989 YBOX3_HUMAN
5007	ASESSSEKDDYEIFVK20220000000000000	sp P49841 GSK3B_HUMAN
5008	STPLASPSPPGR20000002000000	sp P49840 GSK3A_HUMAN
		sp P18583 SON_HUMAN
		sp Q8WYP5 ELYS_HUMAN

5009	KSLSPGVSER202000000	sp Q9UKV3 ACINU_HUMAN
5010	GSDALSETSSVSHIEDLEK2002000000000000	sp Q6P996 PDXD1_HUMAN
5011	LRLSPSPTSQR200020000	sp P02545 LMNA_HUMAN
5012	SGKDSEAWVPDSEER2000200000000000	sp Q96F63 CCD97_HUMAN
5013	DVTNFTVGGFAPMSR2000	sp 000267 SPT5H_HUMAN
5014	SAEPRPELPGQETGTNSR200	sp Q9Y4B5 MTCL1_HUMAN
5015	KGSFSALVGR2000000000	sp Q13619 CUL4A_HUMAN
5016	GLNPDGTPALSTLGGFSPASKPSSPR2E+20	sp Q9UK58 CCNL1_HUMAN
5017	SRSGEGEVSGLMR20000000000000	sp Q13263 TIF1B_HUMAN
5018	SNVSSPATPTASSSSSTPTR200000000000000000	sp Q12888 TP53B_HUMAN
5019	ESLKEEDESDDNM2000000000000000	sp P25788 PSA3_HUMAN
5020	SNSQSDSHDEEVSPPPNPVK200000200000000000	sp P31321 KAP1_HUMAN
5021	LAGGQTSQPTPLTSPQR200020000	sp 095155 UBE4B_HUMAN
5022	THSFENVSCHLPDSR2000000000000000	sp Q5VZ89 DEN4C_HUMAN
5023	KGSISSTQDTPVAVEEDCSLASSK20020000000000000000	sp Q9BZ71 PITM3_HUMAN
5024	ASSSPEPPAVAELSDSGQEPEDGGPAPGEASQGSSEGAQGLASASR2E+34	sp Q9C0C9 UBE20_HUMAN
5025	VMTIPYQMPASSPVICAGGQDR100020000000000000	sp Q15365 PCBP1_HUMAN
5026	KPSQTLQPSEDLADGK2000000000000000	sp Q13029 PRDM2_HUMAN
5027	RTSDPSAAVNHVSSTTSLGENYEDDLVNSDEVMK2E+30	sp Q8NHH9 ATLA2_HUMAN
5028	HASSSPESPAPAPGSHR222002000000002000	sp Q9UQ35 SRRM2_HUMAN
5029	GLSLSPQSPPER200200000	sp Q7Z2Z1 TICRR_HUMAN
5030	NMTVEQLLTGSPTSPTVEPEKPTR20000000000000	sp Q9C0C9 UBE20_HUMAN
5031	I IYDSSESEETLQVK3000200000000000	sp P35251 RFC1_HUMAN
5032	MPGMSPANPSLHSPVPDASHSPR200000002000	sp 060244 MED14_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
5033	GGSYSQAASSDSAQGSVDVSLTACK2000000000000000	sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
		sp P01892 1A02_HUMAN
5034	ISSKSPGHMVILDQTK2002000000000000	sp Q9NWH9 SLTM_HUMAN
5035	ISAESPTGLPSHK2000000000	sp Q96C24 SYTL4_HUMAN
5036	SPSGHPHVR2000000000	sp Q8WX93 PALLD_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
5037	GGSYSQAASSDSAQGSVDVSLTACK2000000000	sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
5038	SLDSESEDEEDDYQQR20000020000000000000	sp Q13442 HAP28_HUMAN
5039	RVSTDLPEGQDVYTAACNSV IHR2E+20	sp Q15751 HERC1_HUMAN
5040	SYLEGSSDNQLK2000000	sp Q9UHI6 DDX20_HUMAN
5041	AGLESGAEPGDGSDTTK200	sp 060832 DKC1_HUMAN
5042	ETSVSKEDTDHEEK20000002000000	sp 075475 PSIP1_HUMAN
5043	SPSDSSTASTPVAEQIER202000000000000000	sp Q16643 DREB_HUMAN
5044	SVCPGGSSPGSSSGGR200000000000	sp Q53LP3 SWAHC_HUMAN
5045	YFEADPPGQVAASPDPTT2000000	sp 043598 DNPH1_HUMAN
5046	LPNNSRPSTPTINVLESK22000000000000	sp Q15910 EZH2_HUMAN
5047	KRSSSEDAESLAPR22200000000000	sp Q4G0J3 LARP7_HUMAN
5048	LSNGSFSAPSLTNSR2002000000000000	sp 094806 KPCD3_HUMAN
5049	GSSDGRGSDSEDLPHR200002020000000000	sp Q9UPQ0 LIMC1_HUMAN
5050	WDQTADQTPGATPK2000	sp 075533 SF3B1_HUMAN
5051	HAPTYTIPLSPVLSPTLP AEAPTAQVPPSLPR2. 02E+19	sp Q9NPI6 DCP1A_HUMAN

5052	TENGTCPSPPLSPAAALGSGSAAAVPK2.000002E+22	sp Q12948 FOXC1_HUMAN
5053	DSAYISSPLGSHQVDFDR2000000000000	sp Q9HCD6 TANC2_HUMAN
5054	YRPENTPEPVSTSVSHYGAEPTTVSPCPSSSAK2000000000	sp P07947 YES_HUMAN
5055	ALVEFESNPEETREPGSPSVQR20000000	sp Q9H6F5 CCD86_HUMAN
5056	SSGHSSSELSPDAVEK22000000020000000	sp Q9UQ35 SRRM2_HUMAN
5057	VSSPLPSPSAMTDAANSQAAA2.000002E+20	sp Q8WXI9 P66B_HUMAN
5058	TFGHNTMDAVPR2000020000000	sp P55011 S12A2_HUMAN
5059	NRPTSISWDGLDSGK200000000000	sp P30086 PEBP1_HUMAN
5060	VEQATKPSFESGR2000000	sp P38159 RBMX_HUMAN
5061	SSLSSHSHQSQIYR220000000000000	sp O15027 SC16A_HUMAN
5062	LLSSNEDDANILSSPTDR20000000000002000	sp O75448 MED24_HUMAN
5063	DGDSYDPYDFSDTEEMPVHTPK300200000100000000	sp P25205 MCM3_HUMAN
5064	NAEEEESEAEEGD2000000000	sp Q7L1Q6 BZW1_HUMAN
5065	LQQQLQGGSR200	sp Q9UP83 COG5_HUMAN
5066	LLSPQQPALQR2000000000	sp Q14814 MEF2D_HUMAN
5067	QQPPEPEWIGGESTSPSDK200000	sp Q8NE71 ABCF1_HUMAN
5068	ELSNSPLR20000	sp Q9UQ35 SRRM2_HUMAN
5069	RPDHSGGSPSPPTSEPAR200002000000	sp Q8TAD8 SNIP1_HUMAN
5070	AASSSPGSPVASSPSR2000000000000000	sp Q92538 GBF1_HUMAN
5071	MNEEISSDSESESLAPR2200000000000	sp O43818 U3IP2_HUMAN
5072	DHSPTPSVFNSDEER2020000000000	sp Q6UN15 FIP1_HUMAN
5073	LLESSLSSEGEPEYK20000000000	sp P08648 ITA5_HUMAN
5074	LGSSGLGSASSIQAAVR20200000000	sp Q7Z6Z7 HUWE1_HUMAN
5075	MSPNETLFLESTNK20000	sp Q9HB90 RRAGC_HUMAN
5076	VHEPPREDAAPTKPAPPAPPPQNLQPESDAPQQPGSSPR2000	sp Q9NQL2 RRAGD_HUMAN
5077	GGWDHGDTSIQSSR2002000	sp Q9Y618 NCOR2_HUMAN
5078	SHSESASPSALSSPNLSPTGWSQPK22200002000000000	sp Q9P206 K1522_HUMAN
5079	SRSSSVGSSSYPIPAVSR220020000000000000	sp P04049 RAF1_HUMAN
5080	SKTPSSPCASK220000000	sp Q15149 PLEC_HUMAN
5081	KLSVPTSDEEDEVPAKPR200200000000000000	sp Q9NQS7 INCE_HUMAN
5082	YSVLNDDYFADVPLR20000	sp Q8NE71 ABCF1_HUMAN
5083	SGKNSQEDSESDKDKV200020020000000	sp Q1ED39 KNOP1_HUMAN
5084	SQSSDLLER2000000000	sp Q9H1E3 NUCKS_HUMAN
5085	EVHDELEDLSPPPPPLSPPTTSPHK20000020000020000	sp Q8IVT2 MISP_HUMAN
5086	SPYKFPSSPLR200000020000	sp Q9NQX3 GEPH_HUMAN
5087	SPGSSWQEYGPSDTGSR200000000000000000	sp P06400 RB_HUMAN
5088	QRGSETGSETHESDLAPSDK202020000000000000	sp Q5JSZ5 PRC2B_HUMAN
5089	TIGGGDDSFNTFFSETGAGK20000000000000	sp P48634 PRC2A_HUMAN
5090	DDSPDEVSTFTVQQPPPSR200000000000	sp Q71U36 TBA1A_HUMAN
5091	HTDDEMTGYVATR20300000	sp P68363 TBA1B_HUMAN
5092	MLPHAPGVQQAIPEDAVHEDSGDEDGEDPDKR200000000000	sp Q6PEY2 TBA3E_HUMAN
5093	ISLEQPPNGSDTPNPEK200000000	sp Q9BQE3 TBA1C_HUMAN
5094	GLAEVQQDGEAEEGATSDGEK200000	sp Q13748 TBA3C_HUMAN
5095	QLMEQDASSPSAQVIGLK20000000000	sp Q8N328 PGBD3_HUMAN
5096	CSDVSELSSPPGYPYHQEPYVCKPEER200000000000000000	sp Q16539 MK14_HUMAN
5097	GLSLVDKENTPPALSGTR2000000000	sp Q92769 HDAC2_HUMAN
5098	RDSGDVDGFEAEGK2000000000000	sp Q9UHI6 DDX20_HUMAN
5099	EAASSPAGEPLR200000000	sp Q02952 AKA12_HUMAN
5100	SESPPAELPSLR2000000000000	sp Q9H4L5 OSBL3_HUMAN
		sp Q9Y478 AAKB1_HUMAN
		sp P31350 RIR2_HUMAN
		sp Q96KR1 ZFR_HUMAN
		sp Q96EB6 SIR1_HUMAN
		sp Q14669 TRIPC_HUMAN

5101	GGHSSVSTESESSSFHSS20	sp P61073 CXCR4_HUMAN
5102	TIAHSPTSFTESSSK200000000000	sp Q03164 KMT2A_HUMAN
5103	YSEFTSTSGTGHNQTR22000000000	sp P48960 CD97_HUMAN
5104	SGSSQELDVKPSASPQER202200000000200000	sp Q9UQ35 SRRM2_HUMAN
5105	TSPADHGGSVSGSESGSAVDSVAGEHSVSGR2.02002E+20	sp Q5T4S7 UBR4_HUMAN
5106	FFTTGSDSESESSLGEELVTKPVGGNYGK2.02E+27	sp B5ME19 EIFCL_HUMAN
5107	SDSEESGSEEEEEEEEEEQPAAQPPTLPVEEK2.0200202E+33	sp Q99613 EIF3C_HUMAN
5108	MSPNETLFLESTNK20000	sp P51532 SMCA4_HUMAN
5109	LDNTPASPPRSPAEPNDIPIAK2.0000002E+19	sp Q9NQL2 RRAGD_HUMAN
5110	LSSAGPRSPYCK20000200000	sp Q9HB90 RRAGC_HUMAN
5111	ALEAASLSQHPPSLCISDSEEEEEER20200000000	sp 095359 TACC2_HUMAN
		sp Q6PL18 ATAD2_HUMAN
		sp 043159 RRP8_HUMAN
		sp P05534 1A24_HUMAN
		sp Q09160 1A80_HUMAN
5112	KGGSYSQAASSDSAQGSVSLTACK20002002000000	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
5113	CVSVQTDPTDEIPTKK2000000000000000	sp P18583 SON_HUMAN
5114	AQTTPGPSLSGSKSPCPQEK2000000000020000000	sp Q9UQ35 SRRM2_HUMAN
5115	RGTSPRPPEGGLGYSQLGDDDLK2.2E+21	sp P21127 CD11B_HUMAN
		sp Q9UQ88 CD11A_HUMAN
5116	KASVDPSTESSPAPQEGSEQPASPASPLSSR2E+30	sp Q66K14 TBC9B_HUMAN
5117	QFIAAQNLGPASGHGTPASSPSSSSLPSPMPTPR200200000	sp Q8WX93 PALLD_HUMAN
5118	SPSFGGASEASAR2000000000000000	sp Q8WVT3 TPC12_HUMAN
5119	IVSAQSLAEDDVE200200000000	sp Q15388 TOM20_HUMAN
5120	SSRSSPELTR2022000000	sp Q9UQ35 SRRM2_HUMAN
5121	GSSPGIQDTLEAEDGAFETDEAPEDR20000000000000000000	sp Q9UNX4 WDR3_HUMAN
5122	ACASPSAQVEGSPVAGSDGSQPAVK2.0000002E+22	sp Q9UFC0 LRWD1_HUMAN
5123	VISDESIDGGSDVEFKPDTK20000200000000000	sp P52701 MSH6_HUMAN
5124	QNPEQSADEDAEK2000000000	sp B5ME19 EIFCL_HUMAN
		sp Q99613 EIF3C_HUMAN
5125	DGDSYDPYDFSDTEEMPQVHTPK2.0003E+21	sp P25205 MCM3_HUMAN
5126	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2.0002E+25	sp Q8WUZ0 BCL7C_HUMAN
5127	SPPPEVDPTSTK200000002000000	sp P46013 KI67_HUMAN
5128	GAAEEAELESDDEEKPVKQDDFPK20000000000000000	sp Q9UBB9 TFP11_HUMAN
5129	TASESISNLSEAGSIK20002000	sp P30622 CLIP1_HUMAN
5130	AASPSQSVR202002000	sp Q8IYB3 SRRM1_HUMAN
5131	TKPTQAAGPSSPKPPTPEETK2.000000000002E+19	sp P18887 XRCC1_HUMAN
5132	CSVSLSNVEAR2000000000	sp P47712 PA24A_HUMAN
5133	QSSLTFQSSDPEQMR2000000000000000	sp P47712 PA24A_HUMAN
5134	STSAPQMSPGSSDNQSSSPQAQK2.00002E+23	sp Q53SF7 COBL1_HUMAN
5135	ASETVSEASPGSTASQTGVPTQVVQQVQGTQQR20000000000000000000	sp Q14157 UBP2L_HUMAN
5136	SYSPDGKESPSDKK2000002000000	sp P22670 RFX1_HUMAN
5137	LIHGEDSDSEEEEEGRGSSGCSEAGGAGHEEGR2.020000000022E+27	sp P43243 MATR3_HUMAN
5138	GPTSSPCEEEGDEGEEDRTSDLR2.2E+20	sp Q9C0C9 UBE20_HUMAN
5139	GSPSPEKPDVEK202000000000	sp Q86VM9 ZCH18_HUMAN
5140	NSDLLTSPDVGLLK2000000000	sp Q93074 MED12_HUMAN
5141	ASDKDITVSK2000000000	sp P05412 JUN_HUMAN
		sp P49790 NU153_HUMAN
5142	TEPAAEAEAAAGPSESPSPAEEELPGSHAEPVPAQGEAPGEQAR2.02E+33	sp P55884 EIF3B_HUMAN
5143	VQDTSNTGLGEDIHQLSK20000000000000000	sp Q8IWA0 WDR75_HUMAN
5144	EQYVPPRSPK2000	sp Q71RC2 LARP4_HUMAN

5145	LSSASTGKPLSVEDDFEK20002000000000000000	sp 075152 ZC11A_HUMAN
5146	HEAPSSPISGQPCGDDQNASPSK2E+19	sp 075376 NCOR1_HUMAN
5147	SSIEDAQCPGLPDLIEENHVVNK2E+22	sp Q15398 DLGP5_HUMAN
5148	SPPAVSPSPSTDHSSHFSNFNDSR2.000002E+26	sp P42694 HELZ_HUMAN
5149	SPGNLTER200000000	sp Q9H0F6 SHRPN_HUMAN
5150	SNSNAVSAIPINK2000000000000	sp P05549 AP2A_HUMAN
5151	SNSEVEDVGPTSHNR20000000000000000000	sp Q13206 DDX10_HUMAN
5152	SPSMVSPSPGWVSPK20000002000002000	sp Q86UU0 BCL9L_HUMAN
5153	KLSVLGK200000	sp Q8NFC6 BD1L1_HUMAN
5154	AQTPPISSLPTSPSDEVGR2000000002000000000	sp Q15751 HERC1_HUMAN
5155	EEGPPPPSPDGASSDAEPEPPSGR20000000000000000000	sp Q9NTJ3 SMC4_HUMAN
5156	LDSSPSVSTLAAK2000000000000	sp Q13428 TCOF_HUMAN
5157	QLPALDGLMGPESPPAQEEEEAPVSPHK2000000000020000	sp Q8N9T8 KRI1_HUMAN
5158	RYSPPPPKR202000000	sp Q8IYB3 SRRM1_HUMAN
5159	SLSPGVS202000000	sp Q9UKV3 ACINU_HUMAN
5160	SLSSPTVTLAPLEGAK20220000000000000000	sp Q96PU5 NED4L_HUMAN
5161	DSGSDQDLDGAVRASDLEDEESAAR2.02000000000002E+25	sp Q86VM9 ZCH18_HUMAN
5162	GHTASESDEQQWPEEK2000000000000000	sp Q9NTI5 PDS5B_HUMAN
5163	ASVALTPTAPSR20000200	sp P28290 SSFA2_HUMAN
5164	SKLSPSPSLR20200000	sp Q5T200 ZC3HD_HUMAN
5165	MSGGSTMSSGGNTNNSNSK2E+19	sp Q86U70 LDB1_HUMAN
5166	AGMSSNQSISSPVLDVPR1000020220000000000	sp Q9UQ35 SRRM2_HUMAN
5167	STSAPQMSPGSSDNQSSSPQPAQQK1.00000000002E+19	sp Q14157 UBP2L_HUMAN
5168	SPISDNSGCDAPGNSNPSPSSAESEK2.002002E+29	sp Q99590 SCAFB_HUMAN
5169	ATSPSSSVSGDFDDGHHSVSTPGPSR2.2E+22	sp Q86U86 PB1_HUMAN
5170	RYPSSISSPQK200020000	sp Q14157 UBP2L_HUMAN
5171	KPIKSPSKDASSGK20200000000	sp Q13523 PRP4B_HUMAN
5172	AENNSEVGASGYGVPGPTWDR20000000000000000000	sp Q08211 DHX9_HUMAN
5173	SESPCESPYPNEK2000000000000	sp Q8NI27 THOC2_HUMAN
5174	GSGSQSSVPSVDQFTGVGIR20000000000000000000	sp Q9UQE7 SMC3_HUMAN
5175	SGSSQELDVKPSASPQER2000000000000200000	sp Q9UQ35 SRRM2_HUMAN
5176	WAHDKFSGEIEEDDESGTENREEK2.00000000002E+20	sp Q9Y2W1 TR150_HUMAN
5177	ILDEDSWSDGEQEPITVDQTWR20200000000000000000	sp Q9H9L4 KANL2_HUMAN
5178	RSPSPYYSR202000000	sp P62995 TRA2B_HUMAN
5179	ASSPSLIER20000000	sp Q13595 TRA2A_HUMAN
5180	LPSTSGSEGVPR202000000000	sp A1L170 CA226_HUMAN
5181	TSLDIEEYSDTEVQK20000000	sp P29317 EPA2_HUMAN
5182	APQQQPPPPQPPPPQPPPPQPPPPPSYSPAR20000	sp P51580 TPMT_HUMAN
5183	KAEDSDSEPEPEDNVR2000000000000	sp Q9BUJ2 HNRL1_HUMAN
5184	ESKDKPEIEDVGSDEEEEK200000000	sp Q9H0D6 XRN2_HUMAN
		sp Q14568 HS902_HUMAN
		sp P30505 1C08_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp Q29960 1C16_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
5185	GGSCSQAASSNSAQGSDESLIACK2000000000	sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30510 1C14_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
5186	NYQQNYQNSGESGEKNEGSESAPEGQAQR2020000000000	sp P67809 YBOX1_HUMAN

5187	LSPSPSSR200000	sp Q03252 LMNB2_HUMAN
		sp P20700 LMNB1_HUMAN
5188	NATDLQNSSMSEEELTK2000000000	sp P40855 PEX19_HUMAN
5189	TASISSPSEGTPTVGSYGCTPQSLPK2.02E+25	sp Q6PKG0 LARP1_HUMAN
5190	KESESEDSSDEPLIKK2000022000000000	sp P35659 DEK_HUMAN
5191	EGDPVSLSTPLETEFGSPSELSR2002000	sp Q9BUR4 WAP53_HUMAN
5192	QERLSPEVAPPAHR200000000000	sp Q8TAD8 SNIP1_HUMAN
5193	QSGSSQPMEVQEGYFGSGDDPYSSAEPHVSGVK2E+31	sp Q13263 TIF1B_HUMAN
5194	EHYPVSSPSSPPAQPGVSR2000020000000000	sp 075179 ANR17_HUMAN
5195	AQRQSCTSVGR20000000	sp Q9NYZ3 GTSE1_HUMAN
5196	RHSSDINHVLVTQGR22000000000000	sp Q5TON5 FBP1L_HUMAN
5197	RRSPPPGPDGHAK200000000000	sp P57081 WDR4_HUMAN
5198	HSQSPPEPHSPTEPPAWGSSIVK2.00002E+19	sp Q5VZ89 DEN4C_HUMAN
5199	KPSGSPDLWKLSPDQR2000000200000	sp Q96JM3 CHAP1_HUMAN
5200	LSHSDEKPYQCPVCQQR2000000000000000	sp P56270 MAZ_HUMAN
5201	VSEEQTPPSPAGAMSTAMGR20000000001000	sp Q16666 IF16_HUMAN
5202	STSAPQMSPGSSDNQSSSPQPAQQK2.000001E+25	sp Q14157 UBP2L_HUMAN
		sp Q9Y4E1 FA21C_HUMAN
5203	SRPTSFADELAAR220000000000	sp Q641Q2 FA21A_HUMAN
		sp Q5SNT6 FA21B_HUMAN
5204	SLSPGKENVSALDMEK2000000000000000	sp Q9H307 PININ_HUMAN
5205	TSAVSSPLLDQQR2000000000	sp Q5T200 ZC3HD_HUMAN
5206	KADTEGKSPSHSSSNRPFTPPTSTGGSK2000020000000000	sp Q15648 MED1_HUMAN
5207	STSAPQMSPGSSDNQSSSPQPAQQK1.0000000022E+19	sp Q14157 UBP2L_HUMAN
5208	TPSSSSTLAYSRL200000000000	sp Q6Q0C0 TRAF7_HUMAN
5209	HFEASCGQLSPAR20000	sp Q96N21 AP4AT_HUMAN
5210	VNPHKVPASSVDSNIPSSQGYK200220020000000000	sp Q9UHB7 AFF4_HUMAN
5211	QSGDGGESTEPVENK2000000000000000	sp 015234 CASC3_HUMAN
5212	EGSQGELTPANSQSR2000000000	sp Q13098 CSN1_HUMAN
5213	LDSSEMDSSENYDTMSSPLPGK2.000020000001E+20	sp Q9NTI5 PDS5B_HUMAN
5214	KPSGSPDLWK2000000000	sp Q96JM3 CHAP1_HUMAN
5215	LNQPGTPTR20000	sp P36507 MP2K2_HUMAN
5216	TAATAAAAAAEPAPPPPPPEEDPEQDSGPEDLPLVR200000000000	sp P06400 RB_HUMAN
5217	RPPSPDIVLSDNEQPSSPR200000020000002000	sp Q86YP4 P66A_HUMAN
5218	TEARSSDEENGPPSSPDLDR220000000000000000	sp Q96B36 AKTS1_HUMAN
5219	RGEADAPFSEPGTTSTQRPSPETATK220000000	sp Q9UQ35 SRRM2_HUMAN
5220	ESGVVAVSPEKSESPQKEDGLSSQLK2.0002E+19	sp Q96T58 MINT_HUMAN
5221	SRDESASETSTPSEHSAAPSPQVEVR2.020222E+22	sp Q92614 MY18A_HUMAN
5222	SVASSQPAKPTK20002000000000	sp Q07157 ZO1_HUMAN
5223	LQSPLEPHSK2000000000	sp 015211 RGL2_HUMAN
5224	FCDSPTSLEMR2002000000	sp 015014 ZN609_HUMAN
5225	HELSPQK200000	sp Q9BXP5 SRRT_HUMAN
		sp E9PAV3 NACAM_HUMAN
5226	VQGEAVSNIQENTQTPTVQEESEEEEEVDGTEVVK2E+23	sp Q13765 NACA_HUMAN
		sp Q68EM7 RHG17_HUMAN
5227	SPSPPTQHTGQPPGQPSAPLSAPR20000000000000000000	sp Q8IX03 KIBRA_HUMAN
5228	ETNTETPAPSPTVVRPK20002000000000	sp Q92508 PIEZ1_HUMAN
5229	TASELLDR20000000	sp P07948 LYN_HUMAN
5230	GKDSLDDGVDLK2000000000	sp Q96AY4 TTC28_HUMAN
5231	SQSPVTVKPKPPAR200000000000000000	sp Q9Y4R8 TELO2_HUMAN
5232	TPQGSPSPNTPCLPEAAVSQPGSAVASDWR2E+24	sp Q6Y7W6 PERQ2_HUMAN
5233	ALSSGGSITSPPLSPALPK20002000000	sp Q8NBJ9 SIDT2_HUMAN
5234	VDSMSSVEEDDYDTLTDIDSDKNVIR2.022E+24	sp P02545 LMNA_HUMAN
5235	LSPSPTSQR2000000000	sp Q9C0C2 TB182_HUMAN
5236	GSGGLFSPSTAHPDGAALGQR2E+20	

5237	SSPNPFVGSPPK20000	sp P98082 DAB2_HUMAN
5238	KRSPSPSPTPEAK202000000000	sp Q8TAQ2 SMRC2_HUMAN
5239	KVVEAVNSDSDSEFGIPK202020000000	sp Q02880 TOP2B_HUMAN
5240	HSLSGSSPGMK20000000000	sp Q9UQ35 SRRM2_HUMAN
5241	RASPPDPSPSAASASER200000020000000000	sp Q9Y2F5 ICE1_HUMAN
5242	KGSISSTQDTPVAVEEDCSLASSK2. 2E+20	sp Q9BZ71 PITM3_HUMAN
5243	SPGMLEPLGSSR2001000000000	sp Q9UQ35 SRRM2_HUMAN
5244	VAGGAAPSKPASAK2000	sp Q14978 NOLC1_HUMAN
5245	ASASGSGAQVGGPISSGSSASSVTVTR22000000	sp P02545 LMNA_HUMAN
5246	LQEEGGGSDDEETGSPSEDGMQ SAR200000000000000	sp Q13610 PWP1_HUMAN
5247	TSSKESPIPSPTSDRK22000200000000	sp Q01082 SPTB2_HUMAN
5248	SSSPTAGPSTEGAEGPEEK220000000000000000	sp Q8NHM5 KDM2B_HUMAN
5249	SPAGPAATPAQAQAASPR2000000002000	sp Q13428 TCOF_HUMAN
5250	MSHSSSDTDINEIHTNHK200000000000	sp P46100 ATRX_HUMAN
5251	DTQSPSTCSEGLLGWSQK200000000000000000	sp Q9C0C2 TB182_HUMAN
5252	TNLTSQSSTTNLPGSPGSPGSPGSPGSPGSPVK200200200000000000	sp Q6IN85 P4R3A_HUMAN
5253	EDDDVIVNKPHVSDEEEEEPPFYHHPFK2000000000000000	sp P18583 SON_HUMAN
5254	SPQPVCSPAGSEGTK20000000002000000	sp Q6KC79 NIPBL_HUMAN
5255	EWYQSTIPQSPSPAPDDPEEGR2000000200000000000	sp Q08499 PDE4D_HUMAN
5256	HGGSPQPLATTPLSQEPVNPPEASPTR2. 000002E+25	sp Q9UQ35 SRRM2_HUMAN
5257	YYSIDDNQNK200000000	sp Q8NI08 NCOA7_HUMAN
5258	LTVSDGESGEEK2000200000	sp P46100 ATRX_HUMAN
5259	TSPPCSPANLSR200020000000	sp P49585 PCY1A_HUMAN
5260	TGSSSPGGPPKPGSQLDSMLGSLQSDLNK2E+26	sp P49023 PAXI_HUMAN
5261	EQSEVSVSPR2000	sp Q7L4I2 RSRC2_HUMAN
5262	SRSPSPELNK2020000000000	sp Q9UQ35 SRRM2_HUMAN
5263	LAGGQTSQPTTPLTSPQR2000020000	sp O95155 UBE4B_HUMAN
5264	NSLLAGDDDTMSVISGSSR12002002000	sp Q8N3U4 STAG2_HUMAN
5265	LLKPGEPESEYTDDEEDTKDHNK200000000000000	sp O15173 PGRC2_HUMAN
5266	SGSSQELDVKPSASPQER20000000020000000	sp Q9UQ35 SRRM2_HUMAN
5267	RTSLGEVSK20000000	sp Q9ULJ8 NEB1_HUMAN
5268	HASSPESPKPAPAPGSHR20002000000000000	sp Q9UQ35 SRRM2_HUMAN
5269	DESASETSTPSEHSAAPSPQVEVR2. 020222E+22	sp Q92614 MY18A_HUMAN
5270	VPDEEENEESDNEK200000	sp P11388 TOP2A_HUMAN
5271	FEPESPGFESR20000000	sp Q9H5H4 ZN768_HUMAN
5272	VQIPVSRPDPPEVSDNEEDSYDEEIHDP23000000000	sp Q07157 ZO1_HUMAN
5273	SPSPLSGHVAQAFPTK20200000000000000	sp Q6P0Q8 MAST2_HUMAN
5274	HAPSLHGSTELLPLSR2000000200	sp Q9H5N1 RABE2_HUMAN
5275	AGTATSPAGSSPAVAGGTQ RPAEDSSSSEESDSEEEK2E+35	sp Q13428 TCOF_HUMAN
5276	SSDEENGPPSSPDLDR2000000022000000	sp Q96B36 AKTS1_HUMAN
5277	TKPTQAAGPSSPQKPPTPEETK2000002000000	sp P18887 XRCC1_HUMAN
5278	SLESVLSLGRPTGGGSSPPEIR2E+23	sp Q13470 TNK1_HUMAN
5279	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2. 00000020002E+32	sp Q8WUZO BCL7C_HUMAN
5280	SPPAPGLQPMR200000000100	sp P15408 FOSL2_HUMAN
5281	FPVPELSDQEGDSSSEED2000002220000	sp Q9BRX2 PELO_HUMAN
5282	TSADAQEPASPVVSPQQSPPTSPTWR2000200020020000000	sp Q96D71 REPS1_HUMAN
5283	DSDYVYPSLESDENPIFK2000000000	sp O75151 PHF2_HUMAN
5284	VVDYSQFQESDDADEDYGRDSGPPTK20000000000000000	sp Q9H1E3 NUCKS_HUMAN
5285	YTGEEDGAGGHSPAPPQTEECLR2000000000000	sp O94762 RECQ5_HUMAN
5286	TNLTSQSSTTNLPGSPGSPGSPGSPGSPGSPVK200200200000000	sp Q6IN85 P4R3A_HUMAN
5287	EPTPSIASDISLPIATQELR202000000000000000	sp A0FGR8 ESYT2_HUMAN
5288	AFNLHLSSESEDETEHPVPIILSNEDGR2. 022000002E+24	sp Q8IWU2 LMTK2_HUMAN
5289	VPKSPEHSAEPIR20000000000	sp Q9ULH0 KDIS_HUMAN
5290	NSSPGEASLLEK20000000000	sp Q76FK4 NOL8_HUMAN

5291	SVSPPPKR200000000	sp Q13247 SRSF6_HUMAN
5292	ASSPSPLTIGTPESQR202000000000000	sp Q9NPI6 DCP1A_HUMAN
5293	NLNSNLFSVNR2000000000	sp P52948 NUP98_HUMAN
5294	RSPSAPAATSSASSSTSSYK2E+19	sp Q15345 LRC41_HUMAN
5295	LLPSVWSESEDGAGSPPPYR2000000	sp Q9UIS9 MBD1_HUMAN
5296	HSASPVVFTSAR2000002000	sp O43166 SI1L1_HUMAN
5297	VVLGDGVQLPPGDYSTTPGGTLFSTTPGGTR3000000000020000000	sp Q13541 4EBP1_HUMAN
5298	GLYDGPVCEVSVTPK2000	sp Q16555 DPYL2_HUMAN
5299	VSPASPGP200020000	sp Q9Y4U1 MMAC_HUMAN
5300	EISPGSGPEIR20020000000	sp O43491 E41L2_HUMAN
5301	LAELPAAQPSAEDSDTEDDSEAEQTER202000200000000	sp O95714 HERC2_HUMAN
5302	ALQAPHSPSK20000	sp Q96AQ6 PBIP1_HUMAN
5303	RHSDPWGR2000000	sp Q92576 PHF3_HUMAN
5304	DGDSYDPYDFSDTEEMPQVHTPK200000100000000	sp P25205 MCM3_HUMAN
5305	SDSASSEPVGIYQGF2002000000000000000	sp Q05655 KPCD_HUMAN
5306	KAPAGQEEPPTSSPLSAEQQLDR2200000000000	sp P13051 UNG_HUMAN
5307	NAPPGGDEPLAETESESEAEAGFSPVVDVK2.0202E+19	sp Q8NC44 F134A_HUMAN
5308	RKGSSEGEDEEGGER2020000000000000000	sp Q6PD62 CTR9_HUMAN
5309	DGDSYDPYDFSDTEEMPQVHTPK3000000000000002000	sp P25205 MCM3_HUMAN
5310	ISSSSF200000000	sp P05787 K2C8_HUMAN
5311	ESYTEEIVSEASHVSGISR200000000000000	sp Q5QJE6 TDIF2_HUMAN
5312	MLPHAPGVQQAIPEDAIPESGDEDEDDPKR1E+33	sp Q13547 HDAC1_HUMAN
5313	SPSSKPSLAVAVDSAPEVAVK2000000000000000000	sp O60508 PRP17_HUMAN
5314	TDCEPVGLEPAVEQSSAASELAEASSEELAEAPTEAPSPEAR2.0000000000002E+42	sp Q15424 SAFB1_HUMAN
5315	SFSISPSR202020000	sp Q14151 SAFB2_HUMAN
5316	LFEDDSDNEKLFDEEEDSSEKLFDDSDER2.000000000022E+23	sp P18583 SON_HUMAN
5317	GSQHSQEQPAPQPSGGDPSPPQER2000000	sp O43719 HTSF1_HUMAN
5318	SHTSEGAHLDITPNSGAAGNSAGPK2.2E+23	sp Q8WXE0 CSK12_HUMAN
5319	AASQGSPLLPAQR2000000000000000000	sp Q92597 NDRG1_HUMAN
5320	ETPAATEAPSSTPK2000	sp Q5TCZ1 SPD2A_HUMAN
5321	YGSPQR200000	sp P80723 BASP1_HUMAN
5322	SITSTPLSGK20000002000	sp O15234 CASC3_HUMAN
5323	AGSISTLDSLDFAR2022002000000	sp Q9P0K7 RAI14_HUMAN
5324	SPSPAPPPR2000000000	sp Q9P260 K1468_HUMAN
5325	TGSGSPFAGNSPAR20000	sp Q8IYB3 SRRM1_HUMAN
5326	RQSVSPPYK20200000	sp Q9UPT8 ZC3H4_HUMAN
5327	IVEPEVGEDSEVEGDWR2000000000	sp Q9NYV4 CDK12_HUMAN
5328	ASPSQGSDERESLTR2000002000020000	sp P55081 MFAP1_HUMAN
5329	YSHSYLSDSDTEAK200000000	sp P37275 ZEB1_HUMAN
5330	KPSDSLVAASSR2000000000	sp Q92614 MY18A_HUMAN
5331	NSCNVLHPQSPNNSNR20000000	sp Q5T0W9 FA83B_HUMAN
5332	HVSPVTPPR20020000	sp Q7Z333 SETX_HUMAN
5333	SGSGISVISSTVDQR2000000000000000	sp Q13443 ADAM9_HUMAN
5334	HLSTPSSVSPEPQDSAK2000000000	sp Q15811 ITSN1_HUMAN
5335	AIGGIILTASHNPGPNGDFGIK2000000000000000	sp Q9NZN5 ARHGC_HUMAN
5336	DGGSGNSTIIVSR20020000000	sp P36871 PGM1_HUMAN
5337	APSTSPSFEQTQETYTVAHEENVNR2000000000000000000	sp Q7Z6Z7 HUWE1_HUMAN
5338	TEFLHSQNSLSPR202000	sp Q9BUT9 F195A_HUMAN
5339	TEPAEAEAAASGPSESPSPAEEELPGSHAEPVPAQGEAPGEQAR2E+29	sp Q96FS4 SIPA1_HUMAN
5340	EVENEDDDDDSDKEK200000	sp P55884 EIF3B_HUMAN
5341	SSSQRSPSPGNHTSNSSNASNATVVPQNSSAR2.02000002E+28	sp Q9P287 BCCIP_HUMAN
5342	LHQLSGSDQLESTASHR200000000000000	sp Q9BTA9 WAC_HUMAN
5343	SHSGTSPDNTAPPPPPRPQPSHSR2.02E+23	sp O60271 JIP4_HUMAN
		sp Q96D71 REPS1_HUMAN



5344	HKLTVSDGESGEEK2000200000	sp P46100	ATRX_HUMAN
5345	SSTPLPTISSAENTR200020000000000	sp P42166	LAP2A_HUMAN
5346	RSSQPSPTAVPASDSPPTK202000000000000	sp P42167	LAP2B_HUMAN
5347	ALQSPALGLR20000000	sp Q3KQU3	MA7D1_HUMAN
5348	SLSPLGGR202000000	sp P46940	IQGA1_HUMAN
5349	RASPSKPASAPASR200000000000000	sp Q14004	CDK13_HUMAN
5350	SLESINSR200000000	sp P27816	MAP4_HUMAN
5351	VSPAHSPPENGLDK200020000000000	sp P62888	RL30_HUMAN
5352	TLSTHSVPNISGATCSAFASPFGCPYSHR2. 2E+27	sp Q04726	TLE3_HUMAN
5353	YSPTSPTYSPK200300000000	sp P28290	SSFA2_HUMAN
5354	EPDATPPSPR2000	sp P24928	RPB1_HUMAN
5355	SSPNPFVGSPPK200000020000	sp P28715	ERCC5_HUMAN
5356	YSHSYLSDSDTEAK20000202000000	sp P98082	DAB2_HUMAN
5357	SAYQEYSDSDVPEELK20200000000	sp Q92614	MY18A_HUMAN
5358	SPGTPRPPGASK2000000000000	sp Q96JG6	CC132_HUMAN
5359	SPQLSLSPRASP200002000	sp Q9Y3X0	CCDC9_HUMAN
5360	SLNLVDSQPILLEK200000000000000	sp 095785	WIZ_HUMAN
5361	ISQSQPVR20000000	sp P46019	KPB2_HUMAN
5362	AQSSPASATFPVSVQEPPTKPR200000000000000	sp 095071	UBR5_HUMAN
5363	GGLNTPHESDFSGVTPQR2000000000020000	sp P56524	HDAC4_HUMAN
5364	QEQINTEPLEDTVLSPTK20000	sp Q99459	CDC5L_HUMAN
5365	HPEPVPEEGSEDELPPQVHK2000000000000	sp 015446	RPA34_HUMAN
5366	KPIKSPSKDASSGK22000	sp Q7Z3C6	ATG9A_HUMAN
5367	RHSHSHSPMSTR202020000000	sp Q13523	PRP4B_HUMAN
5368	APSPTKPLR20000000	sp P62995	TRA2B_HUMAN
5369	NSGIWESPELDR2000000000000	sp Q96FF9	CDC45_HUMAN
5370	SPTGPSNSFLANMGTVAHK20000100000000	sp 015085	ARHGB_HUMAN
5371	LGSLAR200000	sp Q96I25	SPF45_HUMAN
5372	SQTPPGVATPPIPK200000000000000	sp Q5SW79	CE170_HUMAN
5373	VQIPVSRPDPVSDNEEDSYDEEIHDP20000020000000000	sp Q00G26	PLIN5_HUMAN
5374	VQQEIDSDSPVAR200200000	sp Q15648	MED1_HUMAN
5375	KESKEETSIDVAGKPNEVTK2. 000002E+19	sp Q07157	ZO1_HUMAN
5376	SEACPCQPDGSPPLAEEEEK2000000000000	sp Q6T4R5	NHS_HUMAN
5377	GPGSPRTPPAPGPPSFEEQLR2. 0002E+19	sp P53985	MOT1_HUMAN
5378	SDHEDNDPSVDVQHSGSEAPNDDDEGHR2. 00000002000002E+29	sp Q14160	SCRIB_HUMAN
5379	TESFHSYR2000000	sp Q6ZTN6	AN13D_HUMAN
5380	SSFSQHAR20000000	sp Q8WVC0	LEO1_HUMAN
5381	GGWDHGDQSIQSSR2000200	sp Q9Y597	KCTD3_HUMAN
5382	SLNSVVTSEPKHSLLVGDFR2. 0020002000002E+21	sp P02545	LMNA_HUMAN
5383	GGSTGGGGFDPPPAYHEVVDAEK2E+21	sp Q9P206	K1522_HUMAN
5384	SGSTSSLSYSTWTSSHSDK20000000000000000	sp Q04656	ATP7A_HUMAN
5385	RGSQKSTDSPGADAELPESAAR2002000000000000000	sp Q9UPU5	UBP24_HUMAN
5386	NMTVEQLLTGSPSPTVEPEKPTR1. 000000000022E+23	sp Q9ULD2	MTUS1_HUMAN
5387	LTVENSPK2000	sp Q9Y4K1	AIM1_HUMAN
5388	HKSGSMEEDVDTSPGGDYTSPSSPTSSSR1. 000000000003E+25	sp Q9C0C9	UBE20_HUMAN
5389	LASPSGSTSSGLEVVAPGTSAPGGGPGTLDDSATICR2. 0222E+34	sp 043583	DENR_HUMAN
5390	GQGTTPSGPGVVR2000000000000	sp P08651	NFIC_HUMAN
5391	KKDELSDYAEK2000000	sp Q13263	TIF1B_HUMAN
5392	SPQFINLK200000000	sp Q9UMN6	KMT2B_HUMAN
5393	LGSPVDNLEMDRDDLTDVSVFTR2E+22	sp P46100	ATRX_HUMAN
5394	QSGSPTLDTAPNGR2000000000000	sp Q92576	PHF3_HUMAN
5395	ASVALTPTAPSR200000000200	sp Q5VUA4	ZN318_HUMAN
		sp Q9H2J7	S6A15_HUMAN
		sp P28290	SSFA2_HUMAN

5396	SVSASSGPCSPAAPHPPPPR2. 02002E+21	sp Q86UY5 FA83A_HUMAN
5397	RGTGQSDSDIWDITALIK200200200000000000	sp Q16637 SMN_HUMAN
5398	APQTSSSPPPVR22000000	sp Q8IYB3 SRRM1_HUMAN
5399	TPAPPEPGSPAPGEGPSGR200000000000	sp P0C1Z6 TFPT_HUMAN
5400	SGSPSDNSGAEEMEVSLAKPK200200000000000000	sp P31749 AKT1_HUMAN
5401	SAGGSSPEGGEDSDREDGNYCPPVK2. 0000002E+20	sp Q9H6Z4 RANB3_HUMAN
5402	LTPVSLNSPIK20000	sp P46100 ATRX_HUMAN
5403	ESSPIPSPTSDR20002000000	sp Q01082 SPTB2_HUMAN
5404	KGSPTPGFSTR2000000000	sp Q6ZRV2 FA83H_HUMAN
5405	SSPPLRTPDVLESSGPAVR2. 200002E+19	sp 075676 KS6A4_HUMAN
5406	SPGSTPTTPTSSQAPQK2000000	sp P35658 NU214_HUMAN
5407	LQTTDNLPMSPPEEFDEVSR20000000000	sp P42224 STAT1_HUMAN
5408	IESDTEETQDTSVDHNETGNTGESSVEENEK2002002000000000	sp Q6PL18 ATAD2_HUMAN
5409	VPGEQGSDEEHCK20000000	sp Q9UNS1 TIM_HUMAN
		sp Q641Q2 FA21A_HUMAN
5410	SRPTSFADLAAR20002000000000	sp Q5SNT6 FA21B_HUMAN
		sp Q9Y4E1 FA21C_HUMAN
5411	SKPSPQKQALGDGIAPPQK200000000000000000	sp Q9H9J4 UBP42_HUMAN
5412	SYQNSPSSDDGIRPLPEYSTEK202000000000000000	sp Q15648 MED1_HUMAN
5413	ALSSGGSITSPPLSPALPK20000000000	sp Q6Y7W6 PERQ2_HUMAN
5414	SPTKAPHLQLIEGK2000000000000	sp 043491 E41L2_HUMAN
5415	QGSITSPQANEQSVTPQR200200000000000000	sp Q9UQ35 SRRM2_HUMAN
5416	GFSVVADTPELQR2000000	sp Q14847 LASP1_HUMAN
5417	RQSSGSATNVASTPDNR20000000000	sp Q7Z460 CLAP1_HUMAN
5418	LEPVYSPGSPPGDPR20002000000000	sp 075676 KS6A4_HUMAN
5419	GVGSEPGPTAPYLGR20000000000000	sp Q9C0B5 ZDHC5_HUMAN
5420	HSVTAATPPPSPTSGESDLSNLLQSPSSAK2. 002E+19	sp Q96QU8 XPO6_HUMAN
5421	TSAQHALTSVSRGSSLK20020000	sp P61073 CXCR4_HUMAN
5422	TEEPLGSPHSGTVESPGEGPQDTR2. 000000002E+19	sp Q9NWS9 ZN446_HUMAN
5423	RRSPSPAPPPR2020000000	sp Q8IYB3 SRRM1_HUMAN
5424	GDTVSAAPCSAPLAR20000000000	sp Q15477 SKIV2_HUMAN
5425	NYQQNYQNSESGEKNESSESAPEGQAQQR2. 000000002E+19	sp P67809 YBOX1_HUMAN
5426	GPSQEEEDNHSDEEDR2000000	sp Q86VM9 ZCH18_HUMAN
5427	ATPATAPGTSPR2000	sp Q96P47 AGAP3_HUMAN
5428	LASPSGSTSSGLEVVAVEGTSAPGGPGTLDDSATICR2. 02E+34	sp Q13263 TIF1B_HUMAN
5429	LSPSPTSQR2000000	sp P02545 LMNA_HUMAN
5430	RGSSGSVDETLFALPAASEPVIR200020000000000000	sp Q9NQC3 RTN4_HUMAN
5431	RRSDGVDGFEAEGK2000000000000	sp Q96KR1 ZFR_HUMAN
5432	ALSSGGSITSPPLSPALPK200000000000000000	sp Q6Y7W6 PERQ2_HUMAN
		sp A4D2H0 CTGEF_HUMAN
		sp Q86UF2 CTGE6_HUMAN
5433	SNSELEDEILCLEK20000000000000	sp 015320 CTGE5_HUMAN
		sp P0CG41 CTGE8_HUMAN
		sp A4FU28 CTGE9_HUMAN
		sp Q8IX94 CTGE4_HUMAN
5434	LSPSPTSQR202000000	sp P02545 LMNA_HUMAN
5435	QEAKPQQAAGMLSPK2000	sp Q13428 TCOF_HUMAN
5436	ESASPTIPNLDLLEAHTK200000000000000000	sp 075151 PHF2_HUMAN
5437	TPLDLFAHFGPEPGDHSPLPPSAPSPTR20000	sp Q8N1G0 ZN687_HUMAN
5438	NQSPTEAEKPASSSLPSSPPPQLLTR2E+24	sp Q99567 NUP88_HUMAN
5439	SQTPPGVATPPIPK200000002000000	sp Q15648 MED1_HUMAN
5440	EPTSSENEGISTIPSPVTSPLSR20020000000	sp Q8NEL9 DDHD1_HUMAN
5441	ANSPSLFGTEGKPK2000000000000	sp Q96B97 SH3K1_HUMAN
5442	KEDDSSQSTCVVDTTTR2202200000000000	sp Q5SNT2 TM201_HUMAN

5443	SKDASPINR2000200000	sp Q13523 PRP4B_HUMAN
5444	SKTPSSPCASK20020000000	sp Q9NQS7 INCE_HUMAN
5445	LCDFGSASHVADNDITPYLVS200	sp Q13523 PRP4B_HUMAN
5446	SPISDNSGCDAPGNSNPSLSPSSAESEK2.002E+29	sp Q99590 SCAFB_HUMAN
5447	RPDHSGGSPSPPTSEPAR200200002000000	sp Q8TAD8 SNIP1_HUMAN
5448	AFKPEETSSNSDPPSPVVLNNSHPVPR2000200000000000000	sp 060293 ZC3H1_HUMAN
5449	TSQVGAASAPAKESPRK20000	sp Q13428 TCOF_HUMAN
5450	HSDSGTSEASLSPSSPPSRPR220000000	sp Q7Z4S6 KI21A_HUMAN
5451	IIPLEEGLQLPSPTATSQPLESDAVECLNYQHYK2E+19	sp P61978 HNRPK_HUMAN
5452	GSISLNSSPR20002000	sp Q9NYL2 MLTK_HUMAN
5453	SPQRSPLQSAESSPTAGK2000200000022000000	sp Q8IWB9 TEX2_HUMAN
5454	LLKPGEEPSEYTDEEDTKDHNK3200000000000	sp 015173 PGRC2_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
5455	GGSYSQAASSDSAQGSVSLTACK20000000000000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
5456	RDSLTTGSSDLYK20000000000	sp Q14671 PUM1_HUMAN
5457	IIPELVGSPTQSTSSR2000000000	sp Q9NSI6 BRWD1_HUMAN
5458	DSGHGSTSVDSEGFSDPDTGSHCSSEYAASSPGDR200000	sp P54278 PMS2_HUMAN
5459	NSSLGSPSNLCGSPGIR202000020000000	sp Q9C0B0 UNK_HUMAN
5460	QEEGADASEEDPTPAGEEDVK2000000000000000	sp Q86X53 ERIC1_HUMAN
5461	GRDSYGGPPR20000000	sp P38159 RBMX_HUMAN
		sp Q96E39 RMXL1_HUMAN
5462	ASSFSGISILTR200000000000	sp Q9Y2K5 R3HD2_HUMAN
5463	LISDSVER20000	sp Q5T200 ZC3HD_HUMAN
5464	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2.0020002E+32	sp Q8WUZ0 BCL7C_HUMAN
5465	LEEPINEETPSQDEEEGVSDVESER20002000	sp Q9BYW2 SETD2_HUMAN
5466	YQEDSDPERSDYEEQQLQK2000020000000000	sp Q6P6C2 ALKB5_HUMAN
5467	VISDESSEDIIGSDVEFKPDTK2000000000000000000	sp P52701 MSH6_HUMAN
5468	YLEEDNSDESDAEGEHGDGAEIEAPPAGPRPGPEPAGLGR2.002E+34	sp Q9H1C4 UN93B_HUMAN
5469	AKPPADPAAAAASPCR20000	sp Q9GZT9 EGLN1_HUMAN
5470	HFSTLKDDDLVEFSDLESEDDERPR2000200000000	sp Q9HCK8 CHD8_HUMAN
5471	DNSPEPNDPEEPQEVSSTPSDKK2E+21	sp Q9HB58 SP110_HUMAN
5472	EEQTDTSDGESVTHHIR2000000000000	sp P36915 GNL1_HUMAN
5473	ETVQTTQSPTPVEK20000000	sp Q9UPN6 SCAF8_HUMAN
5474	SLLDASEEAIKK2000000000000	sp P18206 VINC_HUMAN
5475	HSTENDSPTNVQQ20000000	sp P49354 FNTA_HUMAN
5476	QPTPPFFGR20000000	sp Q96PK6 RBM14_HUMAN
5477	AQVLATIHGHAGAFPAAGDAGEGAPGGGSSPER20000	sp Q86WR7 PRSR2_HUMAN
5478	SVSASSGPCSPAAPHPPPPR2.0000002E+19	sp Q86UY5 FA83A_HUMAN
5479	SGFGEISSPVIR2000002000000	sp P57740 NU107_HUMAN
5480	NLATSADTPPSTVPGTGK2000000000000000	sp Q96Q15 SMG1_HUMAN
5481	EAEQAASEAAGDTPGSSPSSLYEEPLGQPPR220000000000000	sp Q86TC9 MYPN_HUMAN
5482	LQLATLEAGPGAAGAGSE200	sp C9JI98 TM238_HUMAN
5483	APSTSPSFEQTQETYTVAHEENVR2.2E+21	sp Q9BUT9 F195A_HUMAN
5484	RSDSLLSFR2002000	sp Q00587 BORG5_HUMAN
5485	VSPAPGSAPETPEDK2000000000	sp Q13459 MYO9B_HUMAN
5486	SLNSVVTSEPDKHSLLVGDFR2.0020002E+21	sp Q04656 ATP7A_HUMAN
5487	KLSNSSSVSPLILSSNLPVNNK2E+21	sp Q9BVC5 ASHWN_HUMAN
5488	GSISSTSEVHSPNVGLR22000000000000	sp Q8WWQ0 PHIP_HUMAN
5489	VSGVDGYETEGIR2000000000000	sp Q14566 MCM6_HUMAN

5490	SYDVPPPPMEPDHPFYSNISK2E+21	sp P18669 PGAM1_HUMAN
5491	LSPERSSDESSENEPHLPHGVSQR2.000220002E+24	sp Q8N0Y7 PGAM4_HUMAN
5492	IISPGSSTPSSTR2000000	sp Q9BQ52 RNZ2_HUMAN
5493	SVELDLNQAHEETPK2000000000000000	sp Q5VT52 RPRD2_HUMAN
5494	NSTELAPPLPVR20000000000	sp Q14680 MELK_HUMAN
5495	HGTPDPSPR20002000	sp Q9H7D0 DOCK5_HUMAN
5496	QQSQAASSSVTSPIK20000	sp Q9BRD0 BUD13_HUMAN
5497	FIHQQPQSSSPVYGSSAK2000000000	sp O60583 CCNT2_HUMAN
5498	SDSVLPASHGHLPGAGSLER20000	sp P49023 PAXI_HUMAN
5499	SHMSGSPGPGSNTAPSTPVIGGSDKPGMEEK2E+29	sp Q8N4C8 MINK1_HUMAN
5500	GDDLGAQDGVSSPAASPGHVVR200020000000	sp A0FGR8 ESYT2_HUMAN
5501	SGSSSPDSEITELK200020000000000	sp Q86X02 CDR2L_HUMAN
5502	RPPKSPEEEGQMSS20000000020	sp P17812 PYRG1_HUMAN
5503	RREEGPPPPSPDGASSDAEPEPPSGR200000200000000000	sp Q9BRL6 SRSF8_HUMAN
5504	ISLSDMPR20000000	sp Q9NTJ3 SMC4_HUMAN
5505	HSSISPVRLPLNSSLGAELSR2.02E+19	sp Q9ULU4 PKCB1_HUMAN
5506	KLENSPLGEALR200000000	sp Q9NYV4 CDK12_HUMAN
5507	HKSGSMEEDVDTSPGGDYTSPSSPTSSSR33000020000000	sp Q9NX40 OCAD1_HUMAN
5508	HQASDSENEELPKPR20000000000	sp P08651 NFIC_HUMAN
5509	SQSSDTEQQSPTSGGGK20002000000000	sp Q96ST2 IWS1_HUMAN
5510	ASVALTPTAPSR200000	sp P23588 IF4B_HUMAN
5511	SPSPPTQHTGPPGQPSAPQLSAPR2.002E+24	sp P28290 SSFA2_HUMAN
5512	TNKSPEAKPLPGK20000000000	sp Q68EM7 RHG17_HUMAN
5513	NMPGLSAATLASLGGTSSR200	sp P46087 NOP2_HUMAN
5514	MAPTPIPTRSPSDSSTASTPVAEQIER200000000000000000	sp Q32MZ4 LRRF1_HUMAN
5515	VSSPLSPLSPGIK200200000	sp Q16643 DREB_HUMAN
5516	LTQAEISEQPTMATVVPQVPTSPK20000	sp Q9P2B4 CT2NL_HUMAN
5517	ARNSGTQSDGEEKQPLVMER2020000000000000000	sp Q70E73 RAPH1_HUMAN
5518	GLMAGGRPEGQYSEDEDTDDEYK30000020000000	sp Q6P1M3 L2GL2_HUMAN
5519	SSTPSSPTGTSSSDSGHHIGWGER2.2E+21	sp Q9NPQ8 RTC8A_HUMAN
5520	TNSGGGDGPHISSK2000000000000	sp P46019 KPB2_HUMAN
5521	ECLTGESESSSEDEFEK200202020000000	sp Q96EV2 RBM33_HUMAN
5522	YSPPAISPLVSEK30000020000000	sp Q56P03 EAPP_HUMAN
5523	NFIGNSNHGSQSPR2000	sp Q14694 UBP10_HUMAN
5524	REPESPPASAPIPLFGADTIGQR2000000000000000	sp Q03112 EVI1_HUMAN
5525	ATTPPNQGRPDSPVYANLQELK2.000000002E+20	sp Q96AY4 TTC28_HUMAN
5526	HGTDLWIDNMSSAVPNHSPEKK200000	sp Q8IWW6 RHG12_HUMAN
5527	IFDEVMGCFSDSPQSPTFPEAGHTSLYDEDKVR2.0002E+24	sp Q8N6H7 ARFG2_HUMAN
5528	RPACDPEPGESGSSSDEGCTVVRPEK222000000000000	sp Q13085 ACACA_HUMAN
5529	GGDSIGETPTPGASK202000000	sp O15541 R113A_HUMAN
5530	GGRSPDEVTLTSIVPTR2000000000	sp O75533 SF3B1_HUMAN
5531	MLPHAPGVQMQAIPEDAVHEDSGDEDGEDPDKR1E+33	sp Q9UKS6 PACN3_HUMAN
5532	ATSSSNPSSPAPDWYK2000200000000	sp Q92769 HDAC2_HUMAN
5533	LPSPTSPFSSLSQDQAATSK200000000000	sp P46100 ATRX_HUMAN
5534	SSPSARPPDVPQQPQAAKSPSPVQGK2E+24	sp Q8WWI1 LMO7_HUMAN
5535	TEARSSDEENGPPSSPDLDR22000000022000000	sp Q96JP5 ZFP91_HUMAN
5536	WSGNEPTDEFSDLPLR2000000000000000	sp Q96B36 AKTS1_HUMAN
5537	SLSSSPQAQPPRPAELSDEEVAELFQR2.2E+24	sp Q96II8 LRCH3_HUMAN
5538	QEISRLQK200000	sp Q4V328 GRAP1_HUMAN
5539	RGSSGSVDETLFALPAASEPVIR2.002E+21	sp Q5TB80 CE162_HUMAN
5540	SRTPLISR202000000	sp Q9NQC3 RTN4_HUMAN
5541	APYQENDGYCPDLELSDSEAESDGNK202000000000	sp Q9UQ35 SRRM2_HUMAN
5542	GSRPPLILQSLSLPCSSPR200002000	sp Q92613 JADE3_HUMAN
		sp Q6KC79 NIPBL_HUMAN

5543	ASDPGLPAEEP200000000000	sp P27708 PYR1_HUMAN
5544	SGSETPDGLSPGK200000020000	sp Q8IWI9 MGAP_HUMAN
5545	SGSSQELDVKPSASPQER2000000000000000	sp Q9UQ35 SRRM2_HUMAN
5546	DGQDAIAQSPEK20000	sp Q9NWH9 SLTM_HUMAN
5547	TPTPSTPGDTQNPAPVPPAEDGIK2E+26	sp Q14839 CHD4_HUMAN
5548	SLSSSIGSNSTYLTSK2200000000000000	sp Q86SQ4 GP126_HUMAN
5549	SFNYSPTSSTSEVSSTSASK200000000000	sp Q5VT52 RPRD2_HUMAN
5550	YSPTSPTYSPTSPK2000000002000	sp P24928 RPB1_HUMAN
5551	RPTETNPVTSNSDEECNETVK200000020000000000	sp P46100 ATRX_HUMAN
5552	SASPYPHSLSSPQR2020000000000000	sp Q5T200 ZC3HD_HUMAN
5553	DAEPPSPTAGPPR2000000000	sp O43306 ADCY6_HUMAN
5554	SPSGPVKSPPLSPVGTTPVK2.0000002E+20	sp Q9BVC5 ASHWN_HUMAN
5555	GGPTSHSPVPPASSPSPLSGSALCGGK20200000000000	sp P49848 TAF6_HUMAN
5556	APSPVVEHPR200000000	sp Q86WR7 PRSR2_HUMAN
5557	ASSPSPLTIGTPESQR20000000000000	sp Q9NPI6 DCP1A_HUMAN
5558	LSSDLTR2000000	sp Q7Z6E9 RBBP6_HUMAN
5559	AHASLMSGR20000	sp Q14207 NPAT_HUMAN
5560	ILDNPTEGAPVLSPPADVSTFLAFPSPEK20000000000000000000	sp P46060 RAGP1_HUMAN
5561	ASVSQTSQASASSPR2000	sp Q8TDY2 RBCC1_HUMAN
5562	NPSSSTLPGGGVQNPADR20000000000000000000	sp Q9UJY4 GGA2_HUMAN
5563	TKPTQAAGPSSPKPPTPEETK20000000000000	sp P18887 XRCC1_HUMAN
5564	LSTSDPEEGDPLALGPESPGEPQPPQLK2.2E+26	sp Q5T1M5 FKB15_HUMAN
5565	ERPISLGFPLPAGDGLLTPDAQK2000000	sp O60271 JIP4_HUMAN
5566	IVVGSVAPGNTAPSPSSPTSPTSATTSMLENNPHAI2.2E+36	sp P49757 NUMB_HUMAN
5567	LSTTPSPTSSLHEDGVEDFR2.00000002E+19	sp O94842 TOX4_HUMAN
5568	ESSPEKEAEEGCEK20000000000000	sp O43719 HTSF1_HUMAN
5569	NSPTPATSTESLSPLHNVALR20020000000000000000	sp Q7Z6B7 SRGP1_HUMAN
5570	HNDIVSDSDAEDRGTLAELTAAHFGGGGGLLHK2.0000002E+27	sp P78316 NOP14_HUMAN
5571	GSSDGRGSDSEDLPHR22000000000000000000	sp Q9UPQ0 LIMC1_HUMAN
5572	SNTENLSQHFR2000000000	sp Q9UHB6 LIMA1_HUMAN
5573	GRSGSAAQAEGLCK20200000000000	sp Q92685 ALG3_HUMAN
5574	SEDGYHSDGDYGEHDYR20000020000000000000	sp P52756 RBM5_HUMAN
5575	KQSFDDNDSELEDK20000000	sp O60841 IF2P_HUMAN
5576	RQSSGSATNVASTPDNR200000000000000000	sp Q7Z460 CLAP1_HUMAN
5577	LGSSGLGSASSIQA AVR20020000000	sp Q7Z6Z7 HUWE1_HUMAN
5578	SPHQLLSPSSFSPSATPSQK2020000000000000	sp Q14181 DPOA2_HUMAN
5579	DLLSDLQDISDSER202000	sp P21127 CD11B_HUMAN
5580	SCPELTSGPR200000000000	sp Q9UQ88 CD11A_HUMAN
		sp Q13111 CAF1A_HUMAN
		sp P60709 ACTB_HUMAN
		sp P63261 ACTG_HUMAN
		sp P62736 ACTA_HUMAN
		sp P63267 ACTH_HUMAN
		sp P68032 ACTC_HUMAN
5581	SYELPDGQVITIGNER20000000000000000000	sp P68133 ACTS_HUMAN
		sp Q6S8J3 POTEE_HUMAN
		sp A5A3E0 POTEF_HUMAN
		sp Q9BYX7 ACTBM_HUMAN
		sp Q562R1 ACTBL_HUMAN
5582	TLLAMQAALLGSSSEEELESEN1.000000222E+19	sp P28715 ERCC5_HUMAN
5583	SLSMDSQVPVYSPSIDLK20000000000000000000	sp Q9ULJ3 ZBT21_HUMAN
5584	RNPSSSTLPGGGVQNPADR20000000000000000000	sp Q9UJY4 GGA2_HUMAN
5585	RPSSTSVPLGDK20000000	sp Q9NXR1 NDE1_HUMAN

5586	TDHGAEIVYKSPVVSGDTSR200020020000	sp P10636 TAU_HUMAN
		sp TAU_HUMAN
5587	LSQSSQDSSPVR200000000000	sp 095251 KAT7_HUMAN
5588	KNSTGSGHSAQELPTIR200000000000	sp 094763 RMP_HUMAN
5589	NMPGLSAATLASLGGTSSR100000000000000000	sp Q32MZ4 LRRF1_HUMAN
5590	GSEGYLAATYPTVGGTSPR2000	sp P21359 NF1_HUMAN
5591	FSEMMNMGDEDVDLPEVDGADDDSDSDEK1.1E+30	sp Q15185 TEBP_HUMAN
5592	SISSTVQR200000000	sp Q6PJF5 RHDF2_HUMAN
5593	STFSTNYR200000000	sp P05783 K1C18_HUMAN
5594	DDSDDESQSSHTGK200000000000	sp 094874 UFL1_HUMAN
5595	YFQSPSR20200	sp Q16629 SRSF7_HUMAN
5596	RQSPLPPQK20000000	sp Q01105 SET_HUMAN
5597	KPSPSESPKPFPAVSPEPR202000000000200000	sp Q96JM3 CHAP1_HUMAN
5598	IQSSLSVNSDISKK2002000000	sp Q9P1T7 MDFIC_HUMAN
5599	IDENSDKEMEVEESPEK20001000020000	sp Q8N3X1 FNBP4_HUMAN
5600	KRSPSPPTPEAK200000200000	sp Q8TAQ2 SMRC2_HUMAN
5601	EDKSPSEESAPTTSPESVSGVSSGSSGR2.02E+27	sp P54725 RD23A_HUMAN
5602	VPSPTPAK20000000	sp Q9UQ35 SRRM2_HUMAN
5603	ESPSPAPKPR2000000000	sp Q8IYB3 SRRM1_HUMAN
5604	KHSEEAFTPLKCSPK200000000002000	sp Q3B726 RPA43_HUMAN
5605	GQGESDPLDHEPAVSPLLPR2000000	sp P00519 ABL1_HUMAN
5606	SATPVNCEQPDILVSSTPINEGQTVLDK2.02E+28	sp P82094 TMF1_HUMAN
5607	IISPGSSTPSSSTR200000000000	sp Q5VT52 RPRD2_HUMAN
5608	VQIPVSRPDPVSDNEEDSYDEEIHDP2000000000000000	sp Q07157 ZO1_HUMAN
5609	TASSDSEEDPEALEK222020000000000	sp Q8N9T8 KRI1_HUMAN
5610	NLVSPAYCTQESR20000000200	sp Q9NUQ3 TXLNG_HUMAN
5611	SIDAVIATASAPPSSSPGR20000	sp Q9NR48 ASH1L_HUMAN
5612	AALGLQSDDEDAAVDIDEQIESMFNSK2E+21	sp Q7L014 DDX46_HUMAN
5613	VQRPPSAASAAPSSSK200000000000	sp Q5SRE5 NU188_HUMAN
5614	KSASSESEANLEAQPQSTVRPEEIPPIPENR2.202E+29	sp Q13427 PPIG_HUMAN
5615	SELSQDAEPAGSQETK200000	sp Q9BVJ6 UT14A_HUMAN
5616	KPIDSLRDSR2000200	sp Q9UQ35 SRRM2_HUMAN
5617	GRLTPSPDIIVLSDNEASSPR2020000002000000000	sp Q8WXI9 P66B_HUMAN
5618	ASPEPPGPESSSR2000000000000	sp Q9COC2 TB182_HUMAN
5619	KPGDGEVSPSTEDAPFQHSPLGK2000000000000000	sp 060318 GANP_HUMAN
5620	INHTVILDDPFDDPPDLLIPDRSPEPTR2000000	sp Q8WUA2 PPIL4_HUMAN
5621	DEEEEKEVENEDDDSDKEK200000	sp Q9P287 BCCIP_HUMAN
5622	HKSESPCESYPNEK2020000000000	sp Q8NI27 THOC2_HUMAN
5623	GRSPQPPAEEDDFDDTLVAIDTYNCDLHFK2E+30	sp Q9BUJ2 HNRL1_HUMAN
5624	SSATSGDIWPLSAYDNSR2000000000002000	sp Q9NYF8 BCLF1_HUMAN
5625	FDHESPGTDEDK2200200000	sp Q13427 PPIG_HUMAN
5626	RSSLSSHSHQSQIYR2202000000000	sp 015027 SC16A_HUMAN
5627	ARESSDSEEEYKK22020000000	sp Q9BYW2 SETD2_HUMAN
5628	EKSPVREPVDNLSPEER2000000000000000	sp Q86U06 RBM23_HUMAN
5629	TPHYGSQTPLHDGSR2000020000000000	sp 000267 SPT5H_HUMAN
5630	NTPSQHSHSIQHSPER20000000020000	sp Q9NYF8 BCLF1_HUMAN
5631	RLSSLRASTSK2200020000	sp P62753 RS6_HUMAN
5632	GSTLDLSDLEAEK200000000000	sp 000767 ACOD_HUMAN
5633	WSSGVGGSGGGSSGR2000000000	sp 095071 UBR5_HUMAN
5634	IFDEVMGCFSDSPQSPFPPEAGHTSLYDEDKVPR1.0000020002E+30	sp Q13085 ACACA_HUMAN
5635	SERGEVSGLMR200000000100	sp Q13263 TIF1B_HUMAN
5636	TESPPSPSSDVEDAR2000020020000000	sp Q9BRK4 LZTS2_HUMAN

		sp A5A3E0 POTEF_HUMAN
		sp Q6S8J3 POTEE_HUMAN
		sp P68133 ACTS_HUMAN
		sp Q562R1 ACTBL_HUMAN
5637	SYELPDGQVITIGNER2000000000000000	sp Q9BYX7 ACTBM_HUMAN
		sp P60709 ACTB_HUMAN
		sp P62736 ACTA_HUMAN
		sp P63261 ACTG_HUMAN
		sp P63267 ACTH_HUMAN
		sp P68032 ACTC_HUMAN
5638	TDSEKPFRRGSQSPK202000	sp Q9Y2W1 TR150_HUMAN
5639	SPQTLAPVGEDAMK200000000000100	sp Q92797 SYMPK_HUMAN
5640	GQSSPPPAPPICLR200000000000	sp O43379 WDR62_HUMAN
5641	LSGGLGAGSCR200000000000	sp Q04695 K1C17_HUMAN
5642	SISTSGPLDKEDTGR2000000000000000	sp Q7Z401 MYCPP_HUMAN
5643	SSDQDSTNKEAEAAGVKPAGVRPR2. 2E+24	sp Q92625 ANS1A_HUMAN
5644	VLSSTSEEDPGVVK2220000000000	sp Q8NI08 NCOA7_HUMAN
5645	ERSDSGGSSEPFDR2020000000000000	sp P48634 PRC2A_HUMAN
5646	TIASDSEEEAGKELSDKK2000020000000000000	sp Q96ST2 IWS1_HUMAN
5647	ESAAPASPAPSPAPSPTAPPQK2E+22	sp Q3KQU3 MA7D1_HUMAN
5648	TSYQPSSPGR20000	sp Q14134 TRI29_HUMAN
5649	ARSWSPPEVSR202000000000	sp Q9NR19 ACSA_HUMAN
5650	YSPTSPTYSPPTPK20000030002000	sp P24928 RPB1_HUMAN
5651	SPGSPPGPELPIETALDDR2E+19	sp Q8TEW0 PARD3_HUMAN
5652	TDDEDFVPSDASPPK20020000	sp P11388 TOP2A_HUMAN
5653	YPLCSADAAVSTTPPGSPPPPPLPEPPVLK2. 2E+20	sp Q9BTC0 DIDO1_HUMAN
5654	TPLSFTNPLHSDDSDSDERNSDGAVTQNK2. 00202E+19	sp Q05209 PTN12_HUMAN
5655	SVSQDLIK2000000000	sp Q9UPQ0 LIMC1_HUMAN
5656	AAESLTAIPEPASPQLLETPIHASQIQK2E+25	sp Q14676 MDC1_HUMAN
5657	SSPGQPEAGPEGAQERPSQAAPAVEAEGPGSSQAPR2E+35	sp P40222 TXLNA_HUMAN
5658	IYQYIQR300000	sp Q13627 DYR1A_HUMAN
		sp Q9Y463 DYR1B_HUMAN
5659	SKFDSDEEEEDTENVEAASSGK200000000000	sp Q8TF01 PNISR_HUMAN
5660	SSSVLSLEGSEK20020002000	sp A1L390 PKHG3_HUMAN
5661	TQSPGGCSAEAVLAR2000000000000000	sp Q96MH2 HEX12_HUMAN
5662	LPNNSRPSTPTINVLESK2000200000000000	sp Q15910 EZH2_HUMAN
5663	SLHSAHSLASR200000200200	sp O15027 SC16A_HUMAN
5664	TYSLTTPAR20000000	sp Q9Y4H2 IRS2_HUMAN
5665	ITPPAAKPGSPQAK20000000200000	sp P35658 NU214_HUMAN
5666	SHHAPMSPGSSGGGQPLAR2000000000000000	sp O14497 ARI1A_HUMAN
5667	SKSPSPR2020000	sp Q9UKV3 ACINU_HUMAN
5668	LSTTPSPTSSLHEDGVEDFR2. 20000022E+19	sp O94842 TOX4_HUMAN
5669	RADDFPVRDDPSDVTDEDEGPAEPPPPPK2002000000000000000	sp Q3YEC7 RABL6_HUMAN
5670	KEESEESDDMGFLFD200000010000000	sp P05387 RLA2_HUMAN
		sp P05386 RLA1_HUMAN
5671	RPPSPDIVLSDNEQPSSPR200000000000022000	sp Q86YP4 P66A_HUMAN
5672	YTPTSPSYSPSSPEYTPTSPK2E+20	sp P24928 RPB1_HUMAN
5673	THSTSSSLGSGESPF SR2220000000000000	sp Q9UGV2 NDRG3_HUMAN
5674	KSPSDTEGLVK200000000000	sp O95297 MPZL1_HUMAN
5675	FHSSDSEEEHKK220200000000	sp Q96ST2 IWS1_HUMAN
5676	FAELPEFRPEEVLPSPTLQSLATSPR2000000002000	sp Q96RK0 CIC_HUMAN
5677	GSNYHLSNDASDVE2000020000	sp Q96QE2 MYCT_HUMAN
5678	TPSPPPFR202000000	sp Q7L4I2 RSRC2_HUMAN
5679	RPSPQPSPR20002000	sp Q9UQ35 SRRM2_HUMAN

5680	ALFKPPEDSQDDESDSDAEEEEQTTKR2020000000000	sp Q13769 THOC5_HUMAN
5681	SLSSSPQAQPPRPAELSDDEEVAELFQR2E+24	sp Q4V328 GRAP1_HUMAN
5682	LSSSSPGATVSPSSSDAEYDKLPDADLAR2.02E+19	sp Q8NI08 NCOA7_HUMAN
5683	VQDTSNTGLGEDIHQLSK20000000000000	sp Q8IWA0 WDR75_HUMAN
5684	TVSPPIR200000	sp Q9UHB6 LIMA1_HUMAN
5685	NVQQDNSEAGTQPQVQTDAAQTSQSPPSPELTSEENK20000000000	sp Q92598 HS105_HUMAN
5686	GLPSPYNMSSAPGSR20000000000000	sp P15924 DESP_HUMAN
5687	IPCESPPLEVVDTTASTK20000000000000	sp P46013 KI67_HUMAN
5688	SGSPSDNSGAEEMEVSLAKPK2.020000200001E+21	sp P31749 AKT1_HUMAN
5689	YSPTSPTYSPPTPK300000030002000	sp P24928 RPB1_HUMAN
5690	ASSLNENVDHSAALLK20000000000000	sp Q96SB3 NEB2_HUMAN
5691	VEMYSGSDDDDDFNKLPK320000000000000	sp O60841 IF2P_HUMAN
5692	ALSSAVQASPTSPGGSPSSSGQR20200000000	sp Q03164 KMT2A_HUMAN
5693	HGSGADSDYENTQSGDPLLGLEGK20000000000000	sp Q9Y2X7 GIT1_HUMAN
5694	SVSEHTSCPAESASDAAPLQR200000000000000	sp Q8WUA7 TB22A_HUMAN
5695	TSSEMSWVDLNIPTPEAPPCYMDVIPEDHR2.002000000002E+29	sp Q9H3M7 TXNIP_HUMAN
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5697	STLESEKPGSPEAAETSPPSNIIDHCEK22000000000000	sp Q96T23 RSF1_HUMAN
5698	KNSSTDQGSDEEGLQK2000020000000000	sp Q12789 TF3C1_HUMAN
5699	NPPPQDYESDDDSYEVLDLTEYAR2000000000000000	sp Q8IVP5 FUND1_HUMAN
5700	VDPSLMEDSDDGPSLPTK10020000000000	sp O15258 RER1_HUMAN
5701	GTEPSPGGTPQSRPVPSPAGPPEGVPEEAQPPR2.0000002E+32	sp Q8WUZ0 BCL7C_HUMAN
5702	AKKPEEESSESEEGSESEEEAPAGTR2.022000202E+19	sp Q13428 TCOF_HUMAN
5703	GHTDTEGRPPSPPTSTPEK20000000000	sp Q00613 HSF1_HUMAN
5704	GQASSPTPEPGVGAGDLPGPTSAPVPSGSGGR2E+31	sp Q14676 MDC1_HUMAN
5705	SSSPILTKPTLK22000000000000	sp Q92794 KAT6A_HUMAN
5706	SRQTPSPDVVLR2020000000	sp Q9UPQ0 LIMC1_HUMAN
5707	DHNVGSPPKSPHASPQHTQTEK200000000000000	sp A0JNW5 UH1BL_HUMAN
5708	ESPFSTSASPLLSGSQHFDVPPR2000000000000000	sp O14980 XPO1_HUMAN
5709	KPSPAASPAK200000	sp Q8N3F8 MILK1_HUMAN
5710	GESAPTLSTSPSPSSPSTSPSPTLGR20200000000000	sp Q9H330 TM245_HUMAN
5711	SPGLCSDSLEK200000000000	sp Q5VUA4 ZN318_HUMAN
5712	SLYASSPGGVYATR2000000000000000	sp P08670 VIME_HUMAN
5713	SDSASSEPVGIYQGFQEK2000000003000000	sp Q05655 KPCD_HUMAN
5714	ALQAPHSPSK200	sp Q96AQ6 PBIP1_HUMAN
5715	DLQSSPPLPLGFLGEGDAVTK2000000000000000	sp Q9UN79 SOX13_HUMAN
5716	ACASPSAQVEGSPVAGSDGSQPAVK2E+20	sp Q9UFC0 LRWD1_HUMAN
5717	SESPKEPEQLR2000000000	sp P09651 ROA1_HUMAN
5718	NSSLGSPSNLCGSPGSI200000000020000000	sp Q9C0B0 UNK_HUMAN
5719	SPGSGSQSSGWHEVEPGMPSPTTLK2.002E+25	sp Q12857 NFIA_HUMAN
5720	SRTPLLPR202000000	sp Q9UQ35 SRRM2_HUMAN
5721	SLMASLTNMSR200	sp Q9ULS6 KCNS2_HUMAN
5722	RPSESDKEDELK202000000000	sp Q02952 AKA12_HUMAN
5723	EAEEGPTGASESQDSPR2000	sp Q5TCZ1 SPD2A_HUMAN
5724	EVSPDPSTGAASPASSAGGSAAR200020000000000000	sp P23769 GATA2_HUMAN
5725	VAQGATEKSPEDK200000	sp P49321 NASP_HUMAN
5726	YSDTTDSDPENEPFDEDQHTQITK3.002202E+24	sp P60484 PTEN_HUMAN
5727	GTEPSPGGTPQSRPVPSPAGPPEGVPEEAQPPR2.000000000002E+29	sp Q8WUZ0 BCL7C_HUMAN
5728	EMLMEDVGSEEEQEEDEAPFQEK1.0000002E+23	sp Q9H1E3 NUCKS_HUMAN
5729	LLKPGEEPSEYTDDEEDTKDHNK200000000000	sp O15173 PGRC2_HUMAN
5730	YHGHSMSDPGVSYR21000002000	sp P29803 ODPAT_HUMAN
5731	KEATPRSR200	sp P08559 ODPA_HUMAN
5732	RQSSGSATNVASTPDNR20000000000000	sp Q99572 P2RX7_HUMAN
		sp Q7Z460 CLAP1_HUMAN



5733	GTSDSSSGNVSEGESPPDSQEDSFQGR20000000000000	sp Q8WUB8 PHF10_HUMAN
5734	LCSSDTLVSEGEENQKPK200000000000000000	sp Q5TOW9 FA83B_HUMAN
5735	INPPSSGGTSSSPIK20000	sp P14859 PO2F1_HUMAN
5736	AMSTTSSPQPGK10000020000000	sp Q9UJU6 DBNL_HUMAN
5737	QEAGDSPPPAPGTPK20000000000	sp P49848 TAF6_HUMAN
5738	SCMLTGTPEVQSAK2000000000	sp Q96AE4 FUBP1_HUMAN
5739	SLGSASPGPGQPPLSSPTR200000000020000	sp Q9C0B5 ZDHC5_HUMAN
5740	SSEPVKETVQTTQSPTPVEK20000000	sp Q9UPN6 SCAF8_HUMAN
5741	NVAEALGHSPK2000	sp Q8IU81 I2BP1_HUMAN
5742	SRSPQAFR202000000	sp Q7Z6E9 RBBP6_HUMAN
5743	SLENTPPFPTPK2000000002000	sp O15085 ARHGB_HUMAN
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5745	LDLTENLTGSK200	sp P46013 KI67_HUMAN
5746	VTLYNTDQDGS DSPR200000	sp Q9UPW0 FOXJ3_HUMAN
5747	ALSSDSILSPAPDAR20000000000	sp Q8IVT2 MISP_HUMAN
5748	RPAEATSSPTSPERPR20000000000	sp Q9BU76 MTA2_HUMAN
5749	AGSPSPQPSGELPR2020000000000	sp Q92619 HMHA1_HUMAN
5750	ALSSLHGDDQDSEDEVLTIPEVK2.00000002E+20	sp P11717 MPRI_HUMAN
5751	TGVPSTASVGK20000	sp P28290 SSFA2_HUMAN
5752	REEQTDSDGESVTHHIR20000000	sp P36915 GNL1_HUMAN
5753	GPTTGEGALDLSVHSPPKSPEGK2000200000	sp O95684 FR10P_HUMAN
5754	LEISPDSSPER200020000	sp Q99959 PKP2_HUMAN
5755	ESAAPASPAPSPAPSPAPPQK200020000000000000	sp Q3KQU3 MA7D1_HUMAN
5756	RLSSSLNPSK200000000	sp Q13535 ATR_HUMAN
5757	WDSYENLSADGEVLHTQGPVDGSLYAK2E+25	sp Q68CZ2 TENS3_HUMAN
5758	GSPLAEGPQAFFPER2000000000000000	sp Q8WUF5 IASPP_HUMAN
5759	KSSSEAKPTSLGLAGGHK220000000000000000	sp Q96RK0 CIC_HUMAN
5760	QRSPSPAPAPAPAAAAGPPTRK2.02E+20	sp Q9H7N4 SFR19_HUMAN
5761	SSSQPSSCCSDPSKPGGNVEGATQSLAEQMR2.002E+29	sp Q13501 SQSTM_HUMAN
		sp P30510 1C14_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
5762	GGSCSQAASSNSAQGSDES LIACK2000000000	sp Q07000 1C15_HUMAN
		sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
5763	APSSDEECFFDLLTK20000000000000	sp Q86YR5 GPSM1_HUMAN
5764	RHSHSHSPMSTR20002002000	sp P62995 TRA2B_HUMAN
5765	RTDAL TSSPGR2220000	sp P49736 MCM2_HUMAN
5766	ASPGHSPHYFAASSPTSPNALPPAR2.003E+20	sp Q8TEM1 PO210_HUMAN
5767	ETINNLDTSSSDFTILQEIEEPSLEPENEK200000000000000000	sp Q15424 SAFB1_HUMAN
5768	VASEAPLEHKPQVEASSPR20000	sp Q8TD19 NEK9_HUMAN
5769	DSGSDTASAIIPSTTPSVDSDES VVK20000	sp Q6WKZ4 RFIP1_HUMAN
5770	HKSGSMEEDVDTSPGGDYTSPSSPTSSSR2.001E+28	sp P08651 NFIC_HUMAN
5771	RSPQQTVPYVVPLSPK2000000000002000	sp Q9UQR0 SCML2_HUMAN
5772	KPPAPSPVQSQSPSTNWS PAVPVK2.000200002002E+19	sp Q8IYB3 SRRM1_HUMAN
5773	AGDRNSEDGVMVTFSSVK200000010000000	sp Q9NY61 AATF_HUMAN
5774	EVDGLLTSEPMGSPVSSK2000000	sp Q8WWM7 ATX2L_HUMAN
5775	HFSEHPSTSK2200	sp O43719 HTSF1_HUMAN
5776	DTSATSQSVNGSPQAEQPSLESTSK2E+20	sp Q86WB0 NIPA_HUMAN

5777	LGAGGGSPEKSPSAQELK200000000000	sp Q9UNE7 CHIP_HUMAN
5778	TDCEPVGLEPAVEQSSAASELAEASSEELAEAPTEAPSPEAR220000000200 0000000	sp Q15424 SAFB1_HUMAN sp Q14151 SAFB2_HUMAN
5779	SVSPPPK20200000	sp Q13247 SRSF6_HUMAN
5780	RLSSTSLASGHSVR202000000000	sp Q9BZL6 KPCD2_HUMAN
5781	KKPSTDEQTSSAEEDVPTCGYLNVLNSNR2.00022E+25	sp Q8N556 AFAP1_HUMAN
5782	MSGGSTMSSGGGNTNNSNSK2000000000000000	sp Q86U70 LDB1_HUMAN
5783	TLNAETPKSSPLPAK20002000000	sp Q7L2J0 MEPCE_HUMAN
5784	RPLRPQVVTDDDGQAPEAK200000000000	sp Q9HDC9 APMAP_HUMAN
5785	YFDSGDYNMAK200000000	sp 043768 ENSA_HUMAN sp P56211 ARP19_HUMAN
5786	EDKSPSEESAPTTSPESVSGVSPSSGSSGR2.2E+19	sp P54725 RD23A_HUMAN
5787	SSPSILAVQR2000000000	sp 014646 CHD1_HUMAN
5788	EPLENSASESSDTELPEK202200000000	sp P78337 PITX1_HUMAN
5789	TLENSPISSCDTSDEGVPVNSAAVLK2.002E+24	sp Q8N556 AFAP1_HUMAN
5790	ESEQESEEEILAQK20000000000000	sp Q9BW71 HIRP3_HUMAN
5791	SFYSSHYAR2000000000	sp Q8NEY8 PPHLN_HUMAN
5792	CDILVQEELLASPK2000	sp Q96JM3 CHAP1_HUMAN
5793	SLSYSPVER20200000	sp Q9UQ35 SRRM2_HUMAN
5794	STSPAGQHHSPISSR2200000000000000	sp Q5T200 ZC3HD_HUMAN
5795	GTEPSPGGTPQSRPVSPAGPPEGVPEEAQPPR2.0002E+21	sp Q8WUZ0 BCL7C_HUMAN
5796	SRVSVSPGR2020000	sp Q8IYB3 SRRM1_HUMAN
5797	SLSVEHLETK20200000000	sp P26045 PTN3_HUMAN
5798	FQEQECPPSPEPTRK20000000	sp P62070 RRAS2_HUMAN
5799	SYSPAHR202000000	sp Q9Y2W1 TR150_HUMAN
5800	GSSPLGPAGLGAEEPAAGPQLPSWLQPER2E+27	sp 043482 MS18B_HUMAN
5801	ISDSESEDPPR20000000000	sp Q96ST2 IWS1_HUMAN
5802	SGSSPSQSPCDLEEPWLPR2.0200000002E+20	sp Q5SYE7 NHSL1_HUMAN
5803	AGPTLTDENGDDLGLPPSPGDTSYQQDQVDDFHEAR3000000000000	sp Q9NQZ2 SAS10_HUMAN
5804	SVSNGTAKPATSENFDL2.02E+20	sp Q96K49 TM87B_HUMAN
5805	ASAGHVAVSSPTPETGPLQR200000000000	sp P42695 CNDD3_HUMAN
5806	TAHNSEADLEESFNEHELEPSSPK2E+20	sp Q8IWS0 PHF6_HUMAN
5807	LCSSSDTLVSEGEENQKPK20000000000000	sp Q5T0W9 FA83B_HUMAN
5808	SPSTTYLHTPTSEDAAIPSK2E+19	sp Q13111 CAF1A_HUMAN
5809	SPQTLAPVGEDAMKTPSPAEDAR20200000000	sp Q92797 SYMPK_HUMAN
5810	LAELPAAQPSAEDSDTEDDSEAEQTER2000002000200000000	sp 095714 HERC2_HUMAN
5811	NIPSPGQLDPDTR20000000000	sp Q9P2D1 CHD7_HUMAN
5812	SAQGSSSPVPSMVQK2000010000	sp P35658 NU214_HUMAN
5813	HTGMASIDSSAPETTSDSPTLSR1E+21	sp A7E2V4 ZSWM8_HUMAN
5814	YSPCEEHPAETNQNEGSESGTIR2E+22	sp Q12774 ARHG5_HUMAN
5815	SAASPVVSSMPER20000010000	sp P18583 SON_HUMAN
5816	FGSTGSTPPVSPTPSER200020000000	sp Q969V6 MKL1_HUMAN
5817	TPSSSPPITPPASETK22000000000000	sp Q9NZJ0 DTL_HUMAN
5818	AAPEASSPPASPLQHLLPGK2000000000000000	sp Q96TA1 NIBL1_HUMAN
5819	KGPGQPSSPQR20000	sp 095400 CD2B2_HUMAN
5820	SSATSGDIWPGLSAYDNSPR2E+19	sp Q9NYF8 BCLF1_HUMAN
5821	TASPPGPPYGK20000000000	sp P54259 ATN1_HUMAN
5822	VVSAPVGKETPSK20000	sp Q13416 ORC2_HUMAN
5823	RPASQKDSGAASEQATAAPNPCSSSR2000000000000000	sp Q9BU23 LMF2_HUMAN
5824	DNQESSDAELSSSEYIK2000022000000	sp Q96JM3 CHAP1_HUMAN
5825	TFSLTEVR2000000	sp 095239 KIF4A_HUMAN
5826	TGVTSTSDSEEGDDQEGEKK2020000000000000	sp 075475 PSIP1_HUMAN
5827	HSSTGDSADAGPPAAGSAR2000000000000000	sp Q6ZU35 K1211_HUMAN
5828	EVPPPPAESEEEEDDGLPK200000000000	sp Q9H6X2 ANTR1_HUMAN



		sp P05534 1A24_HUMAN
		sp Q09160 1A80_HUMAN
5873	KGGSYSQAASSDSAQGSVSLTACK20000000000000	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
5874	AAPPPPALTPDSQTVDSCKTPEVSFLPEEATEEAGVR2.002E+21	sp Q9H7N4 SFR19_HUMAN
5875	GPKPEPPGSGSPAPPR200000000	sp Q9UKJ3 GPTC8_HUMAN
5876	RDSFDDR200000	sp P43243 MATR3_HUMAN
5877	KRSEDESETEDEEEK20002020000000	sp P25205 MCM3_HUMAN
5878	VSCHDSDDDIMR20000000	sp Q76FK4 NOL8_HUMAN
5879	SMSENAVR200000000	sp Q9GZY8 MFF_HUMAN
5880	SDGACDSPSSDKENSSQIAQDHQK2000000000000000	sp Q9UIG0 BAZ1B_HUMAN
5881	GILSLPHQASVSR200000	sp O75925 PIAS1_HUMAN
5882	LQQGAGLESPQGQPEPGAASPQR2000000000000000	sp Q9H6F5 CCD86_HUMAN
5883	DKRPLSGPDVGTQPAGLASGAK200000000000000000	sp O96013 PAK4_HUMAN
5884	GSSDGRGSDSEDLPHR200000202000000000	sp Q9UPQ0 LIMC1_HUMAN
5885	EISSPESCPPEIK22000200000000	sp Q92889 XPF_HUMAN
5886	SVASSQPAKPTK20000000000000	sp Q07157 ZO1_HUMAN
5887	LGASNSPGQPNSVK200000000000	sp P49756 RBM25_HUMAN
5888	ISYIPDEEVSSPSPQR200000	sp Q9Y6M1 IF2B2_HUMAN
5889	EGSPAPLEPEPGAAPK2000000000000000	sp Q01167 FOXK2_HUMAN
5890	AGMSSNQSISSPVLDAVPR122002002000000000	sp Q9UQ35 SRRM2_HUMAN
5891	KPEKPLFSSASPQDSSPR2000	sp O00712 NF1B_HUMAN
5892	LDSSPVSSPR20002000	sp O60293 ZC3H1_HUMAN
5893	TSAQHALTSVSRGSSLK22000	sp P61073 CXCR4_HUMAN
5894	ASQEPSKPGTEVIPAAPR2000000000000000	sp Q96CP2 FWCH2_HUMAN
5895	TSPPLDR20000000	sp Q9UQ35 SRRM2_HUMAN
5896	SSSEYTSFQSR20000000000000	sp Q9BTU6 P4K2A_HUMAN
5897	SRSPIIHSFK20000002000	sp O94913 PCF11_HUMAN
		sp Q15836 VAMP3_HUMAN
5898	DQKLELDDR2000000	sp P23763 VAMP1_HUMAN
		sp P63027 VAMP2_HUMAN
5899	SLTNSHLEKK20002000000	sp Q9H2H9 S38A1_HUMAN
5900	SPSVSPEPAEK2000220000000	sp Q9UQ35 SRRM2_HUMAN
5901	VVPEDGSPEKR200000	sp Q02040 AK17A_HUMAN
5902	NSPVTKTPPR2000020000	sp Q13541 4EBP1_HUMAN
5903	LCDDGPQLPTSPR20000	sp O60504 VINEX_HUMAN
5904	HRGSEEDPLLSPVETWK200000020000000	sp Q13017 RHG05_HUMAN
5905	LSPERSDSESNENEPHLPHGVSQR202000000000000000	sp Q9BQ52 RNZ2_HUMAN
5906	SSATSGDIWPGLSAYDNSPR200000000003000000	sp Q9NYF8 BCLF1_HUMAN
5907	SKPNLPSESR200	sp Q08170 SRSF4_HUMAN
5908	TCNSPQNSTDSVSDIVPDSPPFGALGSDTR2E+27	sp Q14694 UBP10_HUMAN
5909	ELSNSPLRENSFGSPLEFR200000002002000000	sp Q9UQ35 SRRM2_HUMAN
5910	HASSPESPKPAPAPGSHR20000000000000	sp Q9UQ35 SRRM2_HUMAN
5911	LGGAVPFAPPEVSPEQAK2000000	sp Q6ZSR9 YJ005_HUMAN
5912	SPHSPSIATPPSQSSSCDAPMLSTVHSAK2.002020002E+31	sp Q9NYF8 BCLF1_HUMAN
5913	AGAPVGGSPGLAK2000000	sp Q6ZRS2 SRCAP_HUMAN
5914	MVIQGPSSPQGEAMVTDVLEDQK1.0000020000001E+23	sp Q12888 TP53B_HUMAN
5915	IEEVLSPGSPSKSPSK2000200020000	sp Q9UEY8 ADDG_HUMAN
5916	SSSPGKQAVSSLNSSHRSR20000000000000000000	sp Q9UHB7 AFF4_HUMAN
5917	YETPSDAIEVISPASSAPPQEK2000200000000	sp O75376 NCOR1_HUMAN
5918	DFDIAEQNESSDEESLRK2200000000	sp Q96KC8 DNJC1_HUMAN
5919	GPAFNMASPESDFGINLK10200000000000	sp Q09666 AHNK_HUMAN

5920	INPPSSGGTSSSPIK20000000	sp P14859 PO2F1_HUMAN
5921	GSRPPLILQSQLPCSSPR2000	sp Q6KC79 NIPBL_HUMAN
5922	LSLSPISLAESWEDGSPPPQAR2000020000000	sp A1L170 CA226_HUMAN
5923	ELEENDSENSEFEDDGSEKVLDEEGSER2.0020000002E+22	sp 043719 HTSF1_HUMAN
5924	DVLSLLNLSR2000	sp Q96AY4 TTC28_HUMAN
5925	ALVVPEPEPDSNSQER20200000	sp Q5VTR2 BRE1A_HUMAN
5926	RNSTTIMSR20000000	sp Q7Z309 F122B_HUMAN
5927	SATPVNCEQPDILVSSTPINEGQTVLTK2E+28	sp P82094 TMF1_HUMAN
5928	SGPKPFSAPKPKTSPSPK2002000	sp Q01518 CAP1_HUMAN
5929	EEAPASPLRPLYPQISPLK200000000020000	sp P85037 FOXK1_HUMAN
5930	SESPKEPEQLR200000000000	sp P09651 ROA1_HUMAN
5931	NSLPASPAHQLSSSPR200000002000	sp Q8TEK3 DOT1L_HUMAN
5932	SGTNSPPPPFSDWGR200000000000	sp Q9UKT5 FBX4_HUMAN
		sp Q09160 1A80_HUMAN
		sp P05534 1A24_HUMAN
5933	GGSYSQAASSDSAQGSVDVSLTACK20002000000000	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
		sp P24941 CDK2_HUMAN
5934	IGEGTYGVVYK23000000	sp P06493 CDK1_HUMAN
		sp Q00526 CDK3_HUMAN
5935	VNSNGKESPGSSEFFQEAVSHGK22000000000000	sp Q8N108 MIER1_HUMAN
5936	TQLWASEPGTPLPTSLPSQNPILK2000000000000000	sp Q9BXP5 SRRT_HUMAN
5937	RPSQPSPR20000000	sp Q9UQ35 SRRM2_HUMAN
5938	SSGDPEQIKEDSLSEESADAR2.000000000202E+20	sp Q9BVS4 R1OK2_HUMAN
5939	EDEISPPPPNPVVK200000000000	sp P10644 KAPO_HUMAN
		sp Q13247 SRSF6_HUMAN
5940	LIVENLSSR200	sp Q08170 SRSF4_HUMAN
		sp Q13243 SRSF5_HUMAN
5941	ITEVSCKSPQESFK200000000	sp P46013 KI67_HUMAN
5942	IPSEMPSPNFGK20002000000	sp Q9H2J7 S6A15_HUMAN
5943	ALSSGGSITSPPLSPALPK200000000002000000	sp Q6Y7W6 PERQ2_HUMAN
5944	GEGDAPFSEPGTTSTQRPSSPETATK20000000	sp Q9UQ35 SRRM2_HUMAN
5945	DSPGIPPSANAHQLFR2000002000000000	sp P51812 KS6A3_HUMAN
5946	RSPSPAPPPR2020000000	sp Q8IYB3 SRRM1_HUMAN
5947	SSPQLDPLR200000000	sp Q8ND56 LS14A_HUMAN
5948	VQAGPGSPR2000	sp Q6PK04 CC137_HUMAN
5949	GGAPDPSPGATATPGAPAQPSSPDAR2000000000200000	sp 095365 ZBT7A_HUMAN
5950	RLSSASTGKPLSVEDDFEK200200000000000000	sp 075152 ZC11A_HUMAN
5951	AMSTTSISSPQPGK2000000	sp Q9UJU6 DBNL_HUMAN
		sp P42166 LAP2A_HUMAN
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		sp Q86WB0 NIPA_HUMAN
5953	DTSATSQSVNGSPQAEQPSLESTSK200020000000000000	sp Q8WWM7 ATX2L_HUMAN
5954	STSTPTSPGPR200000	sp Q8IYB3 SRRM1_HUMAN
5955	KETESEAEDNLDDLEK2000000000000000	sp Q9H6F5 CCD86_HUMAN
5956	ALVEFESNPEETREPGSPSVQR20000000000000	sp Q9Y3T9 NOC2L_HUMAN
5957	SPDKPGGSPSASR20000002000000	sp P21796 VDAC1_HUMAN
5958	LTFDSSFSPNTGK200000000	sp Q2PPJ7 RGPA2_HUMAN
5959	RSSSPAELDLKDDLQQTQGK220000000000000000	sp Q02241 KIF23_HUMAN
5960	RSSTVAPAQPDGAESEWTDVETR2E+20	sp Q9Y2U5 M3K2_HUMAN
5961	RGSDIDNPTLTVMDISPPSR2000000000000200000	sp Q9H4L7 SMRCD_HUMAN
5962	DTVIIIVSEPEDEESQGLPTMAR200000002000001000	

5963	GPDPGEDLADLEVVVPGSPGDCLPEEADGTDTHLGLPAGASQPALTSSR2E+32	sp 060343	TBCD4_HUMAN
5964	GGGTPDANSLAPPK200000000000	sp Q9BRQ0	PYG02_HUMAN
5965	TEPAAEAEAAASGPSESPSPAAEELPGSHAEPVPAQGEAPGEQAR2E+31	sp P55884	EIF3B_HUMAN
5966	SDGEAKPEPSPSR20000000000000	sp P26358	DNMT1_HUMAN
5967	RPNKQEESESPVERPLK20200000000	sp Q96SB4	SRPK1_HUMAN
5968	LSSLPLR20000	sp Q66K74	MAP1S_HUMAN
		sp P49792	RBP2_HUMAN
		sp Q7Z3J3	RGPD4_HUMAN
		sp O14715	RGPD8_HUMAN
5969	DSLITPHVSR200	sp Q99666	RGPD5_HUMAN
		sp A6NKT7	RGPD3_HUMAN
		sp PODJD0	RGPD1_HUMAN
		sp PODJD1	RGPD2_HUMAN
5970	AGGPTTPLSPTR20000000	sp P20700	LMNB1_HUMAN
5971	RVISDESSEDIGSDVEFKPDK2.020200002E+19	sp P52701	MSH6_HUMAN
5972	TENSLANENQQPIKSEPESEGEPEK2E+22	sp Q8NHM5	KDM2B_HUMAN
5973	NRENSPSSQSAGLSSINK2000000000000000	sp Q9H2Y7	ZN106_HUMAN
5974	SETPPHWR202000000	sp Q13427	PIIG_HUMAN
5975	EVHDELEDLPSPPPPLSPPTTSPHK20000020000200000	sp Q9NQX3	GEPH_HUMAN
5976	SHMSGSPGPGSNTAPSTPVIGGSDKPGMEK2.02E+29	sp A0FGR8	ESYT2_HUMAN
5977	SSATSGDIWPGLSAYDNSPR200000000000002000	sp Q9NYF8	BCLF1_HUMAN
5978	RVSPLNLSSVTP20000000200	sp Q9UJX2	CDC23_HUMAN
5979	APSDSSLGTPSDGRPELR2000000000000000	sp Q15642	CIP4_HUMAN
5980	TPTPSTPGDTQPNTAPVPPAEDGIK2.02E+24	sp Q14839	CHD4_HUMAN
5981	RADLNQGIGEPQSPSR20000	sp Q96C19	EFHD2_HUMAN
5982	ALSSGGSITSPPLSPALPK200000200000000000	sp Q6Y7W6	PERQ2_HUMAN
5983	ELSNSPLRENSFGSPLEFR202000000002000000	sp Q9UQ35	SRRM2_HUMAN
5984	KNSTDLDLSDAPEDPTSPK2200000000002000	sp Q96RK0	CIC_HUMAN
5985	LQLGTPPPLAAR2000000000	sp Q9P2N5	RBM27_HUMAN
5986	SASLSHPGGEGERPAR2000000000000000	sp Q2M3G4	SHRM1_HUMAN
		sp P01891	1A68_HUMAN
		sp P10316	1A69_HUMAN
5987	KGGSYSQAASSDSAQGSVSLTACK20000000000000	sp P05534	1A24_HUMAN
		sp P30447	1A23_HUMAN
		sp Q09160	1A80_HUMAN
		sp P01892	1A02_HUMAN
5988	LDSEMMDHSENYDTMSSPLPGK2000030100000000	sp Q9NTI5	PDS5B_HUMAN
5989	GDQPAASGSDDDDEPPPLPR2000000000000000	sp O00264	PGR1_HUMAN
5990	STSPIIGSPPVR220000000000	sp Q86TB9	PATL1_HUMAN
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5992	GAAAAATGQPGTAPAGTPGAPPLAGMAIVK2000000000000000	sp Q13263	TIF1B_HUMAN
5993	YRPASVPPSPSLSR20002000000	sp P13807	GYS1_HUMAN
5994	KFSAGGSDPPLKR20000000000000	sp O75152	ZC11A_HUMAN
5995	SLSSGESGPGSPTHSHLSR20000000000000000000	sp Q6P0Q8	MAST2_HUMAN
5996	ERSPSPLR2020000	sp Q9Y2V2	CHSP1_HUMAN
5997	LFEDDSDNEKLFDEEEDSSEKLFDDSDER2.00000000022E+23	sp O43719	HTSF1_HUMAN
5998	NRPPLPAGTNSK200	sp P42167	LAP2B_HUMAN
		sp P42166	LAP2A_HUMAN
5999	QRGSETGSETHESDLAPSDK2000200000000000	sp P48634	PRC2A_HUMAN
6000	SLSPGVS2000000	sp Q9UKV3	ACINU_HUMAN
6001	SLPITIEMLK20000000000	sp Q8IZ21	PHAR4_HUMAN
6002	FASENDLPEWK2000000000	sp P43487	RANG_HUMAN
6003	SPQISMSDIDLNLK2000000000000000	sp Q09666	AHMK_HUMAN

6004	KTSSDDESEEDDLLQR220000000000000000	sp Q9Y5J1 UTP18_HUMAN
6005	GQKPALDCSQVSPRPATSPENNASLSDTSPMDSSPSGIPK2.002E+33	sp P18858 DNLI1_HUMAN
6006	QRGSETGSETHESDLAPSDK20200000000000	sp P48634 PRC2A_HUMAN
6007	RPMEEDGEEKSPSKK200000	sp Q12906 ILF3_HUMAN
6008	IPATEQTNQVIEKASDSEEPEEK2000000000	sp Q9C0E8 LNP_HUMAN
6009	TAVQYIESSDSEEIETSELPQK2200000000000000	sp Q14527 HLTF_HUMAN
6010	INPDNYGMDLNSDDSTDDEAHPR10002002200000000	sp Q9NQS7 INCE_HUMAN
6011	GPSQATSPIR20000	sp Q7RTP6 MICA3_HUMAN
6012	VQMTSPSSTGSPMFK10200000001000	sp P49790 NU153_HUMAN
6013	RLSSASTGKPPLSVEDDFEK2200200000000000000	sp O75152 ZC11A_HUMAN
6014	IAASSFR2000	sp Q99961 SH3G1_HUMAN
6015	TSSVSSISQVSPER20000	sp P42356 PI4KA_HUMAN
6016	DTVIIVSEPEDEESQGLPTMAR200200002000001000	sp Q9H4L7 SMRCD_HUMAN
6017	FHSPSTTWSFNK20000020000	sp Q99590 SCAFB_HUMAN
6018	AAHVPENSDEQDVLTVKPV2000000000000000	sp Q9H2Y7 ZN106_HUMAN
6019	SVASNQSEMEFSSLQDMPK20020000000001000	sp Q8N163 CCAR2_HUMAN
6020	IAAPELHKGSDSEDEPTKK202000000000	sp Q92733 PRCC_HUMAN
6021	SDELLGSDSHDGESESNAK2002000000000000	sp P38398 BRCA1_HUMAN
6022	VGSGQGGSPCWK200000	sp Q9UMN6 KMT2B_HUMAN
6023	KPESPYGNLCDAPDSRPVK200000000002000000	sp Q9UKA4 AKA11_HUMAN
6024	LQELESCSGLGSTDDTDVR200000000	sp Q14C86 GAPD1_HUMAN
6025	SSISSISGR200000000	sp Q86SQ0 PHLB2_HUMAN
6026	ESEGEDEDEDEDLSK2000000000000000	sp O95218 ZRAB2_HUMAN
6027	ESAAPASPAPSPAPSPPTAPPQK2.0000000020002E+22	sp Q3KQU3 MA7D1_HUMAN
6028	SLNFQEDDDTSR200	sp Q9H814 PHAX_HUMAN
6029	VESEESGDEEGKKHSSGIVADLSEQSLK2.002E+26	sp Q15435 PP1R7_HUMAN
6030	APAGQEEPTGTPSSPLSAEQDR220000000000	sp P13051 UNG_HUMAN
6031	NSLLAGDDDTMSVISGISSR2000200	sp Q8N3U4 STAG2_HUMAN
6032	GLNTSQESDDDILDESSPEGTQK2.2E+21	sp Q5VV41 ARHGG_HUMAN
6033	ASPPGDLQNP20000000000	sp O00273 DFFA_HUMAN
6034	KPSPSEPEPWKPFPAVSPEPR2000000000000200000	sp Q96JM3 CHAP1_HUMAN
6035	NSNFSVQHPSSTSPTEK200000	sp Q03468 ERCC6_HUMAN
6036	GLMAGGRPEGQYSEDEDTDDEYK1.000000003E+22	sp Q9NPQ8 RIC8A_HUMAN
6037	SIGASPNPFSVHTATAVPSGK2.0002E+21	sp P09884 DPOLA_HUMAN
6038	SYSPDGKESPSDKK230000002000000	sp P43243 MATR3_HUMAN
6039	SLTRSPPAIR202000000	sp Q9UQ35 SRRM2_HUMAN
6040	LRLSPSPTSQR2000000	sp P02545 LMNA_HUMAN
6041	VSPSTSYTPSR20000	sp Q9BTE3 MCMBP_HUMAN
6042	VSITSEVESTSNPSSSSLQK22000000000000	sp Q04656 ATP7A_HUMAN
6043	QGSITSPQANEQSVTPQRR202000000000200000	sp Q9UQ35 SRRM2_HUMAN
6044	DGDSYDPYDFSDTEEEMPQVHTPK300002000100000000	sp P25205 MCM3_HUMAN
6045	GEDSAEETEAKPAVVAPAVVEAVSTPSAAFPDATAEQGPILTK2E+42	sp Q12906 ILF3_HUMAN
6046	MSQPGSPSPK200000	sp Q14839 CHD4_HUMAN
6047	SDATNYASHSPVNR2000000	sp Q96FJ0 STALP_HUMAN
6048	QVPVMQGTGMQGASIQGGSQPGGFSPGQNVTPQDHEK2E+25	sp P33240 CSTF2_HUMAN
6049	GGSPQMDDIK200000000	sp Q13144 E12BE_HUMAN
6050	HHYDSDEKSETR200020000	sp Q9BQ39 DDX50_HUMAN
6051	SLPTTVPESPNYR200000	sp Q6PKG0 LARP1_HUMAN
6052	IQQEESGGEEDSDLSPEER2200002000000000	sp Q6NXS1 IPP2M_HUMAN
		sp P41236 IPP2_HUMAN
6053	TTSFFLNSPEK2000020000	sp Q8WYP5 ELYS_HUMAN
6054	VASETHSEGSEYEELPK2000200000000000	sp Q5JSZ5 PRC2B_HUMAN
6055	AVLSPGSVFSR20000020000	sp P27987 IP3KB_HUMAN

6056	NTADHDESPRPTPGNAPSESSEDIDISSPNVSHDESIK2. 0200002E+28	sp	095251	KAT7_HUMAN
6057	TSSKESPIPSPTSDRK2002200020000000	sp	Q01082	SPTB2_HUMAN
6058	KSPGVEIK20000000	sp	Q9Y597	KCTD3_HUMAN
6059	KGSTFRPHDSFPK200000020000	sp	Q9P206	K1522_HUMAN
6060	TSVLGGGEDGIEVSPPEGMTEPGHSR20000000000000	sp	Q9Y618	NCOR2_HUMAN
6061	SPVSPGSSSPQLQVK20000000200000000	sp	Q8N3F8	MILK1_HUMAN
6062	LSTGTTVEDVQK200000000000	sp	Q86SQ0	PHLB2_HUMAN
6063	TRDDGDEEGLLTHSEEELEHSQDTDADDGALQ2E+19	sp	Q9UKM9	RALY_HUMAN
6064	SLDGVTNDR2000000000	sp	Q9UBM7	DHCR7_HUMAN
6065	GEAERGGSSGDEL20022000000	sp	Q7Z4V5	HDGR2_HUMAN
6066	TPTPSTPGDTQPNTAPVPPAEDGIK2. 2E+22	sp	Q14839	CHD4_HUMAN
6067	KVEEESPGDPDHEASTQGR200000000000000	sp	Q9NZZ2	OGFR_HUMAN
6068	SEHSHSTLPR200200000000	sp	P16144	ITB4_HUMAN
6069	HFSEHPSTSK20200	sp	043719	HTSF1_HUMAN
6070	QAPGVGAVGGSPEREVEGAGYNSEDEYEAAAAR2. 000000000002E+23	sp	Q96G74	OTUD5_HUMAN
6071	GAASTLVPGVSETSASPGSPSVR200200000	sp	Q6ZRS2	SRCAP_HUMAN
6072	TNLTSQSSTNLPSPGSPGSPGSPGSPGSPGSPK20020000000	sp	Q6IN85	P4R3A_HUMAN
6073	LVKPSSSSSGSR2000200	sp	Q9H6B4	CLMP_HUMAN
6074	RHSEQVANGPTPPR20000000000000	sp	A6ND36	FA83G_HUMAN
6075	EWYQSTIPQSPAPDDPEEGR200000000000000000	sp	Q08499	PDE4D_HUMAN
6076	KPTSPVKIPQNTTEYR20000000000000	sp	P25054	APC_HUMAN
6077	ESSEPPAPASSPEAASPTQGPAGTSK2. 0000002E+25	sp	Q92766	RREB1_HUMAN
6078	HDDGTQSDSENAGHR20000000000	sp	Q5SW79	CE170_HUMAN
6079	ESSTSEGADQKMSPEDDSDTKR220000000000	sp	Q15424	SAFB1_HUMAN
6080	YMSQMSVPEQAELEK2000000000	sp	Q15154	PCM1_HUMAN
6081	AGSSTPGDAPPAAEVQGR202000000000000000	sp	Q7Z6Z7	HUWE1_HUMAN
6082	SPGASNFTLTK20000000000000	sp	Q9Y4E8	UBP15_HUMAN
6083	AKSPTSPSPPR20202020000	sp	Q8IYB3	SRRM1_HUMAN
6084	EGEETEGSEDDKENDK20000000000	sp	P49321	NASP_HUMAN
6085	SPFEHSVEHK20000000000	sp	014646	CHD1_HUMAN
6086	EGMNPYDEYADSDQHDAYLER1. 002E+22	sp	Q08945	SSRP1_HUMAN
6087	GSVSQPSTSPPKPTGIFQTSANSSFEVK2. 2E+24	sp	015027	SC16A_HUMAN
6088	RRSSPSARPPDVPQQPQAAK2. 2E+19	sp	Q96JP5	ZFP91_HUMAN
6089	EWYQSTIPQSPAPDDPEEGR200000200000000000	sp	Q08499	PDE4D_HUMAN
6090	SGSPGSSSYEHYESR2020000000000000	sp	Q9P1Y6	PHRF1_HUMAN
6091	RSPSPTPTGSPSR20200000000000	sp	P49450	CENPA_HUMAN
6092	EPSSSLCSIDEEHLTDIEIVR2. 22E+20	sp	Q6GYQ0	RGA1_HUMAN
6093	SRGPTPPAMGPR2000200000000	sp	Q3KQU3	MA7D1_HUMAN
6094	DGDSYDPYDFSDTEEMPQVHTPK2. 0000002000001E+21	sp	P25205	MCM3_HUMAN
6095	RDSSSHEETPGSHPLYGHGECK2E+20	sp	Q8IVH2	FOXP4_HUMAN
6096	ENIKPNETSPSFSK20000	sp	Q99700	ATX2_HUMAN
6097	AGGGAQPSQTATGDPSDEDESDLEEFVR200000200000000	sp	Q2YD98	UVSSA_HUMAN
6098	LKSVEDEMDSPGEEPFTYTGQR2. 0000102E+20	sp	Q12857	NFIA_HUMAN
6099	YHGHSMSDPGVSYR1200002000	sp	P08559	ODPA_HUMAN
		sp	P29803	ODPAT_HUMAN
6100	SPTPTVNPR20000000	sp	Q8WY36	BBX_HUMAN
6101	KRTPSPSYQR202000000	sp	Q5T200	ZC3HD_HUMAN
6102	EKPELSEPHLNGPSSDPEAAFLSR2. 0000000002E+20	sp	Q9H4A3	WNK1_HUMAN
6103	SPGQPSSLR2000000000	sp	Q5T011	SZT2_HUMAN
6104	IETDEEESCDNAHGDNQPAR2. 00002E+19	sp	Q96T23	RSF1_HUMAN
6105	VASETHSEGSEYEELPK2000000000000000	sp	Q5JSZ5	PRC2B_HUMAN
6106	TSPKPAVVETVTTAKPQQIQALMDEVTK2E+27	sp	P56192	SYMC_HUMAN



6107	KPSVPDSASPADDSFVDPGER2E+19	sp	P16333	NCK1_HUMAN
6108	SRSSSVGSSSSYPISPAVSR200220000000000000	sp	Q15149	PLEC_HUMAN
6109	RSLSGSSPCPK20200200000	sp	Q9UQ35	SRRM2_HUMAN
6110	SPSWQRPNQGVSTGR200000000000000000	sp	Q96HC4	PDLI5_HUMAN
6111	SLVHEVGKPPQDVTDDSPSK200000	sp	Q96T58	MINT_HUMAN
6112	GNSRPGTPSAEGGSTSSTLR200000000000000000	sp	P35269	T2FA_HUMAN
6113	TPTSTPVHVK20000000000	sp	Q9ULM3	YETS2_HUMAN
6114	ASVALTPTAPSR200	sp	P28290	SSFA2_HUMAN
6115	LLPSVWSESEDGAGSPPPYR2000002000000	sp	Q9UIS9	MBD1_HUMAN
6116	SQSSEGVSSLSPPSNSLETQSQSLSR2020000000000000	sp	O95155	UBE4B_HUMAN
6117	QLPALDGSLMGPEPQAQEEAPVSPHK2. 01E+21	sp	Q8N9T8	KRI1_HUMAN
6118	RPQSPSPR202000	sp	Q99590	SCAFB_HUMAN
6119	TEMKSPFNPSQDSR100200000000000000	sp	P08651	NFIC_HUMAN
6120	SGSDAGEARPPTPASPR200000000000000000	sp	Q96FS4	SIPA1_HUMAN
6121	HKSGSMEEDVDTSPGGDYTSPSSPTSSSR2E+28	sp	P08651	NFIC_HUMAN
6122	IEIPVPTGQSVSPSPSIPGTPLK202000000000	sp	Q9H4Z3	PCIF1_HUMAN
6123	IGSDPLAYEPK2000000000	sp	P19634	SL9A1_HUMAN
6124	ISSSSFSR200	sp	P05787	K2C8_HUMAN
6125	LSPSWNPK20000000	sp	Q96BY7	ATG2B_HUMAN
6126	SSSPLPTVQLHPQSPTAGK200000000002000000	sp	O60245	PCDH7_HUMAN
6127	DRSPIRGSPR200002000	sp	Q9HCD5	NCOA5_HUMAN
6128	FHSPSTTWSFNK20000000000	sp	Q99590	SCAFB_HUMAN
6129	TDSV I IADQTPPTR20000002020000	sp	P17544	ATF7_HUMAN
6130	DEDESGGQRSEAEQEVRTPTK2000020000000020000	sp	Q9H4G0	E41L1_HUMAN
6131	SLNLEDYKK2000000000	sp	Q92541	RTF1_HUMAN
6132	KSNQIPTEVR2000000000	sp	Q9BWH2	FUND2_HUMAN
6133	IVVGSSVAPGNTAPSPSSPTSPTSDATTSLEMNPHAI PR22000000000000	sp	P49757	NUMB_HUMAN
6134	SKSPSPPR200020000	sp	Q9UKV3	ACINU_HUMAN
6135	SQDQDSEVNELSR2000000000000000	sp	Q86VM9	ZCH18_HUMAN
6136	GYTSDSEVYTDHGRPGK200000000000000000	sp	Q92538	GBF1_HUMAN
6137	KTESPIKLSPATPSR2000020000000	sp	Q9Y6X9	MORC2_HUMAN
6138	TPPVALNSSR20000000200	sp	Q9UQ35	SRRM2_HUMAN
6139	ALEQQAESDSSDTEEKDDDDDEEDVGKR2. 02202E+22	sp	Q9BXY0	MAK16_HUMAN
6140	SFADIGKK200000000	sp	Q9Y6M7	S4A7_HUMAN
6141	IKSSPSPEVVKPK20200000000	sp	Q6U841	S4A10_HUMAN
6142	LLSGPSQESPQTLGK20000000	sp	Q15652	JHD2C_HUMAN
6143	VSGEELHTGPPAPQGPLSVPQGLPTQSLASPPAR200000	sp	Q9BWE0	REPI1_HUMAN
6144	VSSEELHTGPPAPQGPLSVPQGLPTQSLASPPAR200000	sp	Q9UPN7	PP6R1_HUMAN
6145	SYSSPDITQAIQEEER200000000000000000	sp	P40818	UBP8_HUMAN
6146	RQSPSPSTRPIR20000200000	sp	Q8IYB3	SRRM1_HUMAN
6146	LASPSGSTSSGLEVVAPEGTSAPGGGPGTLDDSATICR2. 0000000002E+29	sp	Q13263	TIF1B_HUMAN
6147	REESPMDVDQSPSAQDTQSIASDGTQGEK200000000000000000	sp	Q7Z6Z7	HUWE1_HUMAN
		sp	P05534	1A24_HUMAN
		sp	Q09160	1A80_HUMAN
6148	GGSYSQAASSDSAQGSVSLTACK20002000000000	sp	P30447	1A23_HUMAN
		sp	P01891	1A68_HUMAN
		sp	P01892	1A02_HUMAN
		sp	P10316	1A69_HUMAN
6149	AQGEPVAGHESPK2000	sp	O94979	SC31A_HUMAN
6150	EASPAPLAQGEPR2000000000000000	sp	Q9P1Y6	PHRF1_HUMAN
6151	SRTSPVSR200200000	sp	Q9UQ35	SRRM2_HUMAN
6152	SHISDQSPSSK2000000	sp	P42166	LAP2A_HUMAN

6153	ILSQSTDSLNR200000100	sp Q92974 ARHG2_HUMAN
6154	TASRPDDIPDPSPPK20000000022000	sp Q5VZK9 LR16A_HUMAN
6155	FESPATGILSPR2000	sp P49792 RBP2_HUMAN
6156	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2.000200000002E+29	sp Q8WUZ0 BCL7C_HUMAN
6157	SILKPSTPIPPQEGEEVGESSEEQDNAPK2000000000	sp Q9UDY2 ZO2_HUMAN
6158	DAEYIYPSLESDDDDPALK2002000000000	sp Q9UPP1 PHF8_HUMAN
6159	HSPIAPSSPSPQVLAQK20000020000000000	sp Q9NQS7 INCE_HUMAN
6160	MALPPQEDATASPPR100000000020000	sp Q9UQ35 SRRM2_HUMAN
6161	SVSTTNIAGHFNDESPLGLR2000000000000000	sp Q92974 ARHG2_HUMAN
6162	VDSSTNSSPSPQQSESLPAHTSDFR2000000000	sp Q9BQE9 BCL7B_HUMAN
6163	SAYQEYSDSDVPEELK20000000000	sp Q96JG6 CC132_HUMAN
6164	SSSSVTTSETQPCTPSSSDYDLQR2E+23	sp P50552 VASP_HUMAN
6165	SPTPPSSAGLGSNSAPPIPDSR2200000000000000	sp Q8IWX8 CHERP_HUMAN
6166	TSFDQSDVDIFPSDFPTEPPSLPR2E+19	sp Q02880 TOP2B_HUMAN
6167	TDALTSSPGR2220000	sp P49736 MCM2_HUMAN
6168	MAESPCSPSGQQPPSPSPDEL PANVK2.000000002E+19	sp Q14160 SCRIB_HUMAN
6169	TESPATAAETAASELDNR20000000020000000	sp Q9NTJ3 SMC4_HUMAN
6170	APLKYPVSPSPDK2000	sp P41252 SYIC_HUMAN
6171	NPEPDSDEHTPVEDEEPK20002000000000	sp Q8N5F7 NKAP_HUMAN
6172	ASSPAESSPEDSGYMR2000020000000000	sp Q9Y4H2 IRS2_HUMAN
6173	TAAGEYDSVSESEDEEMLEIR30002020000100000	sp Q9BZE4 NOG1_HUMAN
6174	SSSELVTHLK20000000000	sp Q07960 RHG01_HUMAN
6175	IPSAVSTVSMQNIHPK2002000000000	sp P19634 SL9A1_HUMAN
6176	GPSQEEEDNHSDEEDRASEPKSQDQDSEVNELSR2.00000020002E+24	sp Q86VM9 ZCH18_HUMAN
6177	AIPVSPSAVEEDEDGHTVVATAR2E+19	sp Q07157 ZO1_HUMAN
6178	ELSPAAL EK200000000	sp 075683 SURF6_HUMAN
6179	AASVAAATTSPTPR200000	sp Q9Y2D9 ZN652_HUMAN
6180	TPNNVSTPAPSPDASQLASSLSSQK2.0002E+19	sp Q8WWI1 LM07_HUMAN
6181	HCSTYQPTPPLSPASK200000	sp Q9GZR1 SENP6_HUMAN
6182	AALLAQYADVTDDEEADADEKDDSGATTMNGSDK2E+24	sp Q96MW1 CCD43_HUMAN
6183	QSQQPMKPI SPVKDPVSPASQK100020000002000000	sp Q12888 TP53B_HUMAN
6184	SVSEINSDEL SGK2020002000000000	sp P82094 TMF1_HUMAN
6185	ACASPSAQVEGSPVAGSDGSQPAVK2000000000000000	sp Q9UFC0 LRWD1_HUMAN
6186	SSQPSTPQQAPPQPQQGTFVAHK2E+20	sp Q86VM9 ZCH18_HUMAN
6187	TGVTSTSDSEEGDDQEGEKK220000000000000000	sp 075475 PSIP1_HUMAN
6188	SPLSGIPVR2000000000	sp Q5M775 CYTSB_HUMAN
6189	AGMSSNQSISSPVLDVAVPR20000000000	sp Q9UQ35 SRRM2_HUMAN
6190	SQSPHYFR202000000	sp Q9UKJ3 GPTC8_HUMAN
6191	ERSPALK200000	sp Q9Y2W1 TR150_HUMAN
6192	QQFSPSMTER20000000	sp Q8TDB6 DTX3L_HUMAN
6193	AASDDGSLKSSSPTS YWK200020022000000000	sp Q92614 MY18A_HUMAN
6194	MAPALSGANLTSR2000	sp Q9UQ35 SRRM2_HUMAN
6195	SPTPPSSAGLGSNSAPPIPDSR220000200000000000	sp Q8IWX8 CHERP_HUMAN
6196	THSFENVSCHLPDSR2000000000	sp Q5VZ89 DEN4C_HUMAN
6197	APKPPTDGSTSTPSEDQALGK200000000000000000	sp Q8NHM5 KDM2B_HUMAN
6198	DSSTSYTETKDPSSGQEVATPPVQLQVCEPK3.00000002E+27	sp Q9UKV3 ACINU_HUMAN
6199	LSPGESAYQK2000000000	sp Q6NYC8 PPR18_HUMAN
6200	SFSPKSPLELGEK20000200000000	sp Q15742 NAB2_HUMAN
6201	RVSSDEEHTVDSCISDMK220000000000000000	sp Q70CQ2 UBP34_HUMAN
6202	KSPVFSDESDLDFDISK200000002000000000	sp 000566 MPP10_HUMAN
6203	TSPD TLR2000000	sp Q01130 SRSF2_HUMAN
6204	QQSFCAKPPSPLSPVPSVVK200200000000	sp Q70E73 RAPH1_HUMAN
6205	EHSLQVSPSLEK2000000	sp 095235 KI20A_HUMAN
6206	VETVSPSESPKDTIDK2000000200000000	sp Q99590 SCAFB_HUMAN

6207	NEDESPNKLYTLVTVYVPVTTFK200000000000000000	sp P62899 RL31_HUMAN
6208	SPSPAVPLR2020000000	sp Q6P4R8 NFRKB_HUMAN
6209	SSHSLDYDKDR200000000	sp Q9UK61 F208A_HUMAN
6210	ASPITNDGEDEFVPSDGLDKDEYTFSPGK2E+28	sp Q02880 TOP2B_HUMAN
6211	AVPTWASVQVDDPGSLASVESPGTPK2002000	sp Q8IVT2 MISP_HUMAN
		sp P30510 1C14_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
		sp Q29960 1C16_HUMAN
6212	GGSCSQAASSNSAQGSDESLIACK2E+20	sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P30505 1C08_HUMAN
6213	SLKESEQESEEEILAQKK20000000200000000000	sp Q9BW71 HIRP3_HUMAN
6214	EGMNPSTYDEYADSDQHDAYLER2E+19	sp Q08945 SSRP1_HUMAN
6215	ISSSSF2000000	sp P05787 K2C8_HUMAN
6216	SISGTSTSEKPNMMDTANTSPFK2E+23	sp Q9UJF2 NGAP_HUMAN
6217	VLSSTSEDEPGVVK22020000000000	sp Q8NI08 NCOA7_HUMAN
6218	VADVLDVLENEVDEYSGSSEK30002000	sp P38398 BRCA1_HUMAN
6219	NHSGSRTPVALNSSR20200000000000	sp Q9UQ35 SRRM2_HUMAN
6220	GTDTQTPAVLSPSK200	sp P46087 NOP2_HUMAN
6221	DMSLKDSGSDLSHRPK202000000000	sp O95251 KAT7_HUMAN
6222	KESKEETSIDVAGKPNEVTK2000000000000000	sp P53985 MOT1_HUMAN
6223	RRSQSIEQESQEK20200000000000	sp Q5VTL8 PR38B_HUMAN
6224	RSDSFLNIFPEK20000000000000	sp O94854 K0754_HUMAN
6225	TEVPGSPAGTEGNCQATGPSTVDTQNEPLDMK2E+28	sp Q32MZ4 LRRF1_HUMAN
6226	ASSLEDLVLK200000000000	sp Q15477 SKIV2_HUMAN
6227	KPPAPPSPVQSQSPSTNWSPAVPVKK2.00000002E+20	sp Q8IYB3 SRRM1_HUMAN
6228	SPSDLHISPLAK20000200000	sp O95785 WIZ_HUMAN
6229	TPQSPAPCVLLR200000000000	sp Q9Y613 FHOD1_HUMAN
6230	MIFEGPNKLSR2000	sp O94913 PCF11_HUMAN
6231	MLPHAPGVQMAIPEDAIPESGDEDEDPPDKR1.000000001E+33	sp Q13547 HDAC1_HUMAN
6232	HPSSPSALR200000000	sp O60504 VINEX_HUMAN
6233	RASVRSSEEDAR2000220000000000	sp Q8NC56 LEMD2_HUMAN
6234	KIDAGTMAEPSASPSK20000	sp Q8IXQ3 CI040_HUMAN
6235	SLPAFPTSSLLTQSQK20000000000000000000	sp Q99698 LYST_HUMAN
6236	YMNSDTTSPELR200000	sp Q96ST3 SIN3A_HUMAN
6237	LSGINEIPRPFSPVTSNTSPPPAAPLAR220000000000	sp Q0JRZ9 FCHO2_HUMAN
		sp Q641Q2 FA21A_HUMAN
6238	SRPTSFADLAAR2000000000000000	sp Q5SNT6 FA21B_HUMAN
		sp Q9Y4E1 FA21C_HUMAN
6239	EDELMEFSDLESDSEEKPCAKPR200020000000000000	sp Q9P2D1 CHD7_HUMAN
6240	SPVGKSPSTGTYGSSQK2E+19	sp Q9Y2W1 TR150_HUMAN
6241	YSSQDADEQDWEFQK2000000000000000	sp Q9C0C2 TB182_HUMAN
6242	TVDSQGPTPVCTPTFLER200020002000000000	sp Q9BYG3 MK67I_HUMAN
6243	VETVSPSESPKDTIDK200000000000	sp Q99590 SCAFB_HUMAN
6244	RIPSIVSSPLNSPLDR2000200000	sp P49790 NU153_HUMAN
6245	LSVSSNDTQESGNSSGSPGAK200000000000000000	sp Q9Y6D6 BIG1_HUMAN
6246	EIPEDVDMEEKESEDSDEENDFTEK1.000002002E+19	sp Q8IWA0 WDR75_HUMAN
6247	SSSESEDEDVIPATQCLTPGIR2.22202E+23	sp Q13428 TCOF_HUMAN
6248	AATAARPPAPPAPQPPSPTSPRPTLAR20000000020000	sp Q96B36 AKTS1_HUMAN

6249	FSDSEDEAATVPGNEPTTASTPK2. 02E+22	sp Q8N1G1 REX01_HUMAN
6250	SGTNSPPPPFSDWGR2000000000000	sp Q9UKT5 FBX4_HUMAN
6251	GDMSDEDDENEFFDAPEIITMPENLGHK1. 2E+26	sp P22059 OSBP1_HUMAN
6252	EADIDSSDESIEEDIDQPSAHK2002000000000000	sp Q9P2I0 CPSF2_HUMAN
6253	SSLGQSASETEEDTVSVSK20000000000	sp Q13435 SF3B2_HUMAN
6254	SKSPPKSPREEEGAVSS2020000000000000	sp Q01130 SRSF2_HUMAN
6255	ELASPVSPELR200000000	sp Q5T4S7 UBR4_HUMAN
6256	KPVTVSPTTPTSPTTEGEAS2200000000	sp Q9Y6G9 DC1L1_HUMAN
6257	TGQAGSLSGSPKPFSPQLSAPITTK2020000200000000000	sp Q9ULU4 PKCB1_HUMAN
6258	DGDSYDPYDFSDTEEEMPQVHTPK2. 0003000000001E+21	sp P25205 MCM3_HUMAN
6259	NSLSDPNIDR2000000000	sp Q9NRY4 RHG35_HUMAN
6260	GSPLLGPVVPGPSPIPSVTEK2. 00000000002E+20	sp P49454 CENPF_HUMAN
6261	DDSFDSLDSFGSR200000000000	sp Q9UPQ0 LIMC1_HUMAN
6262	TDSTSDGRPAWMR200000000000	sp Q14204 DYHC1_HUMAN
6263	EHYPVSSPSSPSPAQGGVSR2202000000000000000	sp 075179 ANR17_HUMAN
6264	TPTSPLSPSSQK20002000000	sp Q5VT06 CE350_HUMAN
6265	TPSPPPPPIEDIALGK2000000000000000	sp Q5T200 ZC3HD_HUMAN
6266	SCFESSDPPELK20000000	sp Q9UQ35 SRRM2_HUMAN
6267	TENSLANENQQPIKSEPESEGEPEK200020000000	sp Q8NHM5 KDM2B_HUMAN
6268	SLPASPSTSDFCQTR2000200000000000	sp Q9Y2D8 ADIP_HUMAN
6269	DSSSTDSANEKPALIPR2202000000000000	sp Q9NUQ6 SPS2L_HUMAN
6270	KPSSSLR2200000	sp Q13029 PRDM2_HUMAN
6271	SNSFSDER202000000	sp Q09666 AHNK_HUMAN
6272	IDSNISPK2000	sp Q9H501 ESF1_HUMAN
6273	RRSPSPPTR200000000	sp Q8IYB3 SRRM1_HUMAN
6274	APVPSTCSSTFPEELSPSHQAK200000000	sp Q14980 NUMA1_HUMAN
6275	VNSNGKESPGSSEFFQEAVSHGK2. 00000002E+21	sp Q8N108 MIER1_HUMAN
6276	KTSAVSSPLLDQQR2200000000	sp Q5T200 ZC3HD_HUMAN
6277	TRDDGDEGLLTHSEEELEHSQDTDADDGALQ2000000000	sp Q9UKM9 RALY_HUMAN
6278	SFGNPFEPQAR200000000000	sp Q8N5S9 KKCC1_HUMAN
6279	SSSFSELCHR2200000000	sp Q5T8I3 F102B_HUMAN
6280	GYSSPEPDIQDSSGSEAQSVKPSTR2. 200000002202E+23	sp Q9BZF1 OSBL8_HUMAN
6281	LEDAESSNPNLQSLSTDALPSASK20020000000000	sp Q9Y294 ASF1A_HUMAN
6282	DDPFTQPDQSTPSSFDLSPDQK200000	sp P98082 DAB2_HUMAN
6283	ESAAPASPAPSPAPSPPTAPPQK2. 000000002E+22	sp Q3KQU3 MA7D1_HUMAN
6284	SNSPLVPVPSK200000000000	sp Q13247 SRSF6_HUMAN
6285	EVEEDESEDEEMSEDEEDDSSGEEVVPQKK2. 00001200000022E+25	sp P19338 NUCL_HUMAN
6286	THSTSSSLGSGESPFSTR200200000000	sp Q9UGV2 NDRG3_HUMAN
6287	QSHSGSISPYPK200000200000	sp Q9UQ35 SRRM2_HUMAN
6288	RRSPSPTPTGPSR202000000000	sp P49450 CENPA_HUMAN
6289	AISAPTSPTR20000	sp Q9NQG6 MID51_HUMAN
6290	ESGRPIKPPR2000000000	sp P25440 BRD2_HUMAN
6291	RSLSPLGGR202000000	sp Q14004 CDK13_HUMAN
6292	AYSPTCSPTLGFK3000020000000	sp Q5TGY3 AHDC1_HUMAN
6293	GSKESLVPQR2000000000	sp 095835 LATS1_HUMAN
6294	IDFTPVSAPSPTR200020000	sp Q7Z309 F122B_HUMAN
6295	HSHSHSPMSTR202000000	sp P62995 TRA2B_HUMAN
6296	YSPSQNSPIHHIPSR2000020000000000	sp Q9NYF8 BCLF1_HUMAN
6297	TVFSPTLPAAR200000000	sp Q7Z2W4 ZCCHV_HUMAN
6298	SKTPPPPPQTAQTK2000000000000000	sp P25054 APC_HUMAN
6299	DPSPEADAPVLGSPEKEEAASEPPAAAPDAAPPPDRPIEK2E+39	sp 015234 CASC3_HUMAN
6300	HTGPNSPDTANDGFVR200000000	sp P31943 HNRH1_HUMAN
6301	QPPGVPNGPSSPTNESAPELPQR200000000000	sp P55795 HNRH2_HUMAN
		sp Q8TF74 WIPF2_HUMAN

6302	ALFKPPEDSQDDESDSDAEEEEQTTK200000020000000000	sp Q13769 THOC5_HUMAN
6303	TLSPEFNER20000000	sp Q9BSJ8 ESYT1_HUMAN
6304	SAQGSSSPVPSMVQK2000000000	sp P35658 NU214_HUMAN
6305	TPLSEAEFEEIMNR2000000000000000	sp Q16630 CPSF6_HUMAN
6306	SQGDEAGGHGEDRPEPLSPK2000	sp Q9NZT2 OGFR_HUMAN
6307	RGSPSPPR2020000	sp Q5VUA4 ZN318_HUMAN
6308	LYNSEESRPYTNK2000	sp Q9NYV4 CDK12_HUMAN
6309	HQASDSENEELPKPR20000000000000	sp Q96ST2 IWS1_HUMAN
6310	ASSPTKTSPTTPEASATNSPCTSKPATPAPSEK2. 202E+27	sp Q12873 CHD3_HUMAN
6311	TSSLTQFPSPQSEER2000000000000000	sp Q9H6Z4 RANB3_HUMAN
6312	SLLSHEFQDETDTTEEETLYSSKH2020000000000000	sp 075976 CBPD_HUMAN
6313	STSAPQMSPGSSDNQSSSPQPAQQK1. 00022E+19	sp Q14157 UBP2L_HUMAN
6314	TEDGGEFEEGASENNAKESSEPEKEAEEGCPK2. 0000002E+21	sp 043719 HTSF1_HUMAN
6315	GGSPDLWK2000000	sp Q96JM3 CHAP1_HUMAN
6316	IPSFPPSPEPPSPSAPSIAK20000000000	sp Q12802 AKP13_HUMAN
6317	RPACDPEPGESGSSSDEGCTVVRPEK200220000000000000	sp 015541 R113A_HUMAN
6318	SVPGTTSSPLVGDISP2000	sp Q9UQR0 SCML2_HUMAN
6319	SVCGHLENTSVGNPNPSSAENSFR200020002000000000	sp Q96HH9 GRAM3_HUMAN
6320	SGTPPRQGSITSPQANEQSVTPQR2. 02000002E+24	sp Q9UQ35 SRRM2_HUMAN
6321	KRSPSPSPTPEAK202000200000	sp Q8TAQ2 SMRC2_HUMAN
6322	KNSTGSGHSAQELPTIR200000000000000000	sp 094763 RMP_HUMAN
6323	YTPTSPSYSPSSPEYTPSPK200000000000000000	sp P24928 RPB1_HUMAN
6324	NNSNTCNIENELEDSR200000000000000000	sp Q6PL18 ATAD2_HUMAN
6325	RASPPDPSPSAASASER20000202000000000000	sp Q9Y2F5 ICE1_HUMAN
6326	DEIEGLSSDEEH2200000	sp Q6PD74 AAGAB_HUMAN
6327	EIAIVHSDAEKEQEEEEQK200000000000000000	sp Q9H307 PININ_HUMAN
6328	VSPAPGSAPETPEDK2000002000200000	sp Q13459 MYO9B_HUMAN
6329	EQLSETEGSGEDEPGNDPSETTQK2. 02E+21	sp Q13107 UBP4_HUMAN
6330	GYNHGQGSYSNSYNSPGGGGSDYNYESK200000000000000000	sp Q12906 ILF3_HUMAN
6331	SPGNKSDNIKPK20000200000000	sp Q16637 SMN_HUMAN
6332	TVFPGAVPVLASPPPK200000	sp Q9UBC2 EP15R_HUMAN
6333	EGSIETLSNSSGSTSGSIPR20000000000000000000	sp Q9H6U6 BCAS3_HUMAN
6334	YPLCSADAAVSTTPPGSPPPPPLPEPPVLK2. 02E+21	sp Q9BTC0 DIDO1_HUMAN
6335	SPLSDYMNLDLFSSPK2000001000002000	sp Q9Y4H2 IRS2_HUMAN
6336	FSEMMNMGDEDVDLPEVDGADDDSDSDEK1. 001E+29	sp Q15185 TEBP_HUMAN
6337	RQPQQGESEGSSEDEDGSRPLTR200222000000000000	sp Q6ZRS2 SRCAP_HUMAN
6338	RTDALTSPPGR2020000	sp P49736 MCM2_HUMAN
		sp P30505 1C08_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp Q29960 1C16_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
6339	GGSCSQAASSNSAQGSDESLIACK2000000000	sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp P30510 1C14_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
6340	RISSSSVQPCSEEVSTPQDSLQCK2E+20	sp P42694 HELZ_HUMAN
		sp P20700 LMNB1_HUMAN
6341	LKLSPSPSSR20200000	sp Q03252 LMNB2_HUMAN
		sp P19338 NUCL_HUMAN
6342	AAAAAPASEDEDEDDEDDEDDEDDEDDDEEDDSEEEAMETTPAK2E+34	sp Q9Y2F5 ICE1_HUMAN
6343	SCSSPAVSAVQLPLSPK20000000000000000000	sp 095684 FR10P_HUMAN
6344	GPTTGEGALDLSDVHSPPKSPEGK2E+21	

6345	AGLGSPPERPKTSPGSPR22000000	sp Q9H6F5	CCD86_HUMAN
6346	TSAQHALTSVSR20000	sp P61073	CXCR4_HUMAN
6347	SPEPDTLPLAAGSDHPLPR2E+19	sp Q6PCT2	FXL19_HUMAN
6348	HPSSPSALR22000000	sp O60504	VINEX_HUMAN
6349	SGSTSPLGFAR2020000000	sp O75385	ULK1_HUMAN
6350	SSFSSDPDEK2000000000	sp P46013	KI67_HUMAN
6351	TRTPASINATPANINLADLTR2.0000002E+19	sp Q7KZ85	SPT6H_HUMAN
6352	SFHFDPPLSSGSR200	sp Q9C0B5	ZDHC5_HUMAN
6353	AVGSPLCVPAR200000000	sp Q9NYZ3	GTSE1_HUMAN
6354	RREEGPPPPSPDGASSDAEPEPPSGR2000020000000000000	sp Q9NTJ3	SMC4_HUMAN
6355	TSQEELLAEVVQGQSR2000000000000000	sp Q6PJT7	ZC3HE_HUMAN
6356	RPPSQSLTSSSLSSPDIFDPSPEEDKEEHISLAHR2020000000000000000	sp Q96B97	SH3K1_HUMAN
6357	SNVSSPATPTASSSSSTPTR200000	sp Q12888	TP53B_HUMAN
6358	AIPGDQHPESPVHTEPMGIQGR2000000000000000	sp Q14676	MDC1_HUMAN
6359	ANSIGSTSASSVPNTDDEDSYHQEAYK2.002E+26	sp Q9UH92	MLX_HUMAN
6360	TPSDDEEDNLFAPPK2020000000000000	sp Q9Y4E1	FA21C_HUMAN
6361	DMHCLEASSPTFSK2000000	sp Q641Q2	FA21A_HUMAN
6362	TDSREDEISPPPNPVVK20000000000000000000	sp Q7Z333	SETX_HUMAN
6363	SPSESSPQHPTTPARPR20020000000000000000	sp P10644	KAPO_HUMAN
6364	RLGPSESR20200	sp O14733	MP2K7_HUMAN
6365	STSAPQMSPGSSDNQSSSPQPAQQK1.00002E+19	sp Q8NGJ1	OR4D6_HUMAN
6366	KGGSYSQAASSDSAQGSVSLTACK20002002000000	sp Q14157	UBP2L_HUMAN
6367	VAETHSEGSEYEELPK2000000000000000	sp Q09160	1A80_HUMAN
6368	SLHSAHSLASR200000200000	sp P05534	1A24_HUMAN
6369	QQSLPSLCSTSDPDTPLGAPSTPGR2.002002E+23	sp P30447	1A23_HUMAN
6370	LLPPASP200	sp P01891	1A68_HUMAN
6371	TRNSGIWESPELDR200000000000	sp P01892	1A02_HUMAN
6372	RLSSGKADGHVK22000000000	sp P10316	1A69_HUMAN
6373	LSPSPSSR200000	sp Q5JSZ5	PRC2B_HUMAN
6374	NSPPPPGR20000000	sp O15027	SC16A_HUMAN
6375	KKSPIINESR200000000	sp Q8TF72	SHRM3_HUMAN
6376	FSPHPSPAK200020000	sp Q9Y4R8	TELO2_HUMAN
6377	AEDQTESSCESHR20000000	sp O15085	ARHGB_HUMAN
6378	ECSPTSSLER200000000	sp O75962	TRIO_HUMAN
6379	SVPSSLQASAR200000000000	sp P20700	LMNB1_HUMAN
6380	SQSPHYFR2000000	sp Q03252	LMNB2_HUMAN
6381	YTPTSPSYSPSPEYTPSPK22000000000000000000	sp Q96ER3	SAAL1_HUMAN
6382	GGSYSQAASSDSAQGSVSLTACK3.000000000002E+21	sp Q13523	PRP4B_HUMAN
6383	LSPATPTSEGPK200200000000	sp O14681	EI24_HUMAN
6384	GPPSPAPVMHSPSR2000000000200	sp Q9BU76	MMTA2_HUMAN
6385	SSSNDVDEETAESDTSPVLEK2000000000000000000	sp Q9P1Y6	PHRF1_HUMAN
		sp P83369	LSM11_HUMAN
		sp Q9UKJ3	GPTC8_HUMAN
		sp P24928	RPB1_HUMAN
		sp P01891	1A68_HUMAN
		sp P01892	1A02_HUMAN
		sp P05534	1A24_HUMAN
		sp Q09160	1A80_HUMAN
		sp P10316	1A69_HUMAN
		sp P30447	1A23_HUMAN
		sp Q8N1G0	ZN687_HUMAN
		sp Q13573	SNW1_HUMAN
		sp Q99590	SCAFB_HUMAN

6386	VPPPALS200	sp Q14152 EIF3A_HUMAN
6387	AAPEASSPPASPLQHLLPGK2000000000000000	sp Q96TA1 NIBL1_HUMAN
6388	RSFSISPSR202000200	sp P18583 SON_HUMAN
6389	HDTDPSPR20002000	sp Q9BRD0 BUD13_HUMAN
6390	DSFDDRGPVLDYDHGSR2000000000000000	sp P43243 MATR3_HUMAN
6391	DEPQTVDPVPSFGESPLSPIDMTQER200020001000000	sp P17535 JUND_HUMAN
6392	GGHSSVSTESESSSFHSS2200000000000000	sp P61073 CXCR4_HUMAN
6393	RSPSPTTPGSPSR20002000000000	sp P49450 CENPA_HUMAN
6394	SKPIPIIMPASPQK20000	sp 000429 DNM1L_HUMAN
6395	SWDSSSPVDRPEPEAASPTTR2E+21	sp Q86WB0 NIPA_HUMAN
6396	KAASPSQSVR200002000	sp Q8IYB3 SRRM1_HUMAN
		sp P60709 ACTB_HUMAN
		sp P63261 ACTG_HUMAN
		sp Q562R1 ACTBL_HUMAN
		sp Q9BYX7 ACTBM_HUMAN
6397	SYELPDGQVITIGNER200000000000000000	sp A5A3E0 POTEF_HUMAN
		sp Q6S8J3 POTEE_HUMAN
		sp P62736 ACTA_HUMAN
		sp P63267 ACTH_HUMAN
		sp P68032 ACTC_HUMAN
		sp P68133 ACTS_HUMAN
6398	GASSPYGAPGTPR2000000000000000	sp Q5UIP0 RIF1_HUMAN
6399	VAHSDKPGSTSTASFR2000	sp Q9P0L0 VAPA_HUMAN
6400	LMHNASDSEVDQDDVVEWK2000000000000000	sp Q9H2P0 ADNP_HUMAN
6401	TPESQPDTPPGTPLVSQDEK2000000000000000	sp Q96T60 PNKP_HUMAN
6402	NTDVAQSPEAPK20000000	sp Q9P2E9 RRBP1_HUMAN
6403	ISPIQKPGGLDPVEILQER20000000000000000000	sp P51531 SMCA2_HUMAN
6404	GSYGSDAEEEEYR200000000000	sp Q9UDY2 ZO2_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
6405	KGGSYSQAASSDSAQGSVSLTACK2000000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
6406	EVEEDSEDEEMSEDEEDDSSGEEVVIPQK2.0000200000002E+24	sp P19338 NUCL_HUMAN
6407	SPQPESFK2000000000	sp P46013 KI67_HUMAN
6408	TKSPTDDEVTPSAVVR2000000000000000	sp Q9Y2H5 PKHA6_HUMAN
6409	MSQGPSPKTPTPSTPGDTQPNTPAPVPPAEDGIK2.002E+29	sp Q14839 CHD4_HUMAN
6410	LTSQSEAMALQSIR2000000000000000	sp Q9UPQ3 AGAP1_HUMAN
6411	RLSSASTGKPLSVEDDFEK20000000000000000000	sp 075152 ZC11A_HUMAN
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6413	MYSFDDVLEEGK1020000000000000	sp Q8WWI1 LMO7_HUMAN
6414	GYTSDDDTWEPEIHLEDCK2000000000000000	sp Q99549 MPP8_HUMAN
6415	NKKSPEIHR20000000	sp Q9Y2W1 TR150_HUMAN
6416	HPSSPSALR20000000	sp 060504 VINEX_HUMAN
6417	HSLSGSSPGMK20000000	sp Q9UQ35 SRRM2_HUMAN
6418	DWQDDQSDNQSDYSVASEEGDEFDER2.0002002002E+21	sp Q14839 CHD4_HUMAN
6419	SSSVTTSETQPCTPSSSDYSDLQR2E+24	sp P50552 VASP_HUMAN
6420	SASSDTSEELNSQDSPPK20200000000000000000	sp 014745 NHRF1_HUMAN
6421	SSDQDSTNKEAEAAGVKPAGVRPR2.0002E+23	sp Q92625 ANS1A_HUMAN
6422	RQSVSGLHR2000000000	sp Q8N1G4 LRC47_HUMAN
6423	YTPTSPSYSPSSPEYTPSPK20200000000000000000	sp P24928 RPB1_HUMAN
6424	AQLEPVASPAK200000	sp Q15149 PLEC_HUMAN
6425	LVSFHDDSDEDLLHI2000020000000000	sp P11717 MPRI_HUMAN

6426	LDSSIISEGK20020000	sp P30260 CDC27_HUMAN
6427	TGPPPIPSK20000	sp Q14135 VGLL4_HUMAN
6428	LSSDLTR200000	sp Q7Z6E9 RBBP6_HUMAN
6429	LGSTGSQPNSEAESVPENVPKPLLPPK2.02E+25	sp Q8IZ21 PHAR4_HUMAN
6430	MQNTDDEERPQLSDDER100200000000200000	sp Q8WVC0 LEO1_HUMAN
6431	LASDDRSPPR20000	sp P98175 RBM10_HUMAN
6432	AILGSYSELTPAEYSPQLTR2000000	sp Q9Y6D9 MD1L1_HUMAN
6433	VEQATKPSFER20000	sp Q96E39 RMXL1_HUMAN
6434	WKAEGHSDDESSEGSSESTSR200020200200000000	sp Q9P289 STK26_HUMAN
6435	SSSPELVTHLK220000000000	sp Q07960 RHG01_HUMAN
6436	YRSPYSGPK20000000	sp Q14498 RBM39_HUMAN
6437	GPSLNPVLDYDHGSR200	sp P43243 MATR3_HUMAN
6438	GSETGSETHESDLAPSDK2000002000000000000	sp P48634 PRC2A_HUMAN
6439	NDHLTSTSSPGVIVPESENK2000000000000000000	sp O75496 GEMI_HUMAN
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6441	TVSTSSQPEENVDR22000000000	sp P42684 ABL2_HUMAN
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		sp Q9TNN7 1C05_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
6444	GGSCSQAASSNSAQGSDESLIACK2000000000	sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
		sp P30510 1C14_HUMAN
6445	RATGDGSSPELPSLER2000220000000000	sp Q9H2Y7 ZN106_HUMAN
6446	TEPAAEAEAAASGPSESPSPAEEELPGSHAEPVPAQGEAPGEQAR2.0020202E+36	sp P55884 EIF3B_HUMAN
6447	KRESESEDETTPAAPQLIK2000000200000000000	sp O60832 DKC1_HUMAN
6448	VATIQQCQSPTNYLQGLDGKPIIAAPVFTK2E+22	sp Q86TC9 MYPN_HUMAN
6449	TQMAEVLPSPR1000002000	sp P11388 TOP2A_HUMAN
6450	YGPADVEDTTGSGATDSKDDDDIDLFGSDDEESEEAKR2E+30	sp P24534 EF1B_HUMAN
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6454	MQSLQLDCVAVPSSR1020000000002000	sp Q13546 RIPK1_HUMAN
6455	SASASHQADIK202000000000	sp Q14739 LBR_HUMAN
6456	SPAHTPLSR200	sp Q6P4Q7 CNNM4_HUMAN
6457	FQTGNKSPEVLR2000000	sp Q96S66 CLCC1_HUMAN
6458	QSHSEPSLQSK2000000000	sp Q9UQ35 SRRM2_HUMAN
6459	ALFKPPEDSQDDESDDAEQQTTKR2000000200000000000	sp Q13769 THOC5_HUMAN
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6461	SSSESEDEDVIPATQCLTPGIR2.2202E+22	sp Q13428 TCOF_HUMAN
6462	NLVSPAYCTQESR20000000000	sp Q9NUQ3 TXLNG_HUMAN
6463	IEEVLSPEGSPSK200020000	sp Q9UEY8 ADDG_HUMAN
6464	LEISPDSSPER200220000	sp Q99959 PKP2_HUMAN
6465	SDSLLSFR2000000	sp Q00587 BORG5_HUMAN
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6467	RPSVGSQSNQAGQGK20000000000	sp Q9UPP1 PHF8_HUMAN
6468	SGSMEEDVDTSPGGDYTSPSSPTSSSR1.00000000003E+25	sp P08651 NFIC_HUMAN



6469	ERPDLEAPAPGSPFR20000	sp Q9NQT8 KI13B_HUMAN
6470	QSSTPSAPELGQQPDVNISEWK200000000000000000	sp Q93100 KPBB_HUMAN
6471	ELEKPIQSKPQSPVIQAAAVSPK2000	sp Q9Y6D5 BIG2_HUMAN
6472	KPEKPLFSSASPQDSSPR200002000	sp 000712 NFIB_HUMAN
6473	GAGDGSDEEVDGKADGAEAKPAE200000000000000000	sp P35579 MYH9_HUMAN
6474	SVSEHTSCPAESASDAAPLQR200000000000000000	sp Q8WUA7 TB22A_HUMAN
6475	SINKLSDPDPFK2000002000000	sp P42566 EPS15_HUMAN
6476	ISSKSPGHMVILDQTK2002000100000000	sp Q9NWH9 SLTM_HUMAN
6477	VQTTPSKPGGDR2000000000	sp Q12834 CDC20_HUMAN
6478	TPSPGPVESK20200000000	sp Q5T0W9 FA83B_HUMAN
6479	VSPENLVDK200000000	sp 075376 NCOR1_HUMAN
6480	SPSLDNPTFPNLPSENPLK2E+21	sp Q14653 IRF3_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
6481	GGSYSQAASSDSAQGSVSLTACK2000000	sp P01892 1A02_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
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6484	SLTDPSQESHTEAISDAETSSDISFSGIATR222200200000000	sp Q5QJE6 TDIF2_HUMAN
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6486	RISTSDILSEK2000000000	sp 060293 ZC3H1_HUMAN
6487	SLESLADGGSASPIKDQPMVK2E+20	sp Q9HB96 FANCE_HUMAN
6488	SSSVGSSSYPISPAUSR23002000000	sp Q15149 PLEC_HUMAN
6489	GNSRPGTPSAEGGSTSSTLR202000000000000	sp P35269 T2FA_HUMAN
6490	SEEEASSTEKPTK200000000000000	sp Q9UGY1 NOL12_HUMAN
6491	GGISVQVNSIKFDSE200	sp Q14978 NOLC1_HUMAN
6492	KPHDCESSTVSEEDYFSSHR2020000000000	sp Q9H4H8 FA83D_HUMAN
6493	HELSPQKR2000000	sp Q9BXP5 SRRT_HUMAN
6494	RPPTPCSDPER200200000	sp Q9BW71 HIRP3_HUMAN
6495	KRSEGFMDR200000000	sp Q7Z4V5 HDGR2_HUMAN
6496	LPSVENRPK20000000	sp 015417 TNC18_HUMAN
6497	DGSLNQSSGYR2000000000	sp P06241 FYN_HUMAN
6498	EPTPVLGSGAAAAGR200000000000000	sp P42167 LAP2B_HUMAN
		sp P42166 LAP2A_HUMAN
6499	LPSKELPDSSSPVANNIR200000000000000000	sp Q9BWT3 PAPOG_HUMAN
6500	CSSPVDTECSHAEGSR200000000000000	sp 075170 PP6R2_HUMAN
6501	ILLPSFQDSEDDDDILPNMDK200000000000000	sp Q6P0N0 M18BP_HUMAN
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6503	SLTNSHLEK2000200000	sp Q9H2H9 S38A1_HUMAN
6504	SGSPSDNSGAEEMEVSLAKPK2.0200200001E+19	sp P31749 AKT1_HUMAN
6505	RPAPAVSPGSKPGPPGSPRPWK200000000002000000	sp Q96JM3 CHAP1_HUMAN
6506	ASSHSSQTQGGGSVTK200000000000000	sp P02545 LMNA_HUMAN
6507	TRSPSPDDILER2020200000000	sp Q13523 PRP4B_HUMAN
6508	AASAPAKESPR2000	sp Q13428 TCOF_HUMAN
6509	LSPENNVLTQK20000000000	sp Q16514 TAF12_HUMAN
6510	SPHSPSIATPPSQSSSCSDAPMLSTVHSAK2.020002E+28	sp Q9NYF8 BCLF1_HUMAN
6511	LAEAPSPAPTPSPTPVEDLGPQTSTSPGRLSPDFAEELR200002000000000	sp Q14160 SCRIB_HUMAN
6512	VETVSPSESPK202000	sp Q99590 SCAFB_HUMAN
6513	GHHVTDSENDEPLNLDSESEELHR2.0200000000002E+23	sp Q96ST2 IWS1_HUMAN
6514	VISHDRDSPPPPPPPPPPPPPQPSLK2E+24	sp Q69YN4 VIR_HUMAN
6515	RDSFDNCSLGESSK2000000000000	sp Q5UIP0 RIF1_HUMAN



6569	ILDELKDDSTHESLSQESESEEDGIHVDSQK200202000000000000	sp Q9H6T3 RPAP3_HUMAN
6570	KSLSDSESDSK202000000000	sp Q13185 CBX3_HUMAN
6571	TVDLGISDLEDDC20000000	sp Q8NHQ9 DDX55_HUMAN
6572	EVSSRPSTPGLSVVSGISATSEIPNKIEDLR2.002E+29	sp Q14C86 GAPD1_HUMAN
6573	ELDEEGSDPPLPGR200000000	sp Q9BRJ6 CG050_HUMAN
6574	SQDATFSPGSEQAEKSPGPIVSR2000000000000000	sp Q86WB0 NIPA_HUMAN
6575	QQSTPQAANSPPNLGAK200000000	sp Q9Y4C1 KDM3A_HUMAN
6576	SCFESSDPPELK2000020000000	sp Q9UQ35 SRRM2_HUMAN
6577	AAPQSPSVPK2000000	sp 075694 NU155_HUMAN
6578	LQELESCGLGSTSDTDVDR20200020000000000	sp Q14C86 GAPD1_HUMAN
6579	TEQELPRPQSPDLDSLDR202000200000	sp P54259 ATN1_HUMAN
6580	AENGERSDLEEDNER2000000000	sp Q8WU90 ZC3HF_HUMAN
6581	VSSQAEDTSSSFDNLFIDR200000000000	sp P0CAP2 GRL1A_HUMAN
6582	TPKDSPGIPPSAGAHQLFR2.0002E+19	sp Q9BZD3 GCOM2_HUMAN
6583	RISLSDMPR200000	sp Q15418 KS6A1_HUMAN
6584	ISNLSPEEEQGLWK20000000000	sp Q9ULU4 PKCB1_HUMAN
6585	VSVGDSSEFSPIQEAQQDR20000000000	sp Q5HYJ3 FA76B_HUMAN
6586	KESKEETSIDVAGKPNEVTK200000000000000	sp Q8N2Y8 RUSC2_HUMAN
6587	SHESNAPGSAGGQASEKPR200000	sp P53985 MOT1_HUMAN
6588	SEDEESLAGQK200000000000	sp Q6PJG2 EMSA1_HUMAN
6589	TQTPPVSPAPQTEER20200000000000000	sp Q96EZ8 MCRS1_HUMAN
6590	LLKPGEEPSEYTDDEEDTK320000000	sp Q14247 SRC8_HUMAN
6591	VAASMLSPGGQMEEVAGAVK200000000000000	sp Q15173 PGRC2_HUMAN
6592	DFATPSLHTSDQSPGK20000	sp Q9UMN6 KMT2B_HUMAN
6593	TTSFFLNSPEK20000	sp Q63HN8 RN213_HUMAN
6594	QDFSPERPK2000000	sp Q8WYP5 ELYS_HUMAN
6595	SLVDYENANK20000000000	sp Q6T4R5 NHS_HUMAN
6596	ASEDTTSGSPPK20000	sp Q9UNH7 SNX6_HUMAN
6597	YSPTSPTYSPVYTPSPK300020002000000	sp Q9BZZ5 API5_HUMAN
6598	IPVEGPLSPSR20000	sp P24928 RPB1_HUMAN
6599	HSHSPASSPKPLK20020000000	sp P00512 BCL9_HUMAN
6600	RLSSLRASTSK2200000000	sp Q9HDC5 JPH1_HUMAN
6601	FTSQDASTPASSELNK200000000000000	sp P62753 RS6_HUMAN
6602	KPSQTLQPSEDLADGK200000000000000	sp Q9HAW4 CLSPN_HUMAN
6603	GSISSTSEVHSPNVGLR20000000000000000	sp Q13029 PRDM2_HUMAN
6604	AGSPSPQPSGELPR2000000000000	sp Q8WWQ0 PHIP_HUMAN
6605	SAEPAEALVLACK200000000000000	sp Q92619 HMHA1_HUMAN
6606	ALSPVTSR2000000	sp Q96CW6 S7A60_HUMAN
6607	TSSTCSNESLSVGGTSVTPR2E+19	sp Q96L91 EP400_HUMAN
6608	TPASTPVSGTPQASPMIER2000002010000	sp 060343 TBCD4_HUMAN
6609	FSPTMGR2000000	sp Q9UHR4 BI2L1_HUMAN
6610	TGSSSPGGPPKPGSQLDSMLGSLQSDLNK2E+26	sp P78344 IF4G2_HUMAN
6611	NYQQNYQNSESGEKNEGSESAPEGQAQR3.00000000000002E+24	sp P49023 PAXI_HUMAN
6612	KEEPSQNDISPK2000	sp P67809 YBOX1_HUMAN
6613	APSASDSKADSDGAKPEPVAMAR2.02020002E+21	sp Q9NR30 DDX21_HUMAN
6614	GSSPTPPCSPVQPSK200020000000	sp Q7Z4V5 HDGR2_HUMAN
6615	GSGGLFSPSTAHPD GALGQR2000000000000000	sp Q9NUL3 STAU2_HUMAN
6616	EALGLGPPAAQLTPPPAPVGLR20000000000	sp Q9C0C2 TB182_HUMAN
6617	SSSLDGEHHDGYHR200000000000000	sp Q8IY67 RAVR1_HUMAN
6618	KTESFQNAQAGSNPK2000000000000	sp Q9C0J8 WDR33_HUMAN
6619	DKEPFTFSSPASGR20002000	sp Q92667 AKAP1_HUMAN
6620	EHYPVSSPSSPPAQPGGVSRR200000000000000	sp Q6P1X5 TAF2_HUMAN
6621	SMSVDETDKSPCEAGR20000000000000000	sp 075179 ANR17_HUMAN
		sp Q53SF7 COBL1_HUMAN

6622	DGDSYDPYDFSDTEEMPVHTPK2.000000000001E+21	sp P25205 MCM3_HUMAN
6623	GILAADESTGSIK20000	sp P04075 ALDOA_HUMAN
		sp ALDOA_RABIT
6624	VPHLEEVMSPVTTPTDEDVGHR200200000000000	sp Q3KR37 GRM1B_HUMAN
6625	SAPTSPTATNISSK20000000000	sp Q9H792 PEAK1_HUMAN
6626	IALESEGRPEEQMESDNCSGGDDDWHLSSK1.020002E+19	sp Q13501 SQSTM_HUMAN
6627	SSDGGSEPPDR200000000000	sp Q9HCD6 TANC2_HUMAN
6628	HDTVTVSSDLQFTK200000000	sp P30414 NKTR_HUMAN
6629	AVTTVTQSTPVPGPSVPPPEELQVSPGPR200000	sp P51610 HCFC1_HUMAN
6630	ANSDDERPVASDNDDEKQNSDDEEQQLSDEEK2.00000002E+31	sp Q8WVC0 LEO1_HUMAN
6631	AGLESGAEPGDGSDTTTKK2000000002000000	sp O60832 DKC1_HUMAN
6632	TPNNVSTPAPSPDASQLASSLSSQK2E+20	sp Q8WWI1 LMO7_HUMAN
6633	YSPTSPTYSPK200030000000	sp P24928 RPB1_HUMAN
6634	ECLTGESESSSEDEFK202220000000	sp Q56P03 EAPP_HUMAN
6635	ISSLLEEQFQGGK200000000000	sp P62241 RS8_HUMAN
6636	SGSSQELDVKPSASPQER2002000000000200000	sp Q9UQ35 SRRM2_HUMAN
6637	KPIETGSPK2000	sp Q05519 SRS11_HUMAN
6638	NSFTPLSSNTIR2000000000000	sp O60825 F262_HUMAN
6639	QGSITSPQANEQSVTPQRR200000000200000	sp Q9UQ35 SRRM2_HUMAN
6640	SMVSPVSPGTISVPNSCPASPR2.0002E+21	sp P85037 FO XK1_HUMAN
6641	HSPIAPSSPQVLAQK220000000000	sp Q9NQS7 INCE_HUMAN
6642	AGLSLEASGR2000	sp O60784 TOM1_HUMAN
6643	RKSLDSESDSK202000200000	sp Q13185 CBX3_HUMAN
6644	TSPCQPTISEGPVPSLPIRPSR2.000202E+22	sp O43426 SYNJ1_HUMAN
6645	SPSPGNHTSNSSNASNATVVPQNSSAR20000000000000000	sp Q9BTA9 WAC_HUMAN
6646	DSAYQSI THYRVSASR20000	sp P50402 EMD_HUMAN
6647	SPSLSQSASR200000000000	sp Q9H2P0 ADNP_HUMAN
6648	ESDKEDGRESYDTPSQR2000000000000000000	sp Q9Y6M7 S4A7_HUMAN
6649	TTSPEPQESPTLPSTEGQVVK2.2E+21	sp P53814 SMTN_HUMAN
6650	TLDVSTDEEDK22000000	sp P16383 GCFC2_HUMAN
6651	SQDATFSPGSEQAEK2000000000	sp Q86WB0 NIPA_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
6652	EDALDDSVSSSVHASPLASSPVR20002000200000	sp O14715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
		sp A6NKT7 RGPD3_HUMAN
6653	VIEPPAASCPSR2000	sp Q3MIN7 RGL3_HUMAN
6654	ATSSSSGSLSATGR2000000000000	sp Q03252 LMNB2_HUMAN
6655	SQSAAVTPSSTTSSTR20000000	sp Q16186 ADRM1_HUMAN
6656	TAGSSVTSLDGTR22000000	sp Q92597 NDRG1_HUMAN
6657	GSDYSWSYQTPSSPSSSSSR300000200000000	sp Q765P7 MTSSL_HUMAN
6658	LSSSDRYSDASDDSFSEPR30002002000000	sp Q9NRF8 PYRG2_HUMAN
6659	EEPSENIPQELLSNLGHPESPPR20000	sp Q13309 SKP2_HUMAN
6660	GASSPYGAPGTPR20000000000	sp Q5UIP0 RIF1_HUMAN
6661	KAPSASDSDKADSDGAKPEPVAMAR2.0202020002E+23	sp Q7Z4V5 HDGR2_HUMAN
6662	SPTEDFR20000000	sp Q8IXT5 RB12B_HUMAN
6663	GVHIHQAGGSPASSTSSSSLTNDVAK200000000	sp Q8N122 RPTOR_HUMAN
6664	LSSSSPGATVSPSSSDAEYDKLPDADLAR2.20002E+26	sp Q8NI08 NCOA7_HUMAN
6665	KSSFNVS DVARPEAAGSPPEEGGCTEGTPAK2E+29	sp Q92619 HMHA1_HUMAN
6666	RRSPSPAPPPR20000000	sp Q8IYB3 SRRM1_HUMAN
6667	YPLCSADAAVSTTPPGSPPPPPLPEPPVLK2.0002E+19	sp Q9BTC0 DIDO1_HUMAN
6668	KESESEDSSDEPLIK2002200000000	sp P35659 DEK_HUMAN
6669	MKTPVQYSQQNSPK20000000020000	sp P46013 KI67_HUMAN
6670	ERDHSPTPSVFNSDEER20000000000	sp Q6UN15 FIP1_HUMAN

6671	LTPVSLSNSPIK200020000	sp P46100 ATRX_HUMAN
6672	GSSGPRPVEGTPASR200000000000000	sp Q13112 CAF1B_HUMAN
6673	SPGSGSQSSGWHEVEPGMPSPTTLK2E+25	sp Q12857 NFIA_HUMAN
6674	KPPAPSPVQSQSPSTNWSPAVPVK2002000000000	sp Q8IYB3 SRRM1_HUMAN
6675	GPEAAGLSSPLSFPSQR2000000000	sp Q8TEK3 DOT1L_HUMAN
6676	SLSYSPVER2020200000	sp Q9UQ35 SRRM2_HUMAN
6677	SGSSQELDVKPSASPQER2000000000000000000	sp Q9UQ35 SRRM2_HUMAN
6678	AWLDEDSNLSPSPLR2002000000	sp Q9C086 IN80B_HUMAN
6679	IQSSLSVNSDISKK2002002000	sp Q9P1T7 MDFIC_HUMAN
6680	APQTSSSPPVVR20000000	sp Q8IYB3 SRRM1_HUMAN
6681	STSPAGQHHSPISR2200000000000000	sp Q5T200 ZC3HD_HUMAN
6682	YFQSPSR20000	sp Q16629 SRSF7_HUMAN
6683	SSTPLPTISSAENTR200000000000000200	sp P42166 LAP2A_HUMAN
6684	TVSASSTGDLPK2020002000000	sp P42167 LAP2B_HUMAN
6685	HGSSDISSPR22000	sp P78362 SRPK2_HUMAN
6686	ASLQSGQEGAGDSPGSQFSK2000000000	sp Q9BRD0 BUD13_HUMAN
6687	GGSGSHNWGTVKDELTEPK2000	sp Q12778 FOXO1_HUMAN
6688	KRPACDEPEGESGSSSDEGCTVVRPEK222000000000000	sp Q8NC51 PAIRB_HUMAN
6689	RPGGSSPLNAVPCGPPGSEPPR2000000000000000000	sp O15541 R113A_HUMAN
6690	KPSPSEPEPWKPFPAVSPEPR2000000000000000000	sp P48634 PRC2A_HUMAN
6691	TSPSSLPGR2000000000	sp Q96JM3 CHAP1_HUMAN
6692	HSLSGSSPGMK2200000	sp Q5SNT2 TM201_HUMAN
6693	SKPIPIMPASPQK10020000	sp Q9UQ35 SRRM2_HUMAN
6694	VPVPGPGSGNGPER2000000000	sp O00429 DNM1L_HUMAN
6695	RNSNSPPSPSSMNQR2020000000000000	sp Q6ZRV2 FA83H_HUMAN
6696	FGQAATMEGIGAIGGTPAFNR20000000	sp Q7Z5L9 I2BP2_HUMAN
6697	TAPSSPLTSPSDTR220020000000	sp Q15233 NONO_HUMAN
6698	SQSSDLLER2020000000	sp Q8IWB9 TEX2_HUMAN
6699	GNVVPSPLPTR20000000	sp Q8IVT2 MISP_HUMAN
6700	GPTTGEALDLSVHSPPKSPEGK20000000200000	sp Q9Y2V2 CHSP1_HUMAN
6701	SWDSSSPVDRPEPEAASPTR20000000000200000	sp O95684 FR10P_HUMAN
6702	FDHESPGTDEDK2002000000	sp Q86WB0 NIPA_HUMAN
6703	QEESESPVERPLKENPPNK2020000000000000000	sp Q13427 PPIG_HUMAN
6704	VVVSPTKK200000	sp Q96SB4 SRPK1_HUMAN
6705	GSSPTPPCSPVQPSK220000020000000	sp P19338 NUCL_HUMAN
6706	SPFNPSPPQDSR20000000000	sp Q9NUL3 STAU2_HUMAN
6707	SHSTEPNLSSFLNDPNMK2000000000000000000	sp P08651 NFIC_HUMAN
6708	GPPSPAPVMHSPSRK20000000200000	sp Q9UPQ0 LIMC1_HUMAN
6709	TPTSGPVITK2000000000	sp Q13573 SNW1_HUMAN
6710	RRSFSISPSR2020000000	sp Q6MZP7 LIN54_HUMAN
6711	GSDALSETSSVSHIEDLEK2000002000000000	sp P18583 SON_HUMAN
6712	LSPERSSDESNEPHLPHGVSQR2.00022E+24	sp Q6P996 PDXD1_HUMAN
6713	THDHQLESSLSPVEVFAK20200000000	sp Q9BQ52 RNZ2_HUMAN
6714	TGQAGSLSGSPKPFSPQLSAPITTK2000000000000000000	sp Q9H410 DSN1_HUMAN
6715	YFGSIDSSEAEAGAAR2002200000000000	sp Q9ULU4 PKCB1_HUMAN
6716	STSTPNVHMVSTTLPVDSR2000000000000000000	sp O15534 PER1_HUMAN
6717	ASPGTPLSPGLR2000002000000	sp P04049 RAF1_HUMAN
6718	QKSDAEEDGGTVSQEEEDRKPK2.00000002E+20	sp Q96BD0 SO4A1_HUMAN
6719	SQSAAVTPSSTTSSTR200000000000	sp P27824 CALX_HUMAN
6720	EYDSGDTDEIIAMK200000000000	sp Q16186 ADRM1_HUMAN
6721	VPSPTPAPK200000	sp Q76FK4 NOL8_HUMAN
6722	SEIPSPQSQLEDAEANLVGR2000000000000000	sp Q9UQ35 SRRM2_HUMAN
6723	VTGGVSESPSGFSK2000000000000000	sp Q9ULJ8 NEB1_HUMAN
		sp O14757 CHK1_HUMAN

6724	RSPRPDHPGTPPHK20000000200000	sp P30291 WEE1_HUMAN
6725	SRSPTPSSAGLGSNSAPP PDSR220000200000000000	sp Q81WX8 CHERP_HUMAN
6726	ASSLESSEPPK20000	sp Q9UHB6 LIMA1_HUMAN
6727	EYIPGQPPLSQSSDSSPTR20000000000	sp P07814 SYEP_HUMAN
6728	SYSPDGKESPSDK200000200000	sp P43243 MATR3_HUMAN
6729	DGDKSPMSSLQISNEK2010000000000	sp Q9UER7 DAXX_HUMAN
6730	SQDATFSPGSEQAEKSPGPIVSR200200000200000000	sp Q86WB0 NIPA_HUMAN
6731	GHPAQRPEPSPPEPEPQAPEGSQAGAEGPSSPEASR2E+27	sp Q6ZW31 SYDE1_HUMAN
6732	HGGSPQPLATTPLSQEPVNPPEASPTR2. 0000022E+25	sp Q9UQ35 SRRM2_HUMAN
6733	RPPSPDIVLSDNEQPSSPR200000020000020000	sp Q86YP4 P66A_HUMAN
6734	SSATSGDIWPGLSAYDNSPR2. 0000000000003E+19	sp Q9NYF8 BCLF1_HUMAN
6735	SSSQPSSCCSDPSKPGGVEGATQSLAEQMR2. 002E+29	sp Q13501 SQSTM_HUMAN
6736	EVSSRPSTPGLSVVSGISATSEIPNK2002000000000000	sp Q14C86 GAPD1_HUMAN
6737	ALSSGGSITSPPLSPALPK200002000000000000	sp Q6Y7W6 PERQ2_HUMAN
6738	GRNSATSADEQPHIGNYR20000000000000	sp Q7KZI7 MARK2_HUMAN
6739	TVAISDAAQLPHDYCTTPGGTLFSTTPGGTR3000000000002000000	sp Q13542 4EBP2_HUMAN
6740	LQSFDSPER20000	sp P78332 RBM6_HUMAN
6741	TRLSTASEETVQNR220200000000	sp O43379 WDR62_HUMAN
6742	GWSQEGPVKSPAECR20000002000000	sp Q9C0C2 TB182_HUMAN
6743	IHNVGSPLK20000	sp P52701 MSH6_HUMAN
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6745	QESCPHHPQVLAQQGSGSSPK200000000000000000	sp Q8N1G0 ZN687_HUMAN
6746	QEEDSRSSSEESDSDREALAAMNAAQVKPLGK2. 022200202E+28	sp Q13428 TCOF_HUMAN
6747	RYSPP QR2000000	sp Q8IYB3 SRRM1_HUMAN
6748	GSPSSHLLGADHGLR2000000000000000	sp Q6NWX9 PR40B_HUMAN
6749	YNQSI SLR2000	sp Q9UBH6 XPR1_HUMAN
6750	GRPSKTPSPSQPK202000000	sp Q9NTI5 PDS5B_HUMAN
6751	SWSTATSPLGGERPFR200000200000000000	sp Q9UPQ0 LIMC1_HUMAN
6752	VAHEPVAPPEDKESESEAK2000000	sp O95674 CDS2_HUMAN
6753	TEDGGEFEEGASENNAKESPEKEAEEGCPEK2. 00000002E+21	sp O43719 HTSF1_HUMAN
6754	LVEQANS PK2000	sp O95391 SLU7_HUMAN
6755	RSETPPHWR202000000	sp Q13427 PPIG_HUMAN
6756	EISSPARPCSFEEAMK20000000000000	sp O94916 NFAT5_HUMAN
6757	TSPTVATQTGASATSTR2000000000000000	sp Q6P1L5 F117B_HUMAN
6758	QAPGVGAVGGGSPEREV GAGYNSEDEYEAAAAR2E+23	sp Q96G74 OTUD5_HUMAN
6759	SLNFQEDDDTSR2200	sp Q9H814 PHAX_HUMAN
6760	SPTTPIDPEK20000000000	sp Q08AD1 CAMP2_HUMAN
6761	YSHSYLSDSDTEAK200200000000	sp Q92614 MY18A_HUMAN
6762	ASSHSSQTQGGGSVTK2000000000000200	sp P02545 LMNA_HUMAN
6763	GPSHPLDLGTSSPNTSQIHWPYR200000000000000	sp O60504 VINEX_HUMAN
6764	LDDAHSLSGSGAGEGYEPI SDDELDELILAGDAEK2000000000000000	sp Q5T200 ZC3HD_HUMAN
6765	EAYNDSSDEESSHHGPGVQCAHQ3. 0002E+22	sp O60884 DNJA2_HUMAN
6766	SPSSPELNK20020000000	sp Q9UQ35 SRRM2_HUMAN
6767	NASASFQELEDKK200000000000	sp Q99543 DNJC2_HUMAN
6768	SGSIVELIAGGGSSCSPVLSR2000000000	sp P42858 HD_HUMAN
6769	TEDGGEFEEGASENNAKESPEKEAEEGCPEK2200000000000000	sp O43719 HTSF1_HUMAN
6770	SPAKSPNPSTPR2000200000000	sp O76021 RL1D1_HUMAN
6771	MAEESSSSSSSSPTAATSQQQLK20000000000000	sp Q13620 CUL4B_HUMAN
6772	SEALSSVLDQEAMEIK22000000000000	sp Q12888 TP53B_HUMAN
6773	KLSASYEK2000000	sp Q9Y297 FBW1A_HUMAN
6774	SVSSPTSSNTPTPTK2000000	sp Q5M775 CYTSB_HUMAN
6775	RHSSVSDSQPCEPPSVGTEYSQGASPPQHQHLK2. 2E+31	sp P39880 CUX1_HUMAN
6776	RESPAPKPR2020000000	sp Q8IYB3 SRRM1_HUMAN
6777	LASPMKPVGTPPSSK200000002000000	sp Q15648 MED1_HUMAN

6778	YPLCSADAAVSTTPPGSPPPPPPLPEPPVLK2.00002E+20	sp Q9BTC0 DID01_HUMAN
6779	LDTGPQSLSGK2000	sp P35269 T2FA_HUMAN
6780	TPSDDEEDNLFAPPK2000000000000000	sp Q641Q2 FA21A_HUMAN
6781	EKGPTTGEGALDLSDVHSPPKSPEGK20000000200000	sp Q9Y4E1 FA21C_HUMAN
6782	TPASGLQTPSTPAPGSATR20200	sp 095684 FR10P_HUMAN
6783	TGVTSTSDSEEEGDDQEGEK220000000000000000	sp Q96DF8 DGC14_HUMAN
6784	RPLPVESPDTR2000000	sp 075475 PSIP1_HUMAN
6785	SLSLDPGQSLEPHPEGPQR200000000000000000	sp P57682 KLF3_HUMAN
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6788	RASSPSLIER220000000	sp P01106 MYC_HUMAN
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6790	GCSDPDPDTSYVLTPTHTEEK200000000020000000	sp A1L170 CA226_HUMAN
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6794	HRHSPTGPPGFPR20000000000	sp Q66K14 TBC9B_HUMAN
6795	MLPHAPGVQMIPAIPDAIPEESGDEDEDDPDKR1.00000000002E+24	sp Q96JM3 CHAP1_HUMAN
6796	TEPAAEAEASGPSESPSPAEEELPGSHAEPVPAQGEAPGEQAR2.02E+31	sp P98175 RBM10_HUMAN
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6798	NSNSPPSPSSMNR20000200000000	sp P55884 EIF3B_HUMAN
6799	KSSFNVDVARPEAAGSPPEEGGCTEGTPAK2.0000000000002E+29	sp Q9BUU2 MET22_HUMAN
6800	SRSPLLNDR2020000000	sp Q7Z5L9 I2BP2_HUMAN
6801	EPESPASAPIPLFGADTIGQR2000000000000000	sp Q92619 HMHA1_HUMAN
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6803	SRSPTPPSSAGLGSNSAPPIDSR220000000000000000	sp Q96AY4 TTC28_HUMAN
6804	DHFYSDDDAIEADSEGDAEPCDK3.0000000002E+20	sp 075533 SF3B1_HUMAN
6805	TDSREDEISPPPNPVVK20000000000	sp Q8IWX8 CHERP_HUMAN
6806	SLSRPSLIEQEK20000000000000	sp Q9Y3B9 RRP15_HUMAN
6807	SYSPDGKESPSDKK200000002000000	sp P10644 KAP0_HUMAN
6808	RSSQPSPTAVPASDSPPTK2000000000000000	sp Q12802 AKP13_HUMAN
6809	TFSEPGDHPGMLTSGK2000000000000000	sp P43243 MATR3_HUMAN
6810	TRSPSPDDILER20200000000	sp Q3KQU3 MA7D1_HUMAN
6811	IESDVQEPTPEDDLDIMLGNK2.000002E+20	sp Q9HA47 UCK1_HUMAN
6812	SPGLLDTPPIR200000000000	sp Q13523 PRP4B_HUMAN
6813	STPLASPSPPGR200000000	sp P20042 IF2B_HUMAN
6814	SPNSHDSEPWLLR2000000000000000	sp Q8IY92 SLX4_HUMAN
6815	TQPSSGVDSAVGTLPATSPQSTSVQAK20000000000	sp Q8WYP5 ELYS_HUMAN
6816	AESSGGGTVPSSAGILEQGSPGDGSPPKPK2000000	sp Q5T5Y3 CAMP1_HUMAN
6817	NQSHSSPSVSPSR2202020000	sp Q13428 TCOF_HUMAN
6818	YSPSQNSPIHHIPSR202000000000000	sp Q68EM7 RHG17_HUMAN
6819	DDSHAEDSEDEKEDHKNVR200020000000000000	sp Q7Z6E9 RBBP6_HUMAN
6820	RRSSTVAPAQPDGAESEWTDVETR2.2E+21	sp Q9NYF8 BCLF1_HUMAN
6821	CQTAEADSESDHEVPEPESEMK2.0000002E+20	sp Q9H1E3 NUCKS_HUMAN
6822	AENAEGQTPAIGPDGEPLDETSQMSDLPVK2002000000	sp Q02241 KIF23_HUMAN
6823	SLNSVVTSEPDKHSLLVGDFR200020000000000000	sp Q9BPX3 CND3_HUMAN
6824	RHSHSHSPMSTR20200010000	sp P51532 SMCA4_HUMAN
6825	SHMSGSPGPGSNTAPSTPVIGGSDKPGMEEK2.02E+21	sp Q04656 ATP7A_HUMAN
6826	SPQPDPVKTPTSSK200000002000000	sp P62995 TRA2B_HUMAN
6827	SKPNLPSESER20200	sp A0FGR8 ESYT2_HUMAN
6828	IPSIVSSPLNSPLDR20002000200000	sp P46013 KI67_HUMAN
		sp Q08170 SRSF4_HUMAN
		sp P49790 NU153_HUMAN

6829	EEEEGISQESSEEEQ2002000000	sp P17096 HMGA1_HUMAN
6830	ASSLNENV DHSALLK20000000000000	sp Q96SB3 NEB2_HUMAN
6831	LSPERSSDESSENEPHLPHGVSQR2. 202E+20	sp Q9BQ52 RNZ2_HUMAN
6832	FSPFLQTAEDNTQDEVAGAPADHHGSPDEEQSGPPEDK2000002000000	sp Q92766 RREB1_HUMAN
6833	VNSASSSNPPAEVDPDTILK200000000000000000	sp Q9BY77 PDIP3_HUMAN
6834	LGPQDSDPTEANLESADPELCIR200000000000000000	sp Q27J81 INF2_HUMAN
6835	AGSKIPTPK20000000	sp O60308 CE104_HUMAN
6836	GRDSPYQSR2000000	sp Q13242 SRSF9_HUMAN
6837	VAAQSPSPASAEALLPALPLR2000002000000000000	sp Q9Y4F5 C170B_HUMAN
6838	TSGAGSPQTPPER200200000	sp Q96MH2 HEXI2_HUMAN
6839	AGGLQSDTEDECWSDTEAVPR20200000000000000	sp Q9H6H4 REEP4_HUMAN
6840	ISDSESEDPPR20002000000	sp Q96ST2 IWS1_HUMAN
6841	SHLKSESSDESNTNKK2022002000000	sp Q9H7Z3 NRDE2_HUMAN
6842	AASADSTTEGTPADGFTVLSTK22000000000000000	sp O75367 H2AY_HUMAN
6843	SKSPVLHPPK20200000000	sp Q9NX31 OSER1_HUMAN
6844	LEGVDGGQSPR2000	sp O00418 EF2K_HUMAN
6845	LSLEGDHSTPPSAYGSVK20000000000	sp P07355 ANXA2_HUMAN
6846	TGSGSPFAGNSPAR2000000000000	sp A6NMY6 AXA2L_HUMAN
6847	LPLLPESPGPLR2000000	sp Q9UPT8 ZC3H4_HUMAN
6848	LKEDIENEDEQNSPPKK200000	sp Q14160 SCRIB_HUMAN
6849	SPSVSSPEPAEK2020020000000	sp Q9NTI5 PDS5B_HUMAN
6850	ANTPDSDI TEKTEDSSVETPDNER2. 002002002002E+23	sp Q9UQ35 SRRM2_HUMAN
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6852	TCFSPNR20000	sp Q13433 S39A6_HUMAN
6853	RASPPDPSPSAASASER2000000000000	sp O00303 EIF3F_HUMAN
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6855	EKTIASDSEEEAGK2020000000	sp Q9H3S7 PTN23_HUMAN
6856	LATPPAAQESSR20000000000	sp Q96ST2 IWS1_HUMAN
6857	DDISEIQSLASDHSGR2000000	sp Q9Y4B4 ARIP4_HUMAN
6858	DKDSGSSSPLPK2000200000	sp Q07157 ZO1_HUMAN
6859	GPNYTSGYGTSNELSNPSETESER20020000000	sp Q86VQ1 GLC11_HUMAN
6860	EGSEFSFDGEVAEK20000200000000	sp P51114 FXR1_HUMAN
6861	STQIENQHQAQDTS DLMSPSK20000	sp Q9BVS4 RIOK2_HUMAN
6862	VGVEASEETPQTSSSSARPGTPSDHQSQEASQFER2000000000000000	sp Q9NVR2 INT10_HUMAN
6863	TQLASWSDPTEETGPVAGILD TETLEK2E+23	sp Q6Y7W6 PERQ2_HUMAN
6864	EQGTESRSSTPLPTISSSAENTR2022000000000000000	sp Q15149 PLEC_HUMAN
6865	KNSFGYR200000	sp P42166 LAP2A_HUMAN
6866	ESEDSETQFPDTHLEAYGPCLSPPR20000	sp P42167 LAP2B_HUMAN
6867	DTHEDHDTSTENTDES NHDPQFEP IVSLPEQEIK2E+19	sp Q92615 LAR4B_HUMAN
6868	VSQSALNPHQSPDFK200000	sp Q14676 MDC1_HUMAN
6869	STIGVMVTASHNPEEDNGVK1020000000000000	sp P43487 RANG_HUMAN
6870	RPEGPGAQAPSSPR20000	sp Q9NYL2 MLTK_HUMAN
6871	SLSIEIGHEVK2000000000	sp O95394 AGM1_HUMAN
6872	VPLSNENSQVPQSPEDWR2000000	sp P40222 TXLNA_HUMAN
6873	DFTPVCTTELGR20000000000	sp O15155 BET1_HUMAN
6874	ALSSGGSITSPPLSPALPK200002000000	sp Q8IZT6 ASPM_HUMAN
6875	GSPHYFSPFRPY200002000000	sp P30041 PRDX6_HUMAN
6876	SFDYNYR20000000	sp Q6Y7W6 PERQ2_HUMAN
6877	TLNMTTSPEEK200000	sp Q13242 SRSF9_HUMAN
6878	SFSISPVR202000000	sp O75494 SRS10_HUMAN
6879	SPQKPIVR200000000	sp P49915 GUAA_HUMAN
6880	LVSFHDDSDEDLLHI200000000	sp P18583 SON_HUMAN
		sp P15056 BRAF_HUMAN
		sp P11717 MPRI_HUMAN



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6882	KDDSHAEDSEDEKEDHK2000000000	sp	Q9H1E3	NUCKS_HUMAN
6883	EVQAEQPSSSSPR2000	sp	Q9Y388	RBMX2_HUMAN
6884	KTESPIKLSPATPSR200000020000000	sp	Q9Y6X9	MORC2_HUMAN
6885	STGVSFWTQSDENEQEQQSDTEEGSNKK20000000000	sp	014974	MYPT1_HUMAN
6886	RTSPSSLPR200000000	sp	Q5SNT2	TM201_HUMAN
6887	YEDDGISDDEIEGK200000000	sp	Q9Y2K7	KDM2A_HUMAN
6888	YSPTSPTYSPTSPVYTPTSPK220000000000	sp	P24928	RPB1_HUMAN
6889	KESAPPSLR20000000	sp	Q8WUI4	HDAC7_HUMAN
6890	DDPGQETDSEDEDIIGPMPAK20220000000010000	sp	Q8IXQ4	GPAM1_HUMAN
6891	GSETGSETHESDLAPSDK2020000000000000	sp	P48634	PRC2A_HUMAN
6892	HQDGLPYIDDSPSSPHLSSK200020000000	sp	P11274	BCR_HUMAN
6893	TGSEPALSPAVVR2000000	sp	075815	BCAR3_HUMAN
6894	SMGTGDTPGLEVPSSPLR100000000000020000	sp	Q86WB0	NIPA_HUMAN
6895	GATPPAPPR20000000	sp	Q6SPF0	SAMD1_HUMAN
6896	GSPTGGAQLLK20000000000	sp	Q14684	RRP1B_HUMAN
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6898	LSSVMCDSDSDSDILVR200002002002000000	sp	Q8N4S0	CCD82_HUMAN
6899	SPSPSPAAVNHSSSDISPVSNESTSSSPGK220000	sp	075381	PEX14_HUMAN
6900	RGSLSQEMAK200000000	sp	Q8NFC6	BD1L1_HUMAN
6901	RNSNSPPSPSSMNQR20200002000000	sp	Q7Z5L9	I2BP2_HUMAN
6902	GESAPTLSTSPSPSSPSTSPSPTLGR20000	sp	Q9H330	TM245_HUMAN
6903	VAFDPEQKPLHGVLKPTSSPASSPLVAK202000002000000	sp	Q14684	RRP1B_HUMAN
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		sp	POCL80	GG12F_HUMAN
		sp	POCL81	GG12G_HUMAN
		sp	POCL82	GG12I_HUMAN
6906	YVQPPEMIGPMRPEQFSDEVEPATPEEGEPATQR200000000000	sp	Q13065	GAGE1_HUMAN
		sp	Q13068	GAGE4_HUMAN
		sp	076087	GAGE7_HUMAN
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6908	SPAAPSPALR200002000000	sp	Q9P270	SLAI2_HUMAN
6909	VEMYSGDDDDDFNK10202000000000	sp	060841	IF2P_HUMAN
6910	SLSGSPLK20000	sp	Q9UIG0	BAZ1B_HUMAN
6911	LSQSSQDSSPVR200000020000	sp	095251	KAT7_HUMAN
6912	ERSPPLSPEEK2000200000	sp	Q8N157	AHI1_HUMAN
6913	NATDLQNSSMSEEELTK20000000000	sp	P40855	PEX19_HUMAN
6914	TSSTCSNESLSVGGTSVTPR2. 2E+20	sp	060343	TBCD4_HUMAN
6915	SPSPGLFR202000000	sp	Q9HAW4	CLSPN_HUMAN
6916	INSDDSNLYLTASK2002000000000	sp	Q9BZG1	RAB34_HUMAN
6917	SLPSTSSTSSTKR20000	sp	Q12857	NFIA_HUMAN
6918	EDKPPLAPSGGTEGPEQPPPCPSQTGSPPVGLIK200000000	sp	Q8WWM7	ATX2L_HUMAN
6919	RGSDASDFDISEIQSVCSDEVETVPQTHRPTPR2. 002E+30	sp	Q03001	DYST_HUMAN
6920	TTQSMQDFPVVDSEEEAEEEFQK1. 00000002E+19	sp	Q9UBB9	TFP11_HUMAN
6921	SRSPLAIR202000000	sp	Q9UQ35	SRRM2_HUMAN
6922	QKSDAEDGGTVSQEEEDRKP20000000000	sp	P27824	CALX_HUMAN
6923	GTYPSSPTR20000	sp	Q9Y4P8	WIPI2_HUMAN
6924	TSSAFVGKTPEASPEPK20000000000200000	sp	Q99460	PSMD1_HUMAN
6925	DEILPTTPISEQK20000000	sp	P23396	RS3_HUMAN
6926	KELSEESEDEELQLEEFMLK2002000000000000000	sp	Q99543	DNJC2_HUMAN
		sp	P42167	LAP2B_HUMAN
6927	EQGTESRSSTPLPTISSAENTR2. 02E+20	sp	P42166	LAP2A_HUMAN
6928	GLPPSMER20000	sp	P38159	RBMX_HUMAN

6929	ERPLQQVSPVASVPWR2000000000	sp Q9BY89 K1671_HUMAN
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6931	LSSSPSGSGGGGASSSSHCSGESQCR2E+25	sp Q9Y4X0 AMMR1_HUMAN
6932	QHEAPSNRPLNELLTPQGSPR2000	sp P10398 ARAF_HUMAN
6933	SASVNKEPVSPLGIMR20000000000100	sp Q8WWI1 LMO7_HUMAN
6934	KFSAGGSDPPLK20000000000000	sp O75152 ZC11A_HUMAN
6935	SAPTSPTATNISSK2002000000000000	sp Q9H792 PEAK1_HUMAN
6936	VPCNVEGISPELEK20000000	sp Q86VQ1 GLC11_HUMAN
6937	LQPSSSPENSLDPFPPR2000000000000000	sp Q8WWM7 ATX2L_HUMAN
6938	SSISGSLPSSGSLQAVSSR20000000000000000000	sp Q9BXF6 RFIP5_HUMAN
6939	KPSKYPLCSADAAVSTTPPGSPPPPPLPEPPVLK2.000002E+21	sp Q9BTC0 DIDO1_HUMAN
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6941	VTAEADSSSPTGILATSESK2000000000000000	sp A0MZ66 SHOT1_HUMAN
6942	EPLENSASESSDTELPEK2020200000000000	sp P78337 PITX1_HUMAN
6943	STPLASPSPPGRSPQR2020200020000	sp Q8WYP5 ELYS_HUMAN
6944	VSPSPDEHQEAENAVSSGNR20000000000000000000	sp Q9BXS6 NUSAP_HUMAN
6945	KDPSGASNPADSPLHR2000000	sp P55317 FOXA1_HUMAN
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6947	FDIYDPFHPTDEAYSPPPAPEQK20000000000	sp Q9H7N4 SFR19_HUMAN
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		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
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		sp P30504 1C04_HUMAN
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		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
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6952	ASPSKPASAPASR2000000000000000	sp P27816 MAP4_HUMAN
6953	SAMDSPVPASMFAPESPSPAAR1.00000001E+21	sp O00178 GTPB1_HUMAN
6954	VQEHEDSGDSEVENEAK2000000000	sp Q9Y5J1 UTP18_HUMAN
6955	VDEDSAEDTQSNKGKEVVEVGQK2000000000000000	sp Q9H2G2 SLK_HUMAN
6956	SELSQDAEPAGSQETK200	sp Q9BVJ6 UT14A_HUMAN
6957	EFSGGSYSAK2000000000	sp Q5VT25 MRCKA_HUMAN
6958	TVASDLQK200200000	sp Q7L8J4 3BP5L_HUMAN
6959	EHHPEEGSSGSEVEEIPETPCESQGEELK2.202E+22	sp Q12888 TP53B_HUMAN
6960	VSPSPQESLSSSK202000000000	sp Q5JSH3 WDR44_HUMAN
6961	EVSPDPSTGAASPASSSAGGSAAR2.0002E+23	sp P23769 GATA2_HUMAN
6962	SFLESNYFTKPNLK2000000000000000	sp O60293 ZC3H1_HUMAN
6963	DSGRGDSVSDSGSDALR20000000000000000000	sp Q53EL6 PDCD4_HUMAN
6964	GEFSASPMLK201000	sp Q9UQ35 SRRM2_HUMAN
6965	LEPNSPMHGGTLLLEDVVGGR20000000000000000000	sp Q7Z4H7 HAUS6_HUMAN
6966	ESRQESDPEDDDVKKPALQSSVATSK2.0002E+26	sp Q9H307 PININ_HUMAN
6967	APSASDSKADSDGAKPEPVAMAR2.0002020002E+23	sp Q7Z4V5 HDGR2_HUMAN
6968	SLSPVAAPPLREPR2020000000000000	sp Q8NDT2 RB15B_HUMAN
6969	MLSQDAPTVK2000000000	sp O60934 NBN_HUMAN
6970	EHQTSAYSPPR20000	sp Q96K21 ANCHR_HUMAN
6971	SGSMEEDVDTSPGGDYITSPSSPTSSSR1.0000000000033E+25	sp P08651 NFIC_HUMAN
6972	QLWDSPETAPAAR20000000000	sp Q9Y613 FHOD1_HUMAN

6973	SEACPCQPDSGSPLPAEEEEK2000000000	sp Q14160 SCRIB_HUMAN
6974	APSDTGSLSPWDQDRR2000202000000000	sp O15320 CTGE5_HUMAN
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6976	SPVGKSPSTGSTYGSSQK2.00002002E+19	sp Q9Y2W1 TR150_HUMAN
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6979	RTPSPPPR20200000	sp Q8IYB3 SRRM1_HUMAN
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6986	LASPSGSTSSGLEVVAPEGTSAPGGGPGTLDDSATICR2.2E+32	sp Q13263 TIF1B_HUMAN
6987	NLVSPAYCTQESREEIPGGEAR200000000000	sp Q9NUQ3 TXLNG_HUMAN
6988	SMSENAVR2000000	sp Q9GZY8 MFF_HUMAN
6989	SSLSPSSAYER202000000000	sp Q9BYW2 SETD2_HUMAN
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6993	SSPQLDPLRK2000000000	sp Q8ND56 LS14A_HUMAN
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6995	ESAAPASPAPSPAPSPAPPQK200000000000000000	sp Q3KQU3 MA7D1_HUMAN
6996	SETSSNPSSPEICPNK22002000000000	sp Q684P5 RPGP2_HUMAN
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		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
		sp Q29960 1C16_HUMAN
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		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
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		sp Q29865 1C18_HUMAN
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		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
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7005	SPTLSQVHSPLVTSANLK20002200000000	sp Q86UU0 BCL9L_HUMAN
7006	LVEDERSDREETESSEGEAAAGGAK2.00002022E+21	sp Q96G23 CERS2_HUMAN

7007	SRSSSVGSSSSYPISPAVSR2. 022E+20	sp Q15149 PLEC_HUMAN
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7009	QLLQTLQAAEGEAAAAAGAGAGAAAAGAEGPGSPGVPSPPEAASEPPTGLR2. 000002E+20	sp Q8N196 SIX5_HUMAN
7010	SEAEEGEVRTPTK20000000000200	sp Q9H4G0 E41L1_HUMAN
7011	AGHDGDSGDSDDEEGYFICPITDDPSSNQVNSK2. 0000000003E+29	sp 000418 EF2K_HUMAN
7012	ITVQASPNDK2000000	sp Q5TCX8 M3KL4_HUMAN
7013	DTLSNPQSPQSPYNSPKPQHK203000000000	sp Q96PU5 NED4L_HUMAN
7014	SLSPVAAPPLR2000000000	sp Q8NDT2 RB15B_HUMAN
7015	TDHGAEIVYKSPVSGDTSR200000022000	sp TAU_HUMAN  sp P10636 TAU_HUMAN sp P05534 1A24_HUMAN sp Q09160 1A80_HUMAN sp P30447 1A23_HUMAN sp P01891 1A68_HUMAN sp P01892 1A02_HUMAN sp P10316 1A69_HUMAN
7016	GGSYSQAASSDSAQGSVDVSLTACK2002000000	sp P49790 NU153_HUMAN
7017	DSTSQHDDDNISTTSGFSSRASDKDITVSK2. 222002E+19	sp Q9H6B4 CLMP_HUMAN
7018	LVKPSSSSSGSR200	sp P29375 KDM5A_HUMAN
7019	VVSSVSSPR200000	sp P25440 BRD2_HUMAN
7020	ADTTTPTPTAILAPGSPASPPGSLEPK2002000000000	sp Q8IWB9 TEX2_HUMAN
7021	SPLQSAESSPTAGK200000022000000	sp Q13523 PRP4B_HUMAN
7022	EQPEMEDANSEKSINEENGEVSEDQSQNK2. 002000000002E+20	sp Q5T200 ZC3HD_HUMAN
7023	TPSPSYQR202000000	sp Q9HB19 PKHA2_HUMAN
7024	SQSYIPTSGCR2000000000	sp P49585 PCY1A_HUMAN
7025	TSPPCSPANLSR200000000000	sp Q9UK61 F208A_HUMAN
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7027	SPTESVNTPVGK20000000000	sp Q9UHI6 DDX20_HUMAN
7028	SYLEGSSDNQLK22000000	sp 075691 UTP20_HUMAN
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7036	LDNTPASPPRSPAEPNDIPIAK200020000000000000	sp Q96RL1 UIMC1_HUMAN
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7038	RQSVSPPYKEPSAYQSSTR200300000000000000	sp Q14839 CHD4_HUMAN
7039	MSQPGSPSPKPTPTSTPGDTQPNTAPVPPAEDGIK2. 0002E+35	
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7041	TNSSSSSPVVLK20000000000	sp Q7Z589 EMSY_HUMAN
7042	GTDDLNPVTSTPAKPSSPPPEFSFNTPGK20000000000000	sp Q6NXT4 ZNT6_HUMAN
7043	SHLLSSSDAEGNYR200000000	sp Q12830 BPTF_HUMAN
7044	EVEEDSEDEEMSEDEEDSSGEEVVIPQKK2. 00000200000022E+25	sp P19338 NUCL_HUMAN
7045	GGIDNPAITSDQELDDKK2000000000	sp Q13017 RHG05_HUMAN
7046	VGVEASEETPQTSSSSARPGTSPDHQSQEASQFER2. 00002E+20	sp Q6Y7W6 PERQ2_HUMAN
7047	SSSQRSPSPGNHTSNSSNASNATVVPQNSSAR2. 002E+31	sp Q9BTA9 WAC_HUMAN
7048	SRTSPVTR200200000	sp Q9UQ35 SRRM2_HUMAN
7049	LSPERSDSESNENEPHLPHGVSQR2. 00002E+24	sp Q9BQ52 RNZ2_HUMAN
7050	KRSPSPSPTPEAK200000000000	sp Q8TAQ2 SMRC2_HUMAN
7051	EAYSGCSGPVDECPSPSPVHK200000	sp Q8N556 AFAP1_HUMAN
7052	ESISPQPADSACSSPAPSTGK2200000000	sp Q9GZR1 SENP6_HUMAN

7053	NGDECAYHHPISPCCK20000	sp	Q6PJT7	ZC3HE_HUMAN
7054	SPFNPSPPQDSR20002000	sp	P08651	NFIC_HUMAN
7055	YYEDLDEVSTSSVSQSLESEDAR2200000000000000	sp	P35658	NU214_HUMAN
7056	ESISQPADSACSSPAPSTGK20000000000000000000	sp	Q9GZR1	SENP6_HUMAN
7057	APSPAPSSVPLGSEKPSNVSQDR2E+21	sp	075179	ANR17_HUMAN
7058	ENSSPSVTSANLDHTKPCWYWDK2E+21	sp	075909	CCNK_HUMAN
7059	HNDIVSDSDAEDRGTLAELTAAHFGGGGGLLHK20002000000000000000	sp	P78316	NOP14_HUMAN
7060	WDETPASQMGSTPVLTPGK200000000	sp	075533	SF3B1_HUMAN
7061	TADSSSEDEEEYVVEK20020220000000000000	sp	P45973	CBX5_HUMAN
7062	TLSPTPASATAPTSQGIPTSDEESTPK2000020000000000	sp	Q9NQS7	INCE_HUMAN
7063	TISQSSSLK20000000	sp	Q9UHB7	AFF4_HUMAN
7064	SLSYSPVER2000200000	sp	Q9UQ35	SRRM2_HUMAN
7065	SPQTLAPVGEDAMK20000000000000	sp	Q92797	SYMPK_HUMAN
7066	IDFIPVSPAPSPTR200020000	sp	Q96E09	F122A_HUMAN
7067	DDGDEGLLTHSEEELEHSQTDADDGALQ20020000000000	sp	Q9UKM9	RALY_HUMAN
7068	REESPMVDVQPSAQTQSIASDGTQGEK2.01000002E+28	sp	Q7Z6Z7	HUWE1_HUMAN
7069	VQMTSPSSTGSPMFK12000000001000	sp	P49790	NU153_HUMAN
7070	YSPSQNSPIHHIPSR2000000000	sp	Q9NYF8	BCLF1_HUMAN
7071	EKPELSEPSHLNGPSSDPEAAFLSR200000000000	sp	Q9H4A3	WNK1_HUMAN
7072	AGPDLPSLPSHALEDEGWAAAAPSGAR2000	sp	Q9H3Q1	BORG4_HUMAN
7073	GQKPALDCSQVSPRPATSPENNASLSDTSPMDSSPSGIPK202000000000000000000000	sp	P18858	DNLI1_HUMAN
7074	TENGTCPSPQPPLSPAAALGSGSAAAVPK2.0000000000002E+29	sp	Q12948	FOXC1_HUMAN
7075	RISQSQPVR20000000	sp	095071	UBR5_HUMAN
7076	SNSSSEAVLGQEELSAQAK2E+19	sp	Q9BXF6	RFIP5_HUMAN
7077	LASTSDIEEK220000000	sp	014974	MYPT1_HUMAN
7078	KRPSLPSSPGLPK2002000000000	sp	043294	TGFI1_HUMAN
7079	IADLQSPLFR200000	sp	Q9NZJ4	SACS_HUMAN
7080	SPGKAEAESDALPDDTVIESEALPSDIAAEAR20000000000000000000	sp	Q9GZR7	DDX24_HUMAN
7081	SGTSSPQSPVFR2000200000	sp	Q8N556	AFAP1_HUMAN
7082	TGDLGIPPNEPDRSPSEPIYNSEGK20200000000000	sp	Q15637	SF01_HUMAN
7083	SPPRSPPSPPTHR200002000000000000	sp	Q96HA1	P121A_HUMAN
7084	TPESQPDTPPGTPLVSQDEK2000200000000000	sp	Q96T60	PNKP_HUMAN
7085	GRLSPVPVPR20000000	sp	Q9UKM9	RALY_HUMAN
7086	GDTPGHATPGHGATSSAR20000000000000000000	sp	075533	SF3B1_HUMAN
7087	RDYDDMSPR2000	sp	P61978	HNRPK_HUMAN
7088	LPSDTPDSDASEDPGGAER20020020000000000000	sp	Q9UJK0	TSR3_HUMAN
7089	KPSKYPLCSADAAVSTTPPGSPPPPPLPEPPVLK2.2E+20	sp	Q9BTC0	DIDO1_HUMAN
7090	LEVSPQAQTAR2000000000	sp	Q6WKZ4	RFIP1_HUMAN
7091	LSNCDPPPTYEEATGQVNLQR2E+20	sp	014668	TMG1_HUMAN
7092	TFSATVR200000	sp	Q9Y2V2	CHSP1_HUMAN
7093	IDFIPVSPAPSPTR20000	sp	Q96E09	F122A_HUMAN
7094	RDSSSHEETPGSHPLYGHGECK20000000000000	sp	Q8IVH2	FOXP4_HUMAN
7095	QPSWDSPVSPSTVPAPSPLSAAAVSPSK2.0002E+26	sp	Q9NQC3	RTN4_HUMAN
7096	GYSSPEPDIQDSSGSEAQSVKPSTR2020000000000000	sp	Q9BZF1	OSBL8_HUMAN
7097	KETTSGTSTEPVKNSPAPPQAPGK20000000000000	sp	Q9NYV4	CDK12_HUMAN
7098	AAKPGPAEAPSPTASPSGDASPPATAPYDPR2.02E+19	sp	Q9UPT8	ZC3H4_HUMAN
		sp	Q09160	1A80_HUMAN
		sp	P05534	1A24_HUMAN
		sp	P30447	1A23_HUMAN
7099	GGSYSQAASSDSAQGSVSLTACK2000000	sp	P01891	1A68_HUMAN
		sp	P01892	1A02_HUMAN
		sp	P10316	1A69_HUMAN

7100	NSSQSGGKPGSSPITK2000000	sp Q15648 MED1_HUMAN
7101	HASAPSHVQPSDSEK20000000000000	sp O60343 TBCD4_HUMAN
7102	RLSSSESPQRDPPPPPPPPPLLR2.0002E+21	sp Q9H2G4 TSYL2_HUMAN
7103	SSDEENGPPSSPDLDR2000000022000000	sp Q96B36 AKTS1_HUMAN
7104	TASPLLRPDAPVEK2020000000000000	sp Q9HCK8 CHD8_HUMAN
7105	TLSDDISCPLESDLAYSDDDVPVYENGLSQK2.0000000003E+27	sp Q6PL18 ATAD2_HUMAN
7106	LSMEDSKSPPPK10000200000	sp Q9NQG5 RPR1B_HUMAN
7107	SSLSSHSHQSQIYR20200000000	sp O15027 SC16A_HUMAN
7108	DDSHSAEDESEDEKEDHKNVR2020000000000000000	sp Q9H1E3 NUCKS_HUMAN
7109	FSISPDDESSSYSSNSDFNYSYPTK2022000000000000	sp Q96QD8 S38A2_HUMAN
7110	RESVNLGK2000000	sp P46013 KI67_HUMAN
7111	GNDPLTSSPGR2020000	sp P49736 MCM2_HUMAN
7112	RPAEATSSPTSPERPR2002000000	sp Q9BU76 MMTA2_HUMAN
7113	MSPKPELTEEK2000000000000	sp P41208 CETN2_HUMAN
7114	SRSFDYNYR20000000000	sp O75494 SRS10_HUMAN
7115	QVLTAPEGAGQPRSEDEDSLEEAGSPAGPCPR2E+30	sp Q8TBB5 KLDC4_HUMAN
7116	ATVTPSPVKGK202000000	sp Q9H1E3 NUCKS_HUMAN
7117	SYSSPDITQAIQEEER20000000000000000000	sp P40818 UBP8_HUMAN
7118	LVDVSPDRGSPPSR20000200000	sp Q9NYZ3 GTSE1_HUMAN
7119	KNSSTDQGSDEEGLQK20002000000000	sp Q12789 TF3C1_HUMAN
7120	ASPAPGSGHPEGGAHLDMNSLDR2E+23	sp O94826 TOM70_HUMAN
7121	KGSDALRPPVQGEDEVPK2000000000000000000	sp Q8N3D4 EH1L1_HUMAN
7122	GNKSPSPDGSPAATPEIR2000020000000000	sp O00499 BIN1_HUMAN
7123	AASSAAQGAQGN20000000000	sp O15127 SCAM2_HUMAN
7124	KRSPSPSPTPEAK20000200000	sp Q8TAQ2 SMRC2_HUMAN
7125	GPPSPAPVMHSPSR2000001000200	sp Q13573 SNW1_HUMAN
7126	ESSPEKEAEEGCEK2000000000000000	sp O43719 HTSF1_HUMAN
7127	AYLGYQAVPSGSSGLSTSSSSPPGTPSPADAK2000200000000	sp P17676 CEBPB_HUMAN
7128	RHSHSHSPMSTR200002000000	sp P62995 TRA2B_HUMAN
7129	LTPVSLNSPIK200000020000	sp P46100 ATRX_HUMAN
7130	KVPVPEGLDLDAWINEPLSDSESEDERPR202020000000	sp O14617 AP3D1_HUMAN
7131	STDEYSEADEEEEEEGKPSR2.0002E+20	sp Q96T23 RSF1_HUMAN
7132	AGGPTPLSPTR20020000	sp P20700 LMNB1_HUMAN
7133	SERPPTILMTEEPSSPK1000002000	sp Q5VZK9 LR16A_HUMAN
7134	TSFVSGSDELGPIR2000020000000000	sp Q9NRY4 RHG35_HUMAN
7135	DISEGGESPVVQSDEEEVQVDTALATSHTDR2.00002E+29	sp Q96EV8 DTBP1_HUMAN
7136	GSSPTPPCSPVQPSK20200020000000	sp Q9NUL3 STAU2_HUMAN
7137	SPQRSPLQSAESSPTAGK2000200020002000000	sp Q8IWB9 TEX2_HUMAN
7138	LGGDEGYEEEEEDENSRQSEDDSETEKPEADDPK2.0000000002E+20	sp Q9UHR5 S30BP_HUMAN
7139	TAHNSEADLEESFNEHELEPSSPK2000	sp Q8IWS0 PHF6_HUMAN
7140	LNSDDTYQTALLSGSDEE2020000	sp Q96B21 TM45B_HUMAN
7141	HGTDLWIDNMSSAVPNHSPEK20000	sp Q8N6H7 ARFG2_HUMAN
7142	TSPADHGGVSGSESGSAVDSVAGEHSVSGR2.00202E+23	sp Q5T4S7 UBR4_HUMAN
7143	TRESHRSRSSPTAGPSTEGAEGPEEK2.002022E+24	sp Q8NHM5 KDM2B_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
		sp O14715 RGPD8_HUMAN
7144	DSLITPHVSR2000200	sp Q99666 RGPD5_HUMAN
		sp A6NKT7 RGPD3_HUMAN
		sp PODJD0 RGPD1_HUMAN
		sp PODJD1 RGPD2_HUMAN
7145	GLPLGSAVSSPVLFSVGR200000	sp Q8WUM0 NU133_HUMAN
7146	SGECSPVPMGSFPR200020001000000	sp P25054 APC_HUMAN
7147	SPAVATSTAAPPPSSPLPSK200000020000000	sp Q14157 UBP2L_HUMAN

7148	YSPTSPTYSPVYTPSPK30000030002000	sp P24928 RPB1_HUMAN
7149	SRSPTPQNQDGMTMVEK2000000000000000	sp Q3KR37 GRM1B_HUMAN
7150	IESDVQEPTPEDDLIDLGNK2000000000000000	sp P20042 IF2B_HUMAN
7151	FTGSFDDDDPDRDPYGEVDR2E+19	sp O15027 SC16A_HUMAN
7152	VEVTSPNLDVSLPSMEVDIQAPGAK2E+21	sp Q8IVF2 AHNK2_HUMAN
7153	SPSPLGTSVR20200000000	sp Q86SQ0 PHLB2_HUMAN
7154	AAPPPALTPDSQTVDSCKTPEVSFLPEEATEEAGVR2.002E+30	sp Q9H7N4 SFR19_HUMAN
7155	SHSESASPSALSSSPNLSPTGWSQPK2000000000	sp P04049 RAF1_HUMAN
7156	IYHPSCYEDYQNTSSFDCTSPSPK20000	sp O94913 PCF11_HUMAN
7157	TEMKSPFNPSQDSR20000020002000	sp P08651 NFIC_HUMAN
7158	DGTAPPPQSPGSPGTGQDEEWSDEESPRK2.000000000002E+21	sp Q9UKS6 PACN3_HUMAN
7159	GFGGAGELGQTPQVQAALSVSQENPEMEGPER20000000000000	sp Q8NBJ4 GOLM1_HUMAN
7160	FSSVSSPQPR200000	sp Q9P206 K1522_HUMAN
7161	GESAADSDGWSAPSDLR20000000000000	sp Q9BUL5 PHF23_HUMAN
7162	TVNSGGSSEPSPEVDVSR2000000000	sp Q6PIJ6 FBX38_HUMAN
7163	LPSSPASPSK200000	sp Q9H4G0 E41L1_HUMAN
7164	SVIEGVEDSDISDDEPSVYSA20020000000000	sp Q9NUN5 LMBD1_HUMAN
7165	TVSFSSMPTEK202000000000	sp Q9UPRO PLCL2_HUMAN
7166	ILSSMGNDKSSIAGSEDAEPLAPPIKPIKVTR2.20002E+24	sp Q03164 KMT2A_HUMAN
7167	VQIPVSRDPEPVSNEEDSYDEEIHDP200000030000000000	sp Q07157 ZO1_HUMAN
7168	MLQAISPK2000	sp P49585 PCY1A_HUMAN
7169	MHNLHGTKPPPSEGSDEEEEEDEEEDDEER200000000000000000	sp Q9BQ67 GRWD1_HUMAN
7170	QQPEPEWIGDGESTSPSDK2000	sp Q8NE71 ABCF1_HUMAN
7171	GESLSLPGSPPPDGTEQVIISR2E+19	sp Q9BZQ8 NIBAN_HUMAN
7172	APGYSSPVTTASGTTLR20000000000000	sp Q9HCK8 CHD8_HUMAN
7173	ETQSPEQVK2000000	sp Q9NYF8 BCLF1_HUMAN
7174	AEEQQLPPLSPPSPSTPNHR200000200000	sp Q6WCQ1 MPRIP_HUMAN
7175	SLSSGESGPGSPTHSHLSR2E+19	sp Q6P0Q8 MAST2_HUMAN
7176	QPGYQPPNPHGSPSPAAPASK2000000000	sp Q9BUL9 RPP25_HUMAN
7177	SVSEHTSCPAESASDAAPLQR2E+22	sp Q8WUA7 TB22A_HUMAN
7178	SNSQSDSHDEEVSPTPNPVVK20200000000000000000	sp P31321 KAP1_HUMAN
7179	EKFPEFCSSPSPVEVK20000000	sp Q68E01 INT3_HUMAN
7180	SPSPGNHTSNSSNASNATVVPQNSSAR2.000002E+26	sp Q9BTA9 WAC_HUMAN
7181	GPEDYPEGVEESSGEASK22000000	sp Q2NKX8 ERC6L_HUMAN
7182	SPTGPSNSFLANMGTVAHK20000000000000	sp Q96I25 SPF45_HUMAN
7183	STAPETAIECTQAPAPASEDEK200000	sp Q14839 CHD4_HUMAN
7184	LAPVPSPEPQKPAVSPESVK2000000	sp Q96JM3 CHAP1_HUMAN
7185	EYDSGDTDEIIAMK30000200000100	sp Q76FK4 NOL8_HUMAN
7186	ITPHTSPAPR2000200000	sp Q58A45 PAN3_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
7187	GGSYSQAASSDSAQSDVSLTACKV200020020000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
7188	SKSDGEAKPEPSPPR2000000000002000	sp P26358 DNMT1_HUMAN
7189	SGTSSPQSPVFR20200200000	sp Q8N556 AFAP1_HUMAN
7190	SNPESIAETPPAR2002000200000	sp Q9Y4H2 IRS2_HUMAN
7191	AAESETPGKSPEK20000020000	sp O76021 RL1D1_HUMAN
7192	RLSSSATLLNSPDR2000000020000	sp Q14244 MAP7_HUMAN
7193	SPPVDCPR200000000	sp Q5VZK9 LR16A_HUMAN
7194	TIQEVLEEQSEDEDREK2000000000	sp P18858 DNLI1_HUMAN
7195	VLSTSSLTR200200000	sp P16144 ITB4_HUMAN
7196	SRSGSSQELDVKPSASPQR202200000000200000	sp Q9UQ35 SRRM2_HUMAN

7197	SLSPGVS200000000	sp Q9UKV3 ACINU_HUMAN
7198	LSPERSSDSESNENEPHLPHGVSQR2.0002202E+24	sp Q9BQ52 RNZ2_HUMAN
7199	LLDPSSPPTK200000	sp Q6IAA8 LTOR1_HUMAN
7200	ADTSQEICSPR2000	sp P52948 NUP98_HUMAN
7201	KASGSENEGDPNPR200000000000	sp Q02880 TOP2B_HUMAN
7202	SDAEEDGGTVSQEEEDRKPK2E+20	sp P27824 CALX_HUMAN
7203	TQPDGTSVPGEPASISQR200000000000000	sp Q14980 NUMA1_HUMAN
7204	SISLSQSAENVPASK200000000000000	sp Q4ADV7 RIC1_HUMAN
7205	GNSPPSSGEACR200000000000	sp P41227 NAA10_HUMAN
7206	LSVHDMKPLDSPGR20000	sp Q15643 TRIPB_HUMAN
7207	ASSEGTAAGAGLDSLHK20000000000000000	sp O15143 ARC1B_HUMAN
7208	HSASPVVFTSAR202000002000	sp O43166 SI1L1_HUMAN
7209	SDGACDSPSSDKENSSQIAQDHQK2000000000000000000	sp Q9UIG0 BAZ1B_HUMAN
7210	DHGFVASTSPEDESPGSNPEPDATPFQEGLR2E+24	sp P55211 CASP9_HUMAN
7211	EAGLSQSHDDLSTATATPSVR20000000000000000	sp O14523 C2C2L_HUMAN
7212	QQLEEEAAKPEPEKPVSPPIESK2000000000	sp Q9Y618 NCOR2_HUMAN
7213	IILPDDSSSEENNVNDK220000000000	sp Q8IYD8 FANCM_HUMAN
7214	LTQAEISEQPTMATVVPVQVPTSPK2000	sp Q70E73 RAPH1_HUMAN
7215	DTYSDRSGSSPDSEITELK2002022000000000000	sp P17812 PYRG1_HUMAN
7216	AAEETPGKSPEK200020000	sp O76021 RL1D1_HUMAN
7217	NSPLDCGSASPKNK200000000000000	sp O75179 ANR17_HUMAN
7218	SSDSESNENEPHLPHGVSQR2.02E+19	sp Q9BQ52 RNZ2_HUMAN
7219	APSPPSR200000	sp Q7Z2K8 GRIN1_HUMAN
7220	SSPEQPRPSTSK200000000000	sp P29590 PML_HUMAN
7221	SVWGLAVQNSPK2000	sp P18615 NELFE_HUMAN
7222	VEMYSGSDDDDFNKLK200000000000000	sp O60841 IF2P_HUMAN
7223	TSIETNIR20000000	sp P14859 PO2F1_HUMAN
7224	VSPSPQESLSSSK202000000000000	sp Q9UKI9 PO2F3_HUMAN
7225	LFEDDSNEKLFDEEEDSSEK2000000000022000	sp Q5JSH3 WDR44_HUMAN
7226	EAPGSPPLSPR20002000	sp O43719 HTSF1_HUMAN
7227	SCSSPAVSAVSQLPLSPK200000002000	sp Q86UU0 BCL9L_HUMAN
7228	ASGDPGASGMSPR2000	sp Q9Y2F5 ICE1_HUMAN
7229	FRDQDLASCDR20000	sp Q8NDT2 RB15B_HUMAN
7230	YLVDTGKPNAGSEEISSEDELVEEK2.0000000022E+21	sp Q9Y383 LC7L2_HUMAN
7231	KAPAGQEEPPTPSSPLSAEQQLDR200200000000000	sp Q9NY61 AATF_HUMAN
7232	CSDNSSYEEPLSPISASSSTR200000000000	sp P13051 UNG_HUMAN
7233	SPSFGAGEGLLR200000000000	sp Q8IXK0 PHC2_HUMAN
7234	KPPDSYPPK200000	sp Q96PE2 ARHG_HUMAN
7235	SCSSPAVSAVSQLPLSPK2000000000002000	sp O43184 ADA12_HUMAN
7236	SDGACDSPSSDKENSSQIAQDHQK22000000000000000	sp Q9Y2F5 ICE1_HUMAN
7237	HSWGPGK2000000	sp Q9UIG0 BAZ1B_HUMAN
7238	YSPSPPK20200000	sp Q12802 AKP13_HUMAN
7239	IIPITLLEGLQLPSPTATSQLPLESDAVECLNYQHYK2E+24	sp Q8IYB3 SRRM1_HUMAN
7240	EWNGVSESDSPVK20000	sp P61978 HNRPK_HUMAN
7241	NSPGSQVSNPR200000020000	sp P18858 DNLI1_HUMAN
7242	SPPGAEEPQAPPASR20000000000000000	sp P18858 DNLI1_HUMAN
7243	KRPSLPSSPGLPK2000202000000	sp Q9HOD6 XRN2_HUMAN
7244	TSISDPPESPPLLPPR200002000000000	sp Q86VR2 F134C_HUMAN
7245	SPSSKPSLAVAVDSAPEVAVK2E+19	sp O43294 TGFI1_HUMAN
7246	QIASDSPHASPK20002000	sp Q07889 SOS1_HUMAN
7247	EGKPEIEGKPESEGEPSGSETR200000000000	sp O60508 PRP17_HUMAN
7248	ESISPPADSACSPPSTGK2.02E+20	sp Q9HCD6 TANC2_HUMAN
7249	FGSTGTPPVSPTPSER22000200000000	sp Q96EI5 TCAL4_HUMAN
		sp Q9GZR1 SENP6_HUMAN
		sp Q969V6 MKL1_HUMAN



7250	NWTEDMEGGISSPVKK200000	sp P08651 NFIC_HUMAN
7251	YWEETVPTTAGASPGPPR2000000	sp O15213 WDR46_HUMAN
7252	IFDEVMGCFSDSPQSPTFPEAGHTSLYDEDK2.02E+23	sp Q13085 ACACA_HUMAN
7253	AWLDEDSNLSPLR2020000	sp Q9C086 IN80B_HUMAN
7254	STSAPQMSPGSSDNQSSSPQPAQK1.20022E+19	sp Q14157 UBP2L_HUMAN
7255	KRSPSPSPTPEAK200020000000	sp Q8TAQ2 SMRC2_HUMAN
7256	VVDYSQFQESDDADEDYGR300000000000000000	sp Q9H1E3 NUCKS_HUMAN
7257	SPSPALHPPQK200000000000	sp Q9H6S0 YTDC2_HUMAN
7258	SLRGSDALSETSSVSHIEDLEK2000202200000000000	sp Q6P996 PDXD1_HUMAN
7259	KPHDCESSTVSEEDYFSSHR2220000000000000	sp Q9H4H8 FA83D_HUMAN
7260	TQSAPLPQNAQALQHLVIQQHQHFLEK2E+26	sp P56524 HDAC4_HUMAN
7261	GYYSYPVSVSGSGTAGSR30000000000200	sp Q15149 PLEC_HUMAN
7262	SGYSSPGSPGTPGSR2000200000000	sp TAU_HUMAN
7263	HSLSGSSPGMK20200200000	sp P10636 TAU_HUMAN
7264	SAPASPNHAGVLSAHSSGAQTPELSR2.0002E+27	sp Q9UQ35 SRRM2_HUMAN
7265	QLVAGNSPK2000	sp Q01167 FOXK2_HUMAN
7266	SILKPSTPIPPQEGEEVGESSEEQDNAPK20000000000	sp Q96EU6 RRP36_HUMAN
7267	DGDSYDPYDFSDTEEMPQVHTPK2.000000020001E+21	sp Q9UDY2 ZO2_HUMAN
7268	RTSVSPPEQPYPYR2002000000000	sp P25205 MCM3_HUMAN
7269	TASESISNLSEAGSIKK200020000	sp Q8TEH3 DEN1A_HUMAN
7270	HNSESEVPSSMFILEDDR2000000000000000000	sp P30622 CLIP1_HUMAN
7271	RTFSLTEVR202000000	sp Q6R327 RICTR_HUMAN
7272	RGLRDSHSSEDEASSQTDLSTISK2.022E+21	sp O95239 KIF4A_HUMAN
7273	VSKSPSPASTSTVPNMTDAPTAPK2.0020000001E+19	sp Q5JTV8 TOIP1_HUMAN
7274	SRTPPVTR202000000	sp Q659A1 ICE2_HUMAN
7275	DYDDMSPR2000	sp Q9UQ35 SRRM2_HUMAN
7276	RLSVGSSMR20002000	sp P61978 HNRPK_HUMAN
7277	APSPLGPTR20000000	sp Q8N370 LAT4_HUMAN
7278	VHVQFFDDSPTR200	sp Q0VG06 FP100_HUMAN
7279	KGGSYSQAASSDSAQGSVDVSLTACK2000000000	sp P52701 MSH6_HUMAN
7280	SSSVTTSETQPTPSSSDYSDLQR20000000000000000	sp P01891 1A68_HUMAN
7281	KAPSASDSKADSDGAKPEPVAMAR2.02020002E+21	sp P10316 1A69_HUMAN
7282	HGGSPQLATTPLSQEPVNPPEASPTR2000000020000000	sp P01892 1A02_HUMAN
7283	SRSPGRPPK2020000000	sp P05534 1A24_HUMAN
7284	FNISSHNQSPK2000	sp P30447 1A23_HUMAN
7285	SPTPPLFSLPEAR200000000000	sp Q09160 1A80_HUMAN
7286	RPAVLPLELVGEGSNNTSTDGSLPSTPPPAEEEEDELYR200000000000000	sp P50552 VASP_HUMAN
7287	RKSQDLSLK20000000	sp Q7Z4V5 HDGR2_HUMAN
7288	KESESDSSDDEPLIK2020002000000000	sp Q9UQ35 SRRM2_HUMAN
7289	SSSPTQYGLTK20000000000	sp Q14739 LBR_HUMAN
7290	CRSPGMLEPLGSSR2000000000000	sp Q9H1A4 APC1_HUMAN
7291	KRESESEDETTPAAPQLIK20000000000000000	sp Q96FV9 THOC1_HUMAN
7292	EVSSRPSTPGLSVVSGISATSEDI PNK20000000000000000	sp Q07820 MCL1_HUMAN
7293	QSMDSPIK20000	sp Q14693 LPIN1_HUMAN
7294	LEKSPLAGNK20000000	sp P35659 DEK_HUMAN
7295	KNATQPNAEEESGSSASEEEDTKPKPTK2022202000000000000	sp Q9NYL2 MLTK_HUMAN
7296	EPALPLEPDSGGNTSPGVTANGEAR2000000000000	sp Q9UQ35 SRRM2_HUMAN
		sp O60832 DKC1_HUMAN
		sp Q14C86 GAPD1_HUMAN
		sp Q14562 DHX8_HUMAN
		sp O75151 PHF2_HUMAN
		sp Q68E01 INT3_HUMAN
		sp Q92974 ARHG2_HUMAN

7297	ISMQDVDSLGLGSPK100000002000	sp Q09666 AHNK_HUMAN
7298	NAASFPLRSPQVPCSPAGSEGTK20000020000000000	sp Q6KC79 NIPBL_HUMAN
7299	QFTSSSSIKGSSGLGGSSR22000000000000000	sp Q04695 K1C17_HUMAN
7300	SPTLSQVHSPLVTSPSANLK20002000200000000	sp Q86UU0 BCL9L_HUMAN
7301	SGDETPGSEVPGDK200000020000000	sp Q1KMD3 HNRL2_HUMAN
7302	VVLGDGVQLPPGDYSTTPGGTLFSTTPGGTR20000000000000000	sp Q13541 4EBP1_HUMAN
7303	FSPGAPGGSGSQPNQK20000000000000000	sp Q15942 ZYX_HUMAN
7304	LKDLFDYSPPLHK2000000	sp Q9NYF8 BCLF1_HUMAN
7305	RGSQNSSEHRPPASSTSEDVK2E+19	sp O15014 ZN609_HUMAN
7306	SPSTTYLHTPTSEDAAIIPSK2200000000000000000	sp Q13111 CAF1A_HUMAN
7307	ATSPSSVSGDFDDGHHSVSTPGPSR2. 2E+21	sp Q86U86 PB1_HUMAN
7308	SPARTPPSEEDSAEAER2000200000000000000	sp O43765 SGTA_HUMAN
7309	ASLVALPEQTASEEETPPPLTK200000000	sp Q9UKV3 ACINU_HUMAN
7310	ERDHSPTPSVFNSEER20000000200000	sp Q6UN15 FIP1_HUMAN
7311	GISPVVSEHR200000000	sp Q99700 ATX2_HUMAN
7312	VYESECTDSEELNMK202020200000000	sp Q15054 DPOD3_HUMAN
7313	TDSVIIADQTPPTR2000000002000000	sp P17544 ATF7_HUMAN
7314	AVEESSDEER2200000	sp Q9BW71 HIRP3_HUMAN
7315	KPSDGSPTKPSR200200000000	sp Q92733 PRCC_HUMAN
7316	ADLNQIGEPQSPSR200	sp Q96C19 EFHD2_HUMAN
7317	VPASPLGLER200000000	sp Q7KZI7 MARK2_HUMAN
7318	DGDSYDPYDFSDTEEMPQVHTPK30000200000000000	sp P25205 MCM3_HUMAN
7319	GPSPTSPTPAAAAAPAEQAPR20000000000000000	sp Q86VQ1 GLC11_HUMAN
7320	LDQPVSAAPPSPR20000000	sp Q16204 CCDC6_HUMAN
7321	TVSASSTGDLPK2020200000000	sp P78362 SRPK2_HUMAN
7322	RSPSPPPTR202000000	sp Q8IYB3 SRRM1_HUMAN
7323	SATFPNAGPR20000000000	sp Q86X27 RGPS2_HUMAN
7324	SSSPVTELASR220000000000	sp Q9UQ35 SRRM2_HUMAN
7325	SPACSSLTPSLCK20002000000000	sp Q96R06 SPAG5_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
		sp Q29960 1C16_HUMAN
7326	GGSCSQAASSNSAQGSDESLIACK2000200000000	sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
7327	ETNLDSLPLVDTHSK20000000000	sp P08670 VIME_HUMAN
7328	TDGSISGDRQPVTVADYISR2. 002E+20	sp P51116 FXR2_HUMAN
7329	TLTDEVNSPDSRRR20020000	sp Q8TAQ2 SMRC2_HUMAN
		sp Q09160 1A80_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
7330	GGSYSQAASSDSAQGSVSLTACK2000000000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
7331	GIPHSASPVSPDGVQIPLK20200200000000000	sp Q6PJF5 RHDF2_HUMAN
7332	SYSPDGKESPSDKK32000002000000	sp P43243 MATR3_HUMAN
7333	SGDEGSEDEAPSGED2000020000020000	sp Q9H1E3 NUCKS_HUMAN
7334	HNDIVSDSDAEDRGTLSAELTAAHFGGGGLLHK2. 02E+20	sp P78316 NOP14_HUMAN
7335	SPTSSAIPLQSPR20000000002000	sp Q9ULD2 MTUS1_HUMAN

7336	SILKPSTPIPPQEGEEVGESSEEQDNAPK22000000000	sp	Q9UDY2	ZO2_HUMAN
7337	VDEDSAEDTQSNKGKEVVEVGQK2. 000002E+19	sp	Q9H2G2	SLK_HUMAN
7338	KPIDSLRDSRSLSYSPVER202003200000	sp	Q9UQ35	SRRM2_HUMAN
7339	AVSPAPQSSSR2000000000	sp	Q9P275	UBP36_HUMAN
7340	NSASATPLSGNSSR200	sp	Q9Y608	LRRF2_HUMAN
7341	RPLSSSHEASEGQAK200000000000	sp	Q8IZ21	PHAR4_HUMAN
7342	LEGPPEANADPLGVLINSDSESDKEEKQHSVIPK2020200000000000000	sp	Q9UHK0	NUFP1_HUMAN
7343	LNETELTDLEGQQESPPKNYLCIEEEK20000000000000	sp	Q9ULF5	S39AA_HUMAN
7344	ENSPAVSPTTNSTAPFGLKPR2. 0000002E+19	sp	Q9Y2L9	LRCH1_HUMAN
7345	RESEDDPESEPDDHEK2000002000000000	sp	P16383	GCFC2_HUMAN
7346	RGEGDAPFSEPGTTSTQRPSPETATK2000200000000	sp	Q9UQ35	SRRM2_HUMAN
7347	CAAPRPPSSSPEQR200000	sp	Q5H9R7	PP6R3_HUMAN
7348	QLPALDGLMGPESPPAQEEEEAPVSPHK1. 0002E+19	sp	Q8N9T8	KRI1_HUMAN
7349	SPMSTNSSVHTGSDVEQDAEK2. 000000000002E+21	sp	Q9ULU4	PKCB1_HUMAN
7350	YSPTSPTYSPPTPK20000000000	sp	P24928	RPB1_HUMAN
7351	SAFTPATATGSSPSPVLGQGEK200000000000	sp	Q15811	ITSN1_HUMAN
7352	MSESPTPCSGSSFEETEALVNTAAK2E+22	sp	O95359	TACC2_HUMAN
7353	TTSRSPVLSR202000000	sp	O95819	M4K4_HUMAN
7354	EALAEAALESPPALVR200000000	sp	O14745	NHRF1_HUMAN
7355	SASADNLTLP200000000000	sp	Q8ND76	CCNY_HUMAN
7356	GSISSTSEVHSPPNVGLR2000000000000000	sp	Q8WWQ0	PHIP_HUMAN
7357	GEVISSGSKPLSPGPCLDIPSILSNK2. 000002E+21	sp	Q8TEP8	CE192_HUMAN
7358	NQSHSSPSVSPSR200020000	sp	Q7Z6E9	RBBP6_HUMAN
7359	KRSPTESVNTVPVGK2000000000000	sp	Q8WXG6	MADD_HUMAN
7360	KPPAPSPVQSQSPSTNWSPAVPVK2. 000000022002E+19	sp	Q8IYB3	SRRM1_HUMAN
7361	KEDDSSQSSTCVVDTTTR20222000000000	sp	Q5SNT2	TM201_HUMAN
7362	SWGHEPSEER20000200000	sp	Q9UKJ3	GPTC8_HUMAN
7363	SRASPVSR200200000	sp	Q9UQ35	SRRM2_HUMAN
7364	APYSPTTSK2000000	sp	Q96JH7	VCIP1_HUMAN
7365	LEQTSVRDPSPEADAPVLGSPEKEEAASEPPAAAPDAAPPPDRPIEK2. 0000 0002E+29	sp	O15234	CASC3_HUMAN
7366	EECGDTALSSDEEDLSSWLPSSPAR2. 002E+20	sp	Q6P4R8	NFRKB_HUMAN
7367	VKAQTPPGPSLSGSKSPCPQEK2000000000020000000	sp	Q9UQ35	SRRM2_HUMAN
7368	TPLSFTNPLHSDDSDSDER2000000	sp	Q05209	PTN12_HUMAN
7369	RMSPKPELTEEK200000000000	sp	P41208	CETN2_HUMAN
7370	KTEAPAAPAAQETKSDGAPASDSKPGSSEAAPSSK2E+23	sp	P80723	BASP1_HUMAN
7371	LDAEPKPCSDTNSDSPDEGLDHNHIK2. 020202E+19	sp	Q13129	RLF_HUMAN
7372	MEDLDQSPLVSSSDSPPRPQPAFK2000000020000000000	sp	Q9NQC3	RTN4_HUMAN
7373	YSQGGDDGSSSSGSSVAGSQSTLFLK2000000000000	sp	Q9UH99	SUN2_HUMAN
7374	GNKSPSPPDGSPAATPEIR20000000000000000	sp	O00499	BIN1_HUMAN
7375	ITETGAVKPTGECSGEQSPDTNYEPPGEDK2E+21	sp	P49454	CENPF_HUMAN
7376	DSPTLSNSGIR20000000000	sp	Q9HD20	AT131_HUMAN
7377	KSPAKSPNPSTPR2000200020000	sp	O76021	RL1D1_HUMAN
7378	LSSDENSNPDLSGDENDDAVSIESGTNTERPDTPTNTPNAPGR20000200000 000000	sp	O75530	EED_HUMAN
7379	SLPTTVPESPNYR20000200000	sp	Q6PKG0	LARP1_HUMAN
7380	DGDSYDYPDFSDTEEMPVHTPK300000000100000000	sp	P25205	MCM3_HUMAN
7381	QEAIPDLEDSPPVSDSEEQESAR2000002000000000	sp	Q9BTC0	DIDO1_HUMAN
7382	ATAPQTQHVSPMR20100	sp	P29692	EF1D_HUMAN
7383	SPSVSPKQPVSTSSK2020000000000	sp	Q96JG6	CC132_HUMAN
7384	KSPSGPVKSPPLSPVGTTPVK2000000002000000000	sp	Q9BVC5	ASHWN_HUMAN
7385	KRESESEDETTPAAPQLIK2000200000000000000	sp	O60832	DKC1_HUMAN
7386	HSSNPPLSHVGVWMSR20000000000010000	sp	Q6PKG0	LARP1_HUMAN

7387	KYSASSGGLCEEATAAK200000000000	sp Q9UHV7 MED13_HUMAN
7388	SPQTPSQMVPLPSANPPGPK202010000000000000	sp Q86UU0 BCL9L_HUMAN
7389	ASLSSAPVVLVDHA2000000000000000	sp Q9HBH9 MKNK2_HUMAN
7390	TRDDGDEEGLLTHSEEELEHSQDTDADDGALQ2.000000002E+21	sp Q9UKM9 RALY_HUMAN
7391	YSKPTAPAPSAPPSPSAPEPPK20002000000000	sp P55198 AF17_HUMAN
7392	SSDQDSTNKEAEAAGVKPAGVRPR2.20002E+24	sp Q92625 ANS1A_HUMAN
7393	LTSVSSSVDFDQR2200000000	sp Q5T3F8 CSCL2_HUMAN
7394	VDSPTVTTLK2000000000	sp Q07866 KLC1_HUMAN
7395	VVPQQIHTSPR2000	sp Q9H0E3 SP130_HUMAN
7396	IADPEHDHTGFLTEYVATR20000000	sp P27361 MK03_HUMAN
7397	THCVGDSQSSASSPPATSK20000000	sp Q9H4Z2 ZN335_HUMAN
7398	HGSGADSDYENTQSGDPLLGLEK3002000000000000	sp Q9Y2X7 GIT1_HUMAN
7399	SKGSLEITESQSADAEPKPPKPDLSR2E+27	sp Q9C0B5 ZDHC5_HUMAN
7400	SFSISPVR2020000	sp P18583 SON_HUMAN
7401	NKIEEAPEATPQSPQPGSSPISLSAEENAEGEVS2020000000000200	sp Q9H4L7 SMRCD_HUMAN
7402	VDDKPSSPGDSSK20000000	sp Q9HC52 CBX8_HUMAN
7403	SQSPAASDCSSSSSSASLPSSGR20000000000000	sp O95817 BAG3_HUMAN
7404	LFPEFDDSDYDEVPEEGGAPAR3000000000000000	sp Q96P48 ARAP1_HUMAN
7405	GEGDIHENVDLPGSLGQSEEKVPVAPVPSVAPVPSR200000000000	sp Q9UK76 HN1_HUMAN
7406	ASPEPQRENASPAGTTAEAMSR2.000000002E+23	sp P46379 BAG6_HUMAN
7407	TEARSSDEENGPPSSPDLDR20000000022000000	sp Q96B36 AKTS1_HUMAN
7408	LGAGGSPEKSPSAQELK200000000	sp Q9UNE7 CHIP_HUMAN
7409	TEPHSDCSVDLGISK200200000000	sp Q96RT1 LAP2_HUMAN
7410	RPDYAPMESSDEEDEFQFIKK102200000000000000	sp P55081 MFAP1_HUMAN
7411	SVSHGSNHTQKPDEQR200000000	sp Q9Y520 PRC2C_HUMAN
7412	TRDDGDEEGLLTHSEEELEHSQDTDADDGALQ2.020000000002E+21	sp Q9UKM9 RALY_HUMAN
7413	YFDSGDYNMAK200000000	sp P56211 ARP19_HUMAN
7414	VEQMPQASPLAPR100020000000	sp O43768 ENSA_HUMAN
7415	TVSPGSVSPIHGQGVVENLK2000000000000000	sp Q8N3F8 MILK1_HUMAN
7416	FGEYNSNMSPEEK200000	sp Q9NZM3 ITSN2_HUMAN
7417	ESPTQISIGAFPSTK2000000000000000	sp P78316 NOP14_HUMAN
7418	SPSPDGSPAATPEIR202000000000000000	sp Q5VWN6 F208B_HUMAN
7419	VKGGDDHDDTSDSDGLTLK22020000000000	sp 000499 BIN1_HUMAN
7420	AKPVVSDFDSDEEQDER200000000	sp Q9BTC0 DIDO1_HUMAN
7421	VDKETNTETPAPSPTVVRPK2000002000000000	sp P51531 SMCA2_HUMAN
7422	TAVQYIESSDSEIETSELPQK2002000000000000	sp Q8IX03 KIBRA_HUMAN
7423	PGPTPSGTNVGSSGRSPSK20020000	sp Q14527 HLTF_HUMAN
7424	QQLSPSPGQEAGILPETEK200000000000000000	sp P60468 SC61B_HUMAN
7425	GSSPGPRPVEGTPASR200000000000000000	sp Q17RY0 CPEB4_HUMAN
7426	DRPVSQPSLVGSK2000000000	sp Q13112 CAF1B_HUMAN
7427	HSSISPVR2020000	sp Q9H4A3 WNK1_HUMAN
7428	TTPATGEQSPGAR200000	sp Q9NYV4 CDK12_HUMAN
7429	ACRPPGSPGR20000	sp 000308 WWP2_HUMAN
7430	SRPTSFADELAAR2000000000	sp O95685 PPR3D_HUMAN
7431	GGIDNPAITSDQELDDKK20000000000	sp Q641Q2 FA21A_HUMAN
7432	GSVPLSSPPATHFPDETEITNPVVK2.0002E+25	sp Q5SNT6 FA21B_HUMAN
7433	DHSPTPSVFNSEER200000200000	sp Q9Y4E1 FA21C_HUMAN
7434	GEGDAPFSEPGTTSTQRPSPETATK200000000	sp Q13017 RHG05_HUMAN
7435	SKPPPTYESEEDK2000000000	sp P11388 TOP2A_HUMAN
		sp Q6UN15 FIP1_HUMAN
		sp Q9UQ35 SRRM2_HUMAN
		sp O60885 BRD4_HUMAN

7436	TSRDTSPSSGSAVSSSK2200000000000	sp Q8NEY8 PPHLN_HUMAN
7437	SDSELEVKPAESLLR2000000000000	sp Q92539 LPIN2_HUMAN
7438	TDSREDEISPPPPNPVVK2000002000000000	sp P10644 KAPO_HUMAN
7439	SGPKPFSAPKPKTSPSPKR2020000	sp Q01518 CAP1_HUMAN
7440	SSGHSSSELSPDAVEK22002200020000000	sp Q9UQ35 SRRM2_HUMAN
7441	ILDELDKDDSTHESLSQESESEEDGIHVDSQK2.0200202E+19	sp Q9H6T3 RPAP3_HUMAN
7442	SIEDDEEGLICQSGDVL2E+19	sp Q9HAZ1 CLK4_HUMAN
7443	TVQSPNSSVSPGLAGPVTMTSVHPPIR2.002E+21	sp O15164 TIF1A_HUMAN
7444	ASAVSELSR2002002000	sp Q9Y2W1 TR150_HUMAN
7445	LGLQEGSNSSPVDFVNNK20002000000000	sp Q96R06 SPAG5_HUMAN
7446	HGSSDISSPR200002000	sp Q9BRD0 BUD13_HUMAN
7447	ILIPSDIESSK20000000	sp P30414 NKTR_HUMAN
7448	ALEAASLSQHPPSLCISDSEEEEEER200020200000000	sp O43159 RRP8_HUMAN
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		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
		sp A6NKT7 RGPD3_HUMAN
7450	DSLITPHVSR2000200	sp PODJD0 RGPD1_HUMAN
		sp PODJD1 RGPD2_HUMAN
		sp O14715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
7451	TSSKESPIPSPTSDRK220000000000020000	sp Q01082 SPTB2_HUMAN
7452	HGGSPQLATTPLSQEPVNPSEASPTR2.000002E+25	sp Q9UQ35 SRRM2_HUMAN
7453	STAQQELDGKASPPTPVIVASHTANKEEK2E+28	sp P35606 COPB2_HUMAN
7454	STSAPQMSPGSSDNQSSSPQAQQK2200000000	sp Q14157 UBP2L_HUMAN
7455	AESPESAIESTQSTPQK2000000000000	sp Q9NTI5 PDS5B_HUMAN
7456	TSPENVQDR200000000	sp P49792 RBP2_HUMAN
7457	SSLGQSASETEEDTVSVSK2000200000000000000	sp Q13435 SF3B2_HUMAN
7458	ALPSLNTGSSSPR200000	sp O14545 TRAD1_HUMAN
7459	TSSSETEEKKEKPLILQSLGPK2.2E+22	sp Q9Y5K6 CD2AP_HUMAN
7460	YSPTSPYSPPTPK20000000002000	sp P24928 RPB1_HUMAN
7461	ADTSQEICSPR200002000	sp P52948 NUP98_HUMAN
7462	RNSDRSEDGYHSDGDYGEHDYR2.0020003E+20	sp P52756 RBM5_HUMAN
7463	SAASPVVSSMPER2000000000000	sp P18583 SON_HUMAN
7464	KPSPSEPEPWKPFPAVSPEPR2.0002E+20	sp Q96JM3 CHAP1_HUMAN
7465	FDIYDPFHPTDEAYSPPPAPQK200000000000000	sp Q9H7N4 SFR19_HUMAN
7466	SSSSLGSGSSRDSISSDSETSEPLSCR2.002E+24	sp O75164 KDM4A_HUMAN
7467	GVQVPASPDTVPQPSLR2000000000000	sp Q92613 JADE3_HUMAN
7468	ASSARHSESDITSLIEAMDK2.00022002E+20	sp Q92545 TM131_HUMAN
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7470	RQSSGSATNVASTPDNR2020000000000	sp Q7Z460 CLAP1_HUMAN
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7472	SEPPAELPSLR20000000000	sp Q14669 TRIPC_HUMAN
7473	SNSFSDER200000000	sp Q09666 AHNK_HUMAN
7474	LEGNSPQGSNQGVK2000000	sp P61006 RAB8A_HUMAN
7475	TGSGSPFAGNSPAR200000000020000	sp Q9UPT8 ZC3H4_HUMAN
7476	DHTVSGDEDYCP2000000000	sp Q9UIG0 BAZ1B_HUMAN
7477	RGSPAPANPGR2000000000	sp O75676 KS6A4_HUMAN
7478	SPSPPTIQSK20200000000	sp Q14687 GSE1_HUMAN
7479	DRPHASGTDGDESEEDPPEHKPSK2.0200002E+19	sp Q96RS0 TGS1_HUMAN
7480	AYSPEYR200000	sp Q9UDY2 ZO2_HUMAN
7481	SSLGQSASETEEDTVSVSK2020000000000	sp Q13435 SF3B2_HUMAN
7482	EKPELSEPHLNGPSSDPEAAFLSR2.000000002E+20	sp Q9H4A3 WNK1_HUMAN
7483	IPSEMPSPNFGK102000000	sp Q9H2J7 S6A15_HUMAN

7484	SRSGEGEVSGLMR2000000000100	sp Q13263 TIF1B_HUMAN
7485	YESQLSTNEEKVDTDDR200000000000000	sp Q13433 S39A6_HUMAN
7486	SSQPSPTAVPASDSPPTK22000000000000000	sp Q3KQU3 MA7D1_HUMAN
7487	SGGGSGGSGGVSLK20000000000000000	sp Q9H7N4 SFR19_HUMAN
7488	SPPLPAVIR2000000000	sp Q9Y6X9 MORC2_HUMAN
7489	TSPADHGGVSGSESGSAVDSVAGEHSVSGR2E+20	sp Q5T4S7 UBR4_HUMAN
7490	SLLSHEFQDETDTTEEETLYSSK20000000000	sp 075976 CBPD_HUMAN
7491	VDSSTNSSPSPQQSESLSPAHTSDFR200000000000	sp Q9BQE9 BCL7B_HUMAN
7492	EPVSPMELTGPEDGAASSGAGR2E+19	sp 094953 KDM4B_HUMAN
7493	DDRSDVESSEEDVTCTK200022200000000000	sp Q8N5B7 CERS5_HUMAN
7494	YSDTTSDPENEPFDEDQHTQITK2.02202E+23	sp P60484 PTEN_HUMAN
7495	HGLLLPASPV20000	sp Q96E09 F122A_HUMAN
7496	SSPPLRTPDVLESSGPAVR2.000002E+19	sp 075676 KS6A4_HUMAN
7497	VVDYSQFQESDDADEDYGR200000000000000	sp Q9H1E3 NUCKS_HUMAN
7498	MVIQGPSSPQGEAMVTDVLEDQK1.000002000001E+23	sp Q12888 TP53B_HUMAN
7499	KSPAQSPNPSTPR2000200002000	sp 076021 RL1D1_HUMAN
7500	RLSGSSEDEEDSGKGEPTAK202200000200000000	sp Q9BW71 HIRP3_HUMAN
7501	QFTSSSSIKGSSGLGGSSR22000000000000000	sp Q04695 K1C17_HUMAN
7502	VSEEQTPPSPAGAGMSTAMGR200000000010001000	sp Q16666 IF16_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
7503	EDALDDSVSSSSVHASPLASSPV2000200000	sp 014715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
		sp A6NKT7 RGPD3_HUMAN
7504	KITPPSSPHSLTSTLVSLGHEEALEMAGSK2.2E+25	sp Q9BY89 K1671_HUMAN
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7506	SEPSLEPESFR200200000000	sp Q9C0B5 ZDHC5_HUMAN
7507	GARPPAAGPGDEDEDEEDTAPESALDTSCLK20000000020000	sp Q27J81 INF2_HUMAN
7508	LSSLRASTK200000000	sp P62753 RS6_HUMAN
7509	SPEAKPLPGK20000000000	sp P46087 NOP2_HUMAN
7510	ELDDSSSESSDLQLEGSTLR22020000000000000	sp P29590 PML_HUMAN
7511	TAPSSPLTSPSDTR20020000000	sp Q8IWB9 TEX2_HUMAN
7512	GPSWDPFR2000000	sp P04792 HSPB1_HUMAN
7513	RPGHGSLTNISR20000000	sp Q7Z6B7 SRGP1_HUMAN
7514	STPFIVPSSPTEQGR200000000	sp Q12888 TP53B_HUMAN
7515	SLTSSLENIFSR200000000	sp 060343 TBCD4_HUMAN
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7517	SPQTPSQMVPLPSANPPGPLK20200000000000000	sp Q86UU0 BCL9L_HUMAN
7518	DSALQDTDDSDDDPVLIPGAR2.00000002E+20	sp Q58WW2 DCAF6_HUMAN
7519	EISSSPESCPEEIK2020020000000	sp Q92889 XPF_HUMAN
7520	SPTPPSSAGLGSNSAPPIDSR202000000000	sp Q8IWX8 CHERP_HUMAN
7521	RHSHSHSPMSTR202000000	sp P62995 TRA2B_HUMAN
7522	SSLSPSSAYER2002000000000	sp Q9BYW2 SETD2_HUMAN
7523	FRRSETPPHWR202000000	sp Q13427 PPIG_HUMAN
7524	NDSVIVADQTPPTR20000002020000	sp P15336 ATF2_HUMAN
7525	ERDHSPTPSVFNSEER20200000000000	sp Q6UN15 FIP1_HUMAN
7526	KYSISSDNSDTTDSHATSTSASR2.2002E+19	sp Q9NQC1 JADE2_HUMAN
7527	ILDELKDDSTHESLSQESESEEDGIHVDSQK2.02002E+19	sp Q9H6T3 RPAP3_HUMAN
7528	SLSGSSPCPK20002200000	sp Q9UQ35 SRRM2_HUMAN
7529	LRLSPSPTSQR2002000	sp P02545 LMNA_HUMAN
7530	VQSSSESTSSPSQHEATPGAR2000000000000	sp 075152 ZC11A_HUMAN
7531	SVDFDSLTVR20000000000	sp Q9Y5K6 CD2AP_HUMAN
7532	VESGDPSSLHPPSPGTPSATPASPPASATPSQR200200000000000	sp Q4KMQ1 TPRN_HUMAN
7533	GLPLGSAVSSPVLFPVGR20000200000	sp Q8WUM0 NU133_HUMAN

7534	ITPHTSPAPR2002000000	sp Q58A45 PAN3_HUMAN
7535	VNPHKVPASSVDSNIPSSQGYK200020000000000000	sp Q9UHB7 AFF4_HUMAN
7536	SPLHSSPEK20000	sp Q13459 MYO9B_HUMAN
7537	APEPDLMSPTR20000	sp Q9H7D0 DOCK5_HUMAN
7538	LSVQSNPSPQLR200000	sp 075152 ZC11A_HUMAN
7539	ESAAPASPAPSPAPSPTAPPQK20002000000000	sp Q3KQU3 MA7D1_HUMAN
7540	ERSPSPSFR20200000	sp P49585 PCY1A_HUMAN
7541	SPSTTYLHTPTSEDAAIPSK2. 2E+19	sp Q13111 CAF1A_HUMAN
7542	SATDGNTSTPPTSAK200000000	sp Q13620 CUL4B_HUMAN
7543	SPLHSSPEK220000	sp Q13459 MYO9B_HUMAN
7544	RTDALTSPPGR20002220000	sp P49736 MCM2_HUMAN
7545	SPVSSLQIR2000000000	sp P55197 AF10_HUMAN
7546	EVPWSPSAEK2000000	sp Q5JSZ5 PRC2B_HUMAN
7547	GAPSSPATGVLSPQ GK200000	sp Q5SRE5 NU188_HUMAN
7548	EASRPPEEPSAPSPTLPAQFK2000000000	sp Q9H3P2 NELFA_HUMAN
7549	SPSFGEDYYGPSR200000000000	sp P49761 CLK3_HUMAN
7550	NSDRSEDGYHSDGDYGEHDYR2. 002000002E+20	sp P52756 RBM5_HUMAN
7551	AGGPTTPLSPTR200020000	sp P20700 LMNB1_HUMAN
7552	LTPVSPVQHQGATVNNTNK20020000000000000000	sp Q2KHR2 RFX7_HUMAN
7553	NMVDLVNTHHLHSSSDEDDR1. 0000000000002E+20	sp Q9UPN7 PP6R1_HUMAN
7554	GTPPLTPSDSPQTR20000000200000	sp Q6NZY4 ZCHC8_HUMAN
7555	KASPEPEGEAAGK200000000000	sp Q8IU81 I2BP1_HUMAN
7556	RRSSPSARPPDVPGQPAAK20200000000000000000	sp Q96JP5 ZFP91_HUMAN
7557	QQDLHLESPQRQPEYSPESPR2000000000000000	sp Q9H6F5 CCD86_HUMAN
7558	NKQDDDLNCEPLSPHNITPEPVSK20000000000000	sp Q6VMQ6 MCAF1_HUMAN
7559	KPVTVSPTTPTSPTGEAS22020000000000	sp Q9Y6G9 DC1L1_HUMAN
7560	YVESDDEKPLDDETVNEDASNENSENDITMQSLPK2E+32	sp P46100 ATRX_HUMAN
7561	TIASDSEEEAGKELSDKK20020000000000000000	sp Q96ST2 IWS1_HUMAN
7562	EDYFEPISPDR20000	sp Q69YN4 VIR_HUMAN
7563	EESEESDEDMGFLFD200200010000000	sp Q8NHW5 RLAOL_HUMAN
7564	GSSPTPPCSPVQPSK200000020000000	sp P05388 RLAO_HUMAN
7565	LSSGDPSTSPSLSQTTPSK20000000000000000000	sp Q9NUL3 STAU2_HUMAN
7566	EDEDTPNSPNVDL GK2000200000000	sp Q99504 EYA3_HUMAN
7567	MSCFSRSPMSPTPLDR2020000000	sp P98174 FGD1_HUMAN
7568	GPTSSPCEEEGDEGEEDRTSDLR2E+19	sp Q9UQ35 SRRM2_HUMAN
7569	APAPPPPQPPPSPLIPNR20000000	sp Q86VM9 ZCH18_HUMAN
7570	MSQPGSPSPK10000200000	sp 075128 COBL_HUMAN
7571	LLKPGEEPSEYTD EEDTKDHNKQD20000000000000	sp Q14839 CHD4_HUMAN
7572	GPHHLDNSSPGGSEAR2000000000	sp 015173 PGRC2_HUMAN
7573	TASPPPPPK2000000000	sp Q8WWM7 ATX2L_HUMAN
7574	HLEHAPSPSDVSNAP EVK20000000000	sp Q8IYB3 SRRM1_HUMAN
7575	SKLSPSPSLR20000200000	sp Q9Y4C1 KDM3A_HUMAN
7576	GESAPTLSTSPSPSSPSPTSPSPTLGR20202000000000	sp Q9Y4C1 KDM3A_HUMAN
7577	GLFQDEEDSCSDCSYR202002000	sp Q9Y4C1 KDM3A_HUMAN
7578	LSSSSPGATVSPSSSDA EYDKLPDADLAR2. 0000000000002E+28	sp Q5T200 ZC3HD_HUMAN
7579	ESVSPEDSEK20000000	sp Q9H330 TM245_HUMAN
7580	EEATYGVWAERDSDDERPSFGGK200000000000	sp Q8NEM2 SHCBP_HUMAN
7581	IQQEQESSGEEDSDLSPEER20000200000000	sp Q8NI08 NCOA7_HUMAN
7582	VSSSSPQSGCPSPTIPAGK200000000	sp P35251 RFC1_HUMAN
7583	SPPGSGASTPVGPWDQAVQR2. 00000002E+20	sp Q9UBB9 TFP11_HUMAN
7584	KLSSSSEPYEDEFNDDQS IKK2. 0000003E+20	sp Q6NXS1 IPP2M_HUMAN
7585	LAPVPSPEPQKPAPVSPESVK200000000000000000	sp P41236 IPP2_HUMAN
		sp Q76L83 ASXL2_HUMAN
		sp Q8TE77 SSH3_HUMAN
		sp Q9H4L7 SMRCD_HUMAN
		sp Q96JM3 CHAP1_HUMAN

7586	GSGGLFSPSTAHPD GALGQR2. 0000002E+20	sp Q9C0C2 TB182_HUMAN
7587	IDENSDKEMEVEESPEK20000000020000	sp Q8N3X1 FNBP4_HUMAN
7588	SCVEEPEPEPEAAEGDGD KKGNAEGSSDEEGK22000000	sp P51858 HDGF_HUMAN
7589	QELGSPEER200000	sp Q92974 ARHG2_HUMAN
7590	RPESPSEISPIK20020000	sp Q7Z5K2 WAPL_HUMAN
7591	GNLELRPGGAHPGTCSPSRPGS200000	sp Q96SN8 CK5P2_HUMAN
7592	DTYSDRSGSSPDSEITELK3000002220000000000	sp P17812 PYRG1_HUMAN
7593	LLDQCDLSTSR200	sp Q9Y597 KCTD3_HUMAN
7594	SPIINESR200000000	sp Q13523 PRP4B_HUMAN
7595	TSPSSPAPLPHQEATPR2000000000000000000	sp P04920 B3A2_HUMAN
7596	KVEEDLKADEPSSEESDLEIDK20000000	sp P50502 F10A1_HUMAN
7597	ATISDEEIER20000000	sp Q9UQN3 CHM2B_HUMAN
7598	KPTPFSSR2000200	sp Q8NEV8 EXPH5_HUMAN
7599	KPALPVSPAAR200000	sp Q9BWU0 NADAP_HUMAN
7600	YSSDPTGALTEDSIDD TFLPVPEYINQSVPK2E+29	sp P00533 EGFR_HUMAN
7601	SSSPVTELASR22000000200	sp Q9UQ35 SRRM2_HUMAN
7602	SSIETKPDASQLPK2000000	sp Q3KR37 GRM1B_HUMAN
7603	SQDATFSPGSEQAEK2002000000	sp Q86WB0 NIPA_HUMAN
7604	RKSEDGTPAEDGTPAATGGSQPPSMGR2E+21	sp Q9BQG0 MBB1A_HUMAN
7605	LVTSGAESGNLNTSPSSNQTR200000000	sp Q8NHV4 NEDD1_HUMAN
7606	REPGSGTESDTSPDFHNQENEPSQEDPEDLDGSVQGVKPK2. 022E+33	sp Q9HAF1 EAF6_HUMAN
7607	TSSKESSIPSP TSDRK2220000000000000000	sp Q01082 SPTB2_HUMAN
7608	ASSLNFLNK20000000	sp Q9H0B6 KLC2_HUMAN
7609	HSGGFLSSPADFSQENK20000000000	sp Q7LBC6 KDM3B_HUMAN
7610	RNSSDSAIDNPKPNK20000000000	sp P46100 ATRX_HUMAN
7611	SESFSEDLIPSR200000000000000	sp Q9P219 DAPLE_HUMAN
7612	SPTMEQAVQTASAHLPAPAAVGR1. 000002E+20	sp Q8ND56 LS14A_HUMAN
7613	YDERPGSP LPHR2000000	sp P08621 RU17_HUMAN
7614	VSVSPGR2020000	sp Q8IYB3 SRRM1_HUMAN
7615	TQDPSSPGTTPPQAR200020000000	sp Q13112 CAF1B_HUMAN
7616	SVASSQPAKPTK2000000000	sp Q07157 ZO1_HUMAN
7617	IGEGTYGVVYK23000000	sp P06493 CDK1_HUMAN
		sp Q00526 CDK3_HUMAN
		sp P24941 CDK2_HUMAN
7618	LDNTPASPPRSPAEPNDIPIAK2. 002E+19	sp O95359 TACC2_HUMAN
7619	TPSSSPITPPASETK22000200000000	sp Q9NZJ0 DTL_HUMAN
7620	KDDSHAEDSEDEKEDHK20002000000000	sp Q9H1E3 NUCKS_HUMAN
7621	TQDPSSPGTTPPQAR20002000000	sp Q13112 CAF1B_HUMAN
7622	ASSPAESSPEDSGYMR2000002000000000	sp Q9Y4H2 IRS2_HUMAN
7623	KVLSDESEDEEK202000000	sp Q8WWQ0 PHIP_HUMAN
7624	GILAADESTGSI AK20000	sp ALDOA_RABIT
		sp P04075 ALDOA_HUMAN
7625	HKSDSPESDAER200200000	sp O75400 PR40A_HUMAN
7626	GTAGKSPDLSSQK200000000	sp Q9Y2I7 FYV1_HUMAN
7627	GLQLTPGIGGMQHHFFDDED R2000001000000000000	sp P12270 TPR_HUMAN
7628	ALSSLHGDDQDSEDEVLTIPEVK2. 2E+21	sp P11717 MPRI_HUMAN
7629	SPGSTPTTSSQAPQK200000000	sp P35658 NU214_HUMAN
7630	ALSSAVQASPTSPGGSPSSSGQR2200020000000000	sp Q03164 KMT2A_HUMAN
7631	SHSGTSPDNTAPPPPPRPQPSHSR2E+25	sp Q96D71 REPS1_HUMAN
7632	RRTASPPPPPK2020000000	sp Q8IYB3 SRRM1_HUMAN
7633	SRSPPPVSK20000000	sp Q01130 SRSF2_HUMAN
7634	SNSTSSMSSGLPEQDR2000000000000000	sp P42684 ABL2_HUMAN
7635	NSSDSAIDNPKPNKLPK2020000000000000	sp P46100 ATRX_HUMAN
7636	ASPPDPK2000000	sp Q7L590 MCM10_HUMAN



7637	QILIIACSPVSSVR20002000	sp Q5QJE6 TDIF2_HUMAN
7638	SPAKSPNPSTPR2000200002000	sp O76021 RL1D1_HUMAN
7639	DIDISSPEFK200000	sp Q09666 AHNK_HUMAN
7640	CAAPRPPSSPEQR20000000	sp Q5H9R7 PP6R3_HUMAN
7641	VIKDEALSDGDDLRFPSDLK2000000000000000	sp Q01831 XPC_HUMAN
7642	ALSGTSPDDVQPGPSVGPSSK2E+19	sp Q9Y314 NOSIP_HUMAN
7643	GVDEQSDSSESESEEEKPPEEDKEEEEEK2.022002E+23	sp P35269 T2FA_HUMAN
7644	RSPSPSPTPEAK200000200000	sp Q8TAQ2 SMRC2_HUMAN
7645	TLSSSSMDLSR2000000000	sp Q9H0B6 KLC2_HUMAN
7646	SRTPPAIR202000000	sp Q9UQ35 SRRM2_HUMAN
7647	MAGNEALSPTSPFR20020000	sp Q86WR7 PRSR2_HUMAN
7648	YSQDDDDGSSSSGSSVAGSQSTLFLK200000000000	sp Q9UH99 SUN2_HUMAN
7649	SPPYTAFLGNLPYDVTEESIK2E+21	sp P23588 IF4B_HUMAN
7650	DPTQKPSPHVKPLSAAPVEGSPDR20000	sp Q9BXF6 RFIP5_HUMAN
7651	SKSPPKSPEEEGAVSS202000200000000000	sp Q01130 SRSF2_HUMAN
7652	VDSSTNSSPSPQQSELSAHTSDFR200000000000000000	sp Q9BQE9 BCL7B_HUMAN
7653	FDHESSPGTDEDK2000200000	sp Q13427 PIIG_HUMAN
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7655	YRSPEPDPYLSYR200000000000	sp P49761 CLK3_HUMAN
7656	KPVTVSPTTPTSPTEGEAS2000000	sp Q9Y6G9 DC1L1_HUMAN
7657	EKPELSEPSHLNPGSSDPEAAFLSR200000000000	sp Q9H4A3 WNK1_HUMAN
7658	SPSPSFR20200000	sp P49585 PCY1A_HUMAN
7659	SPAGSPELR2000200000	sp Q96JM3 CHAP1_HUMAN
7660	ARHSDPDSPPR200020000	sp Q9BRD0 BUD13_HUMAN
7661	RSPGEMR2000000	sp Q6NYC8 PPR18_HUMAN
7662	SSPGAVAGLSNAPGTPR2000	sp Q9BWG4 SSBP4_HUMAN
7663	DSAYQSITHYRVPVSR200	sp P50402 EMD_HUMAN
7664	LNQPGTPTR200	sp P36507 MP2K2_HUMAN
7665	RRPSGSEQSDNESVQSGR2020020002002000	sp Q6PD62 CTR9_HUMAN
7666	GSQPPPAESQSSLRR20000	sp O43491 E41L2_HUMAN
7667	LTEGCSFR2000	sp Q71UM5 RS27L_HUMAN
7668	ETAVPGPLGIEDISPNSPDDK200000	sp P42677 RS27_HUMAN
7669	LMLSTSEYSQSPK2000	sp Q12888 TP53B_HUMAN
7670	SRTPPSAPSQR20000000000000	sp Q12888 TP53B_HUMAN
7671	IAAPELHKGSDSEDEPTK200000000000	sp Q9UQ35 SRRM2_HUMAN
7672	ALEQQEAESDSDTEEKDDDDDEEDVGKR2.02E+19	sp Q92733 PRCC_HUMAN
7673	RLSPSSAASSYSFSDLNSTR200200000000000000	sp Q9BXY0 MAK16_HUMAN
7674	VALSDDETKETENMR2000000000100	sp P50402 EMD_HUMAN
7675	NSTGSGHSAQELPTIR2000000000000000	sp Q15054 DPOD3_HUMAN
7676	GSGSPHSLSQK2000000000	sp O94763 RMP_HUMAN
7677	GRLSVASTPISR20020002000	sp Q9BSJ6 FA64A_HUMAN
7678	TRLSPPR20000	sp Q9BXS6 NUSAP_HUMAN
7679	DEILPTPISEQK2000000000	sp Q96PK6 RBM14_HUMAN
7680	HYEDGYPGSDNYGSLSR20000	sp P23396 RS3_HUMAN
7681	SPQLATPGSSHPGEEECR20000000000000000000	sp O60716 CTND1_HUMAN
7682	LEPQELSPLSATVFPK20000000000	sp Q9C0A6 SETD5_HUMAN
7683	TTHLAGALSPGEAWPFESV200000000000	sp Q3T8J9 GON4L_HUMAN
7684	NMVDLVNTHHLHSSSDEDDRK200000000000	sp Q9H869 YYAP1_HUMAN
7685	EYIPGQPPLSQSSDSSPTR20020000	sp Q9Y3X0 CCDC9_HUMAN
7686	GGPGAAPGGASPASSSSAASSPSSGR2000000000000000	sp Q9UPN7 PP6R1_HUMAN
7687	LDSSEMDSHENEDYTMSSPLPGK2.00002E+20	sp P07814 SYEP_HUMAN
7688	TPTHQPPKPGSPGRPACPLSNLLLSYEER2E+20	sp Q9BXX1 KLF16_HUMAN
		sp Q9NTI5 PDS5B_HUMAN
		sp Q96F63 CCD97_HUMAN

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7690	PGTPSGTNVGSGRSPSK20000	sp P60468 SC61B_HUMAN
7691	NSQKHSPTSEPTPPGDALPPVSSPHTHR200000000020000000	sp Q9H6S3 ES8L2_HUMAN
7692	DINTFLGTPVQK200000	sp P46013 KI67_HUMAN
7693	RPGNSTVTSSPPVTPK2000002000	sp Q7Z2Z1 TICRR_HUMAN
7694	QGSGRESPLASR200000	sp Q8WWM7 ATX2L_HUMAN
7695	VVELRPPSR200	sp Q9P1Y6 PHRF1_HUMAN
7696	I IPTLEGLQLPSPTATSQ LPLESDAVECLNYQH YK2E+22	sp P61978 HNRPK_HUMAN
7697	DQTASAPATPLV NK20200000000000	sp Q53SF7 COBL1_HUMAN
7698	TLSNAEDYLDDESD20000000000000	sp Q92882 OSTF1_HUMAN
7699	KHSPSPPTPTESR200000000000	sp Q92922 SMRC1_HUMAN
7700	QSSTPSAPELGQQPDVNI SEWK2E+19	sp Q93100 KPBB_HUMAN
7701	SSPSARPPDVP GQQPQAAK20000000000000000000	sp Q96JP5 ZFP91_HUMAN
7702	SSLAETLDSTGSLDPQR2002000000	sp Q9UGH3 S23A2_HUMAN
7703	VAAQSPSPASAEALLPALPLR20020000000000000000	sp Q9Y4F5 C170B_HUMAN
7704	AGTATSPAGSSPAVAGGTQRPAEDSSSSEESDSEEEK2E+35	sp Q13428 TCOF_HUMAN
7705	LLSTDAEAVSTR2000000000	sp Q15545 TAF7_HUMAN
7706	RHSSETFSSTTTVTPVSPFAHNPK2. 2E+23	sp Q9Y4H2 IRS2_HUMAN
7707	SLGNILQAKPTSSPAK20000	sp Q13428 TCOF_HUMAN
7708	LSSLSSQTEPTSAGDQYDCSR2E+19	sp Q9BY89 KI671_HUMAN
7709	RVSLEPHQPGTPESK200000000200000	sp Q14980 NUMA1_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
7710	GGSYSQAASSDSAQGS DVSLTACK3. 000000000002E+21	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
7711	APLVP P P P P P P P P P P P P L P D P T P P E P E E E I L G S D D E E Q E D P A D Y C K 2 E + 2 5	sp P78362 SRPK2_HUMAN
7712	SLGSSPCPK2200000	sp Q9UQ35 SRRM2_HUMAN
7713	SLGSASPGPGQPPLSSPTR20000	sp Q9C0B5 ZDHC5_HUMAN
7714	DTPRPDHPHDGHSPASR200000	sp Q9C0J8 WDR33_HUMAN
7715	EEEEGISQESSEEEQ2200000	sp P17096 HMGA1_HUMAN
7716	RSSKSFSLDEPPLFIPDNIATIR2. 02E+19	sp Q92576 PHF3_HUMAN
7717	GKSSEPVVIMK2000000100	sp P46013 KI67_HUMAN
7718	KKTVSASESEDR20202000000	sp Q9ULG6 CCPG1_HUMAN
7719	YSPTSPTYSPTSPVYTPTSPK2. 0000000000003E+20	sp P24928 RPB1_HUMAN
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7721	GTPSQSPVVGR200000000	sp P52594 AGFG1_HUMAN
7722	SPEHSAEPIR20000000000	sp Q9ULH0 KDIS_HUMAN
7723	KHSEEAFTPPLK20000000000	sp Q3B726 RPA43_HUMAN
7724	RHSTSSLSPQR22000000	sp Q03164 KMT2A_HUMAN
7725	LVDVSPDRGSPSR20000000200	sp Q9NYZ3 GTSE1_HUMAN
7726	NTSSDNSDEVMPAQSPR20002000010000000	sp Q9HCK8 CHD8_HUMAN
7727	KL GAGEGGEASVSPEK20000	sp Q13428 TCOF_HUMAN
7728	TFSFSDDENKPPSPK20200000000000	sp Q9UHJ3 SMBT1_HUMAN
7729	KASSPSPLTIGTPESQR2002000000000000	sp Q9NPI6 DCP1A_HUMAN
7730	EEDEENDNDNESDHDEADS200000000	sp Q9NRG0 CHRC1_HUMAN
7731	VPSVESLFR20000000	sp Q13439 GOGA4_HUMAN
7732	SSGSLGHRPSQEMDK2000000000000	sp Q5SW79 CE170_HUMAN
7733	VLSSTSEDEPGVVK22220000000000	sp Q8NI08 NCOA7_HUMAN
7734	CPVKQDSVESQLK20000000	sp Q5T8I3 F102B_HUMAN
7735	RASPSKPASAPASR20000000000	sp P27816 MAP4_HUMAN
7736	EGEETEGSEEDDKENDKTEEMPNDVLENK2E+26	sp P49321 NASP_HUMAN
7737	SPISELHSAPLTPVAR200200000000200000	sp Q7Z3B3 KANL1_HUMAN

7738	SPGPPALK200000000	sp Q8IU81 I2BP1_HUMAN
7739	SPSMVPSPGWVASPK200002000000000	sp Q86UU0 BCL9L_HUMAN
7740	KRSESPPAELPSLR2020000000000	sp Q14669 TRIPC_HUMAN
7741	GYQSLSDSPPGSR2000000	sp Q9NPR2 SEM4B_HUMAN
7742	SLKESEQESEEEILAQKK200020000000000	sp Q9BW71 HIRP3_HUMAN
7743	LINSELGSPSR20000	sp Q68CP4 HGNAT_HUMAN
7744	HTPNTSDNEGSDTEVCGPNSPSK20200000000000	sp Q8NEN9 PDZD8_HUMAN
7745	SPTPALCDPPACSLPVASQPPQHLSEAGR2E+29	sp Q9P2R6 RERE_HUMAN
7746	LAGGQTSQPTTPLTSPQR20000	sp Q095155 UBE4B_HUMAN
7747	GRLTPSPDIIVLSDNEASSPR20200000000000020000	sp Q8WXI9 P66B_HUMAN
7748	HHHHDSDSNSPCK2000200000	sp Q9BU76 MMTA2_HUMAN
7749	SQSFHQPSR2000000000	sp Q8WVR3 CGO43_HUMAN
7750	TPLSSLSATSNNSTVACIDR200000000000000000	sp P00533 EGFR_HUMAN
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7752	RSSLLNAK2000000	sp Q8TEU7 RPGF6_HUMAN
7753	SKTPSSPCPASK2000020000000	sp Q9NQS7 INCE_HUMAN
7754	SGPKPFSAPKQTSPPSK200000	sp Q01518 CAP1_HUMAN
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7757	SRSPLELEPEAK2020000000000	sp Q92466 DDB2_HUMAN
7758	SSRSVEDDKEGHLVCR20200000000000000	sp P49761 CLK3_HUMAN
7759	SHSGTSPDNTAPPPPPRPQPSHSR2. 2E+21	sp Q96D71 REPS1_HUMAN
7760	RASSPFR220000	sp Q14978 NOLC1_HUMAN
7761	SGSDAGEARPTPASPR2000000000000000000	sp Q96FS4 SIPA1_HUMAN
7762	SPGSQAPDEGAGGALR200000000000000000	sp P10075 GLI4_HUMAN
7763	GHHVTDSENDEPLNLNASDSESEELHR2. 0000000000202E+21	sp Q96ST2 IWS1_HUMAN
7764	LLDESESAASSR200	sp Q8N108 MIER1_HUMAN
7765	WDQTADQTPGATPK20002000	sp Q075533 SF3B1_HUMAN
7766	RGEGDAPFSEPQTSTQRPSSPETATK2E+19	sp Q9UQ35 SRRM2_HUMAN
7767	SPENSPPVKPK200000000000	sp Q8TF72 SHRM3_HUMAN
7768	RPAEPGGPQEPGLASSPPGR200000	sp Q9BZC7 ABCA2_HUMAN
7769	ALPQTPRPR200000	sp Q9UQ35 SRRM2_HUMAN
7770	LSTPAHSPVLK2000200000	sp Q8IY63 AMOL1_HUMAN
7771	GGSGSHNWGTVK20000000000	sp Q8NC51 PAIRB_HUMAN
7772	VVVSPK20000	sp P19338 NUCL_HUMAN
7773	RYSPPPKR3002000000	sp Q8IYB3 SRRM1_HUMAN
7774	EPLENSASESDTELPEK20202200000000	sp P78337 PITX1_HUMAN
7775	GSPSPEKPDVEK200000000000	sp Q93074 MED12_HUMAN
7776	TVCGSPLAAQSPSAELQAAPPPPGSPSLPGR2. 0000000000002E+20	sp Q969V6 MKL1_HUMAN
7777	TEMKSPFNPSPQDSR10000002020000000	sp P08651 NFIC_HUMAN
7778	TPTPSTPGDTQNTAPVPPAEDGIK2. 0200000000002E+26	sp Q14839 CHD4_HUMAN
7779	FTMDLSDDEDFSDFEKTDDEDFVPSDASPPK2. 00002000002E+26	sp P11388 TOP2A_HUMAN
7780	RPMSSPEMK200000	sp P51610 HCFC1_HUMAN
7781	SQDATFSPGSEQAESKSPGPIVSR2. 02002E+19	sp Q86WB0 NIPA_HUMAN
7782	LPSANQGSFPK20000	sp Q56NI9 ESCO2_HUMAN
7783	VTSSVPLPSGSTSSPGR20000000	sp Q15036 SNX17_HUMAN
7784	SPHQLLSPSSFSPSATPSQK202200000000000	sp Q14181 DPOA2_HUMAN
7785	GAAPTPPGK200000	sp Q13428 TCOF_HUMAN
7786	SVASSQPAKPTK200	sp Q07157 ZO1_HUMAN
7787	AQTPPISSLPTSPSDEVGR200000002000000000	sp Q15751 HERC1_HUMAN
7788	KRNSISDDDTSEDEL202000202000000	sp Q76FK4 NOL8_HUMAN
7789	RGSPASPTR200020000	sp Q6P5Z2 PKN3_HUMAN
7790	ALLQSSASR20000	sp Q9UQ80 PA2G4_HUMAN

7791	KAPSASDSKADSDGAKPEPVAMAR2. 0202000002E+23	sp Q7Z4V5 HDGR2_HUMAN
7792	SRTPPVTR200000000	sp Q9UQ35 SRRM2_HUMAN
7793	YHGHSMSDPGVSYR21000000000	sp P08559 ODPA_HUMAN
7794	STSTPTSPGPR2000200000	sp P29803 ODPAT_HUMAN
7795	TDALTSSPGR2200000	sp Q8WWM7 ATX2L_HUMAN
7796	VEQMPQASPLAPR200000000	sp P49736 MCM2_HUMAN
7797	TPTPSTPGDTQPNTAPVPPAEDGIK2. 0002E+21	sp Q8N3F8 MILK1_HUMAN
7798	RRSPSPPTR202000000	sp Q14839 CHD4_HUMAN
7799	SPSESSPQHPTTPARPR20000020000000000000	sp Q8IYB3 SRRM1_HUMAN
7800	KKSSSEAKPTSLGLAGGHK220000000000000000	sp O14733 MP2K7_HUMAN
7801	SPTAQSPTPSVEADSPDK2E+19	sp Q96RK0 CIC_HUMAN
7802	SDDSSTLPR200000000	sp Q96HH9 GRAM3_HUMAN
7803	ISAPNVDFNLEGPK200000000000000	sp Q9ULE0 WWC3_HUMAN
7804	APQQPPPPQPPPPQPPPPQPPPPPSYSPAR2000000	sp Q09666 AHNK_HUMAN
7805	MSDSESEELPKPQVSDSESEEPPI. 20202E+24	sp Q9BUJ2 HNRL1_HUMAN
7806	AAPPPPPPPPLESSPR20000	sp Q96ST2 IWS1_HUMAN
7807	ALSKPGTAAELR200000000000	sp Q5T4S7 UBR4_HUMAN
7808	AGGGPTLQCPPSSPEK20000	sp Q9BZ29 DOCK9_HUMAN
7809	RESISPQADSACSSPAPSTGK2. 02000002E+20	sp Q96N66 MBOA7_HUMAN
7810	EDTDHEEKASNEVTK200000020000000	sp Q9GZR1 SENP6_HUMAN
7811	LASVPAGGAVAVSAAPGSAAPAAGSAPAAAEK200000000000000000	sp O75475 PSIP1_HUMAN
7812	LGSYSGPTSVSR200000000000	sp P05387 RLA2_HUMAN
7813	RNSNSPPSPSSMNQR200002000000000	sp Q9BZ23 PANK2_HUMAN
7814	MSQPGSPSPK2000200000	sp Q7Z5L9 I2BP2_HUMAN
7815	TAASVLNFIKESPK20000	sp Q14839 CHD4_HUMAN
7816	SDLSSSSGSLSLSHGSSSLEHR200000000000000000	sp Q6N021 TET2_HUMAN
7817	QISSSTGCLSSPNATVQSPK220000000000000000	sp O15013 ARHGA_HUMAN
7818	SVSSYGNIR2002000000	sp Q9Y6Y0 NS1BP_HUMAN
7819	ITEVSCKSPQDPVKPTSSK200000002000000	sp Q8N122 RPTOR_HUMAN
7820	SPSDSSTASTPVAEQIER200000000000000	sp P46013 KI67_HUMAN
7821	SRSHSPMSNR20202000000	sp Q16643 DREB_HUMAN
7822	IRHSSDTSPPR220020000	sp Q13595 TRA2A_HUMAN
7823	SGAQASSTPLSPTR200000000	sp Q9BRD0 BUD13_HUMAN
7824	ILLDAQHESGQSSSR200	sp P02545 LMNA_HUMAN
7825	DDDSLPAETGNHPFFR2000000000000000	sp O60238 BNI3L_HUMAN
7826	KPLTTSGFHHSEEGTSSSGSK200000	sp O95071 UBR5_HUMAN
7827	RRSPSLSSK200000000	sp Q5SW79 CE170_HUMAN
7828	SPMSTNSSVHTGSDVEQDAEK2. 010000000002E+21	sp Q8IYB3 SRRM1_HUMAN
7829	ASTTSLSGDSETEGK20202000000	sp Q9ULU4 PKCB1_HUMAN
7830	RSSLSPSSAYER202000000000	sp Q8N1G2 CMTR1_HUMAN
7831	IGPLGLSPK2000	sp Q9BYW2 SETD2_HUMAN
7832	AGGKPSQSPSQEAAGEAVLGAK200000000000000000	sp P30050 RL12_HUMAN
7833	SAASPVSSMPERASESSSEEKDDYEIFVK220000000000000	sp P00519 ABL1_HUMAN
7834	ASPSPPK20200000	sp P18583 SON_HUMAN
7835	SPSPSPAAVNHSSSDISPVSNSTSSSPGK220000000	sp Q8IYB3 SRRM1_HUMAN
7836	SSPVNNHSPGQTPTLGQPEAR2. 2E+22	sp O75381 PEX14_HUMAN
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7838	SVSSPTSNTPTTK202000000000000000	sp Q96L14 C170L_HUMAN
7839	GPTSSPCEEEDGEDRTSDLR2. 00000000000002E+19	sp Q8WUZO BCL7C_HUMAN
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7841	YSDGDKSDDLVVDVSNEDPATPR200000000000000000	sp Q86VM9 ZCH18_HUMAN
7842	SAGGSSPEGGEDSDREDGNYCPPVK2. 20000002E+21	sp Q8IYB3 SRRM1_HUMAN
		sp Q04726 TLE3_HUMAN
		sp Q9H6Z4 RANB3_HUMAN

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7845	RGSSAAASPGSPPPGR200002000000000	sp P29803 ODPAT_HUMAN
7846	MISTPSPK202000	sp Q86UX6 ST32C_HUMAN
7847	HASSPESPAPAPGSHR222000000000002000	sp Q53F19 CQ085_HUMAN
7848	QESCPHHPQVLAQQGSGSSPK2.02E+20	sp Q9UQ35 SRRM2_HUMAN
7849	SSPSGITHVIASNAR2000000000000	sp Q8N1G0 ZN687_HUMAN
7850	STSAPQMSPGSSDNQSSSPQPAQQK2002200000000000000	sp Q9BQI6 ANR32_HUMAN
7851	SMGGAIIAPPTSLVEK20000000000000000	sp Q14157 UBP2L_HUMAN
7852	TAAELLSQGSQAGGSQTLK2000	sp Q96I25 SPF45_HUMAN
7853	SKSPPKSPEEEGAVSS200020000000020	sp Q14141 SEPT6_HUMAN
7854	TPDGNKSPAPKPSDLRPGDVSSK2.000002E+23	sp Q01130 SRSF2_HUMAN
7855	ASETPPPVAQPKPEAPHPGLETTLQER2E+24	sp Q05682 CALD1_HUMAN
7856	LEGSDDLLEDSDSEEHSR200000200	sp Q6P1N0 C2D1A_HUMAN
7857	LNSPEETAFTPK200000002000	sp Q9NR09 BIRC6_HUMAN
7858	NPPPQDYESDDSYEVLDTLEYAR3000000000000	sp Q9BTX1 NDC1_HUMAN
7859	GPSPPASPTR200020000	sp Q8IVP5 FUND1_HUMAN
7860	HFSEHPSTSK200000200	sp Q6P5Z2 PKN3_HUMAN
7861	KPSDSLVAASSR200020022000	sp O43719 HTSF1_HUMAN
7862	SRTASGSSVTSLDGTR202020000000000	sp Q5T0W9 FA83B_HUMAN
7863	TDSDDLQLYK20000000	sp Q92597 NDRG1_HUMAN
7864	SSPVNRPSSATNK2000000000000	sp Q01433 AMPD2_HUMAN
7865	MSPNETLFLESTNK2000	sp Q96L91 EP400_HUMAN
7866	RGSSSSPEHSASSDSTK220000000000000	sp Q9HB90 RRAGC_HUMAN
7867	YGGVGSQPPVPEPVPSSPSQEPPTK2000000000	sp Q9NQL2 RRAGD_HUMAN
7868	SPASCHEPMADDPSK2000000000000	sp Q9P275 UBP36_HUMAN
7869	LPSSSGLPAADVSPATAEPLSPSTPTR20000000	sp Q04323 UBXN1_HUMAN
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7873	DQQNLPYGVTPASPSGHSQGR20000000000	sp Q7Z460 CLAP1_HUMAN
7874	ESSITSCCSTSSCADDEGVR2E+19	sp Q01082 SPTB2_HUMAN
7875	RSPPAPGLQPMR200000000100	sp Q7KZI7 MARK2_HUMAN
7876	HQSFGAAVLSR2000000000	sp Q9UIC8 LCMT1_HUMAN
7877	LALSPEDKPIR200000000	sp P15408 FOSL2_HUMAN
7878	GVSRSPPK2020000	sp Q5VV41 ARHGG_HUMAN
7879	GTEPSPGTQPSPRVSPAGPPEGVPEEAQPPR2.0000002E+32	sp O15417 TNC18_HUMAN
7880	ASSLESPK20000020000	sp Q8TAD8 SNIP1_HUMAN
7881	SSSSGVPYSPAIPNK20000000	sp Q8WUZ0 BCL7C_HUMAN
7882	HDYDSSSEQAEDSYEASPGSETQR20000000000000000	sp Q9UHB6 LIMA1_HUMAN
7883	FHHTIGGSR200	sp Q9UPT5 EXOC7_HUMAN
7884	TFSLTEVR202000000	sp P98175 RBM10_HUMAN
7885	LSVSVTSRPSTPTLGTPTQTMSVSTK2.02E+24	sp P61313 RL15_HUMAN
7886	KRTPSPSYQR200020000	sp O95239 KIF4A_HUMAN
7887	GPGQPSSPQR20000	sp Q16594 TAF9_HUMAN
7888	QDDGSSASPSVQGAPR2000000000	sp Q5T200 ZC3HD_HUMAN
7889	STSAPQMSPGSSDNQSSSPQPAQQK1.000000022E+19	sp O95400 CD2B2_HUMAN
7890	LSTTPSPTSSLHEDGVEDFR220000000000000	sp Q9NQA3 WASH6_HUMAN
7891	LGSLTPEIVSTPSSPEEDK200000000	sp Q14157 UBP2L_HUMAN
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7893	SWSPPEVSR200000000	sp Q14CS0 UBX2B_HUMAN
7894	DVTPPPETEVLK2000000000000	sp Q9HCD6 TANC2_HUMAN
		sp Q9NR19 ACSA_HUMAN
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		sp Q641Q2 FA21A_HUMAN
		sp Q5SNT6 FA21B_HUMAN
7896	TESPPSPSSDVEDAR20020200000000	sp Q9BRK4 LZTS2_HUMAN
7897	SREDLSAQPVQTK20000000000000	sp Q07157 ZO1_HUMAN
7898	SWSPPEVSR20200000000	sp Q9NR19 ACSA_HUMAN
7899	APSDSSLGTPSDGRPELR2000002000000000	sp Q15642 CIP4_HUMAN
7900	AAESETPGKSPEKKPK20000020000000	sp 076021 RL1D1_HUMAN
7901	SRSRSPPPVSK202000000000	sp Q01130 SRSF2_HUMAN
7902	LHQLSGSDQLESTAHSR200000000000	sp 060271 JIP4_HUMAN
7903	SPSPTPTGPSR20000000000000	sp P49450 CENPA_HUMAN
7904	LDSSEMDHSENYDTMSSPLPGK2.2010000000001E+21	sp Q9NTI5 PDS5B_HUMAN
7905	ETPRPEGGSPAGTPPQPK20000020000000	sp Q9BR39 JPH2_HUMAN
7906	STYFSDEEELSD200000200	sp Q8TBF4 ZCRB1_HUMAN
7907	SKSHESQLGNR2000000000	sp Q8IVT5 KSR1_HUMAN
7908	KSDGACDPSDDKENSQIAQDHQK20020000000000000000	sp Q9UIG0 BAZ1B_HUMAN
7909	KTSEDQAAILPGK200000000000	sp Q12968 NFAC3_HUMAN
7910	SLSPILPGR2000000000	sp Q6ZS75 ARHG1_HUMAN
7911	SLSYSPVER200000	sp Q9UQ35 SRRM2_HUMAN
7912	QSPPLPGR20000000	sp Q9C0B5 ZDHC5_HUMAN
7913	YYGGAGGSPGK20000	sp Q9BUH8 BEGIN_HUMAN
7914	SSGSPYGGYGGSGGGSGGYGSR200000000000	sp P51991 ROA3_HUMAN
7915	TEMDKSPFNPSQDSR20000000002000	sp P08651 NFIC_HUMAN
7916	MALEVGMEDGQLSDSDMTVAPSDRPLQPK1.0000001E+33	sp Q9H814 PHAX_HUMAN
7917	SPSMVPSPGWVASPK210002000002000	sp Q86UU0 BCL9L_HUMAN
7918	NRSPSDDMEDYSPPSLSEVAR2.0202000002E+21	sp Q6KC79 NIPBL_HUMAN
7919	STPLASPSPGRSPQR20000000200020000	sp Q8WYP5 ELYS_HUMAN
7920	SRSSVGSSSSYPIPAVSR2002220000000000	sp Q15149 PLEC_HUMAN
7921	TNSSDSERSPDLGHSTQIPR22020020000000000000	sp Q6VY07 PACS1_HUMAN
7922	TPTPSTPGDTQPNTAPVPPAEDGIK2.000000000002E+26	sp Q14839 CHD4_HUMAN
7923	EEKESEDKPEIEDVGSDEEEKK2.0000000002E+19	sp P07900 HS90A_HUMAN
7924	SSPGAVAGLSNAPGTPR2000000000000000	sp Q9BWG4 SSBP4_HUMAN
7925	SRSRSPTPSSAGLGSNSAPPIDSR2.02020022E+24	sp Q8IWX8 CHERP_HUMAN
7926	IRHSDSDTSPPR200220000	sp Q9BRD0 BUD13_HUMAN
7927	LQQTVPADASPDSK200000	sp Q9NS56 TOPRS_HUMAN
7928	STPLASPSPGRSPQR20000	sp Q8WYP5 ELYS_HUMAN
7929	TTYDSAEENKENLYAGK20000000000000000000	sp Q9HAW4 CLSPN_HUMAN
7930	IPESETESTASAPNSPR20002000	sp Q07889 SOS1_HUMAN
7931	SPPPRSPSPPTH2000020002000000	sp Q96HA1 P121A_HUMAN
7932	SSSLGSTPHEELER22002200000000	sp Q9NRA8 4ET_HUMAN
7933	HTPNTSDNEGSDTEVCGPNPSK20000200000000000000	sp Q8NEN9 PDZD8_HUMAN
7934	ASIANSDGPTAGSQTPPFK20000000	sp 075167 PHAR2_HUMAN
7935	KPSGSGDQSKPNCTK20000000	sp Q8TBM8 DJB14_HUMAN
7936	SPSLGMLSTATR20000000000000	sp Q14687 GSE1_HUMAN
7937	TKCEILSDESSDEDEK200222000000	sp P35659 DEK_HUMAN
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7939	SAGGSSPEGGEDSDREDGNYCPPVKR2000000000000000	sp Q9H6Z4 RANB3_HUMAN
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7941	AQTLPTS VVTITSESSPGK20000	sp Q01082 SPTB2_HUMAN
7942	LSPSPSSR20200000	sp P20700 LMNB1_HUMAN
		sp Q03252 LMNB2_HUMAN
7943	SYQNSPSSDDGIRPLPEYSTEK20220000000000000000	sp Q15648 MED1_HUMAN
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7945	GPGDLMGPQGLSPEEMAR20000000	sp Q86UU0 BCL9L_HUMAN

7946	GYYSYSPVSVSGSGSTAGSR20200000000	sp Q15149 PLEC_HUMAN
7947	SLSSSPVKK20200000	sp Q9H3Q1 BORG4_HUMAN
7948	TPIQVESPQPGLPAGEQLEGLK2.0000002E+23	sp Q99618 CDCA3_HUMAN
7949	SLHSAHSLASR200200	sp O15027 SC16A_HUMAN
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7951	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2E+32	sp Q8WUZO BCL7C_HUMAN
7952	NEEDEGHNSNSPR2000	sp Q14103 HNRPD_HUMAN
7953	VDEDSAEDTQSNHGK200000	sp Q9H2G2 SLK_HUMAN
7954	DMLPSPEQTLSPK200000	sp Q53SF7 COBL1_HUMAN
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7960	KGTTPPRSPEASPK20000200000000	sp Q9HDC5 JPH1_HUMAN
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		sp P01892 1A02_HUMAN
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		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
		sp P30447 1A23_HUMAN
7962	ISDSESEDPPRNQASDSENEELPKPR2.0200000000002E+25	sp Q96ST2 IWS1_HUMAN
7963	AQSVSPVQAPPPGSAQLLPGK2E+20	sp Q96RK0 CIC_HUMAN
7964	DAVFDGSSCISPTIVQQFGYQR20000000000	sp P04049 RAF1_HUMAN
7965	LMELHGEGSSSGK2000	sp P61247 RS3A_HUMAN
7966	TESVSDKEDKPLAPSGGTGPEQPPPPCPSQTGSPPVGLIK2000000000000 0	sp Q8WWM7 ATX2L_HUMAN
7967	SIVPSSPQQR2000000	sp Q9P246 STIM2_HUMAN
7968	WFSNPSR200000	sp Q9NRA8 4ET_HUMAN
7969	TPPEDCEGPFRAEILK2000000000000000000	sp P49790 NU153_HUMAN
7970	SKGPSAAGEQEPDKESGASVDEVAR2E+25	sp P50579 MAP2_HUMAN
7971	LILTDSGSAPTFCSDDEDVAPLSAK2000000000000	sp P78312 F193A_HUMAN
7972	RRSSPSARPPDVPGQQPQAAK2.002E+19	sp Q96JP5 ZFP91_HUMAN
7973	EHYPVSSPSSPSPAQGGVSR3.000000202E+20	sp O75179 ANR17_HUMAN
7974	SHSGTSPDNTAPPPPPRPPQPSHSR2.0202E+25	sp Q96D71 REPS1_HUMAN
7975	IAISPLKSPNK200020000	sp Q6MZP7 LIN54_HUMAN
		sp P20700 LMNB1_HUMAN
7976	LKLSPSSSR20000000	sp Q03252 LMNB2_HUMAN
		sp Q16629 SRSF7_HUMAN
7977	YFQSPSR200	sp Q9Y6E2 BZW2_HUMAN
7978	FVEWLQNAEEEESEGEEN202000000	sp Q53LP3 SWAHC_HUMAN
7979	DLVMGSSPQLK200000	sp Q12965 MYO1E_HUMAN
7980	NQYVPYPHAPGSQR2000	sp Q8IYB7 DI3L2_HUMAN
7981	RPGTQGHGPEKEEEEESDGEPEDSSTS200000000000	sp Q8IWX8 CHERP_HUMAN
7982	SPTPPSSAGLGSNSAPPIDPSR20000200000000000	sp Q02880 TOP2B_HUMAN
7983	KTSFDQDSVDVIFPSDFPTEPPSLPR2.00002E+24	sp Q9BTA9 WAC_HUMAN
7984	DGGTYSYPQENSHNHSALHSSNSHSSNPSNPSK2E+31	sp Q12888 TP53B_HUMAN
7985	SNVSSPATPTASSSSSTPTR2020000000000000	sp Q9Y4H2 IRS2_HUMAN
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7988	SPPKSPEEEGAVSS2000000000000200	sp Q13523 PRP4B_HUMAN
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7996	DPSIKPVTPSPSK2020000	sp Q9H792 PEAK1_HUMAN
7997	IPMPSPSPQPR200000	sp Q8WUF5 IASPP_HUMAN
7998	AGDPGEMPSPTGLGQPK2000000000	sp Q9UIF9 BAZ2A_HUMAN
7999	KHSSDDYYGDISSLESSQK3000000000000000	sp Q9UGU5 HMGX4_HUMAN
8000	APSASDSKADSDGAKPEPVAMAR2. 0002000002E+23	sp Q7Z4V5 HDGR2_HUMAN
8001	TPTPSTPGDTQPNTAPVPPAEDGIK2. 002E+24	sp Q14839 CHD4_HUMAN
8002	VEGSPTEEANLPTELQDNSLSPAASEAGEK2E+27	sp P50851 LRBA_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
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8003	GGSYSQAASSDSAQGSVDVSLTACK20000000000000	sp Q09160 1A80_HUMAN
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		sp P30447 1A23_HUMAN
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8005	SPFNPSPPQDSR2000002000	sp P08651 NFIC_HUMAN
8006	SYPGHVQPATAR20000000000000	sp Q86VS8 HOOK3_HUMAN
8007	GFSQYGVSGSPTK20000	sp Q9NXC5 MIO_HUMAN
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8009	SPSPSPTPEAK202020000000	sp Q8TAQ2 SMRC2_HUMAN
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8012	NTETSKSPEKDVPMVEK200000000000	sp 014617 AP3D1_HUMAN
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8014	KPPDSYPPK230000	sp 043184 ADA12_HUMAN
8015	ASSHSSQTQGGGSVTK20000	sp P02545 LMNA_HUMAN
8016	ILKSPEIQR2000000	sp P36578 RL4_HUMAN
8017	RRSLSPKPR20200000	sp Q13523 PRP4B_HUMAN
8018	RPPGPDSPDGPR2000000	sp P22455 FGFR4_HUMAN
8019	SRGSSQELDVKPSASPQER22000000020000000	sp Q9UQ35 SRRM2_HUMAN
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8021	DLPTIPGVTSPSSDEPPMEASQSHLR200000001000000000	sp Q13541 4EBP1_HUMAN
8022	LAIQGPEDSPSR20000	sp Q15773 MLF2_HUMAN
8023	EMSSDSEYSDDDRTK220200020000000	sp 060841 IF2P_HUMAN
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8025	VATPPNQNK200000000	sp Q6NUQ4 TM214_HUMAN
8026	LSSLPLR2020000	sp Q66K74 MAP1S_HUMAN
8027	KPPAPSPVQSQSPSTNWSAPVPVKK2. 000202000002E+20	sp Q8IYB3 SRRM1_HUMAN
8028	CALSSPSLAFTPIK200000000000	sp Q8NFH5 NUP53_HUMAN
8029	HTDDEMTGYVATR200	sp Q16539 MK14_HUMAN
8030	GFSLEELR2000000	sp P26373 RL13_HUMAN
		sp 043768 ENSA_HUMAN
8031	YFDSGDYNMAK200001000	sp P56211 ARP19_HUMAN
		sp Q9UKV3 ACINU_HUMAN
8032	EKSKSPSPR202020000	sp Q7Z309 F122B_HUMAN
8033	MFVSSGLPPSPVPSR20002000	sp Q8IWS0 PHF6_HUMAN
8034	DRSPHRSSPSDTRPK20000200000000	sp Q96JH7 VCIP1_HUMAN
8035	TVSPSTIR2000000	sp Q8TEQ6 GEMI5_HUMAN
8036	ENSGPVENGVSQEGEEQAREPELPCGLAPAVSR2E+32	sp Q8IYB3 SRRM1_HUMAN
8037	TASPPPPPKR200000000	sp Q86TB9 PATL1_HUMAN
8038	RSTSPIIGSPPVR2020000000000	sp P48634 PRC2A_HUMAN
8039	QRGETGSETHESDLAPSDK200020000000000000	sp Q9Y2L9 LRCH1_HUMAN
8040	ENSPAVSPTTSTAPFGLKPR2. 000002E+19	



8041	RPNKQEESESPVERPLK200000000	sp Q96SB4 SRPK1_HUMAN
8042	LKSSTSFANIQENSN220000000000	sp Q86WC4 OSTM1_HUMAN
8043	YGPADVDTTGGGATDSKDDDDIDLFGSDDEESEEAKR2.00000000002E+23	sp P24534 EF1B_HUMAN
8044	SGKYDLDFK2000000000	sp P06733 ENOA_HUMAN
8045	SPVPSPGSSSPQLQVK200000000	sp Q8N3F8 MILK1_HUMAN
8046	ILEGQQSLSGSPLQPIPASPPPQIPPGLRPR2.000000000002E+25	sp O15446 RPA34_HUMAN
8047	IRLSSASR200	sp Q96MU7 YTDC1_HUMAN
8048	AGSSASSPPSASSSPHPSAAVPAADPADSASGSSNK2E+33	sp Q8NDF8 PAPD5_HUMAN
8049	TPHYGSQTPLHDGSR2000000200000000	sp O00267 SPT5H_HUMAN
8050	EYIPGQPPLSQSSDSSPTR220000000	sp P07814 SYEP_HUMAN
8051	TRESHRSRSSPTAGPSTEGAEGPEEK2.0222E+21	sp Q8NHM5 KDM2B_HUMAN
8052	DLDRPESQSPK2000	sp Q8WUA4 TF3C2_HUMAN
8053	AKSPTPSPPR20002020000	sp Q8IYB3 SRRM1_HUMAN
8054	SPIIHSPK200002000	sp O94913 PCF11_HUMAN
8055	HSSETFSSTTTVTPVSPSFAHNPK2E+23	sp Q9Y4H2 IRS2_HUMAN
8056	DNEESEQPPVPGTPTLR200000	sp O15439 MRP4_HUMAN
8057	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2.002E+32	sp Q8WUZ0 BCL7C_HUMAN
8058	NHSGSRTPPVALNSSR20200000000000	sp Q9UQ35 SRRM2_HUMAN
8059	KSSPSKENESSDSEEEVSRPR200002202000000000	sp Q6KC79 NIPBL_HUMAN
8060	SMPEGFGVSR20000000000	sp O43166 SI1L1_HUMAN
8061	DGGTSYSQENSHNHSALHSSNSHSSNPSNPSK2E+28	sp Q9BTA9 WAC_HUMAN
8062	NHETDGGSAHGDDDDGPHFEPVVPLPDK2E+26	sp P49792 RBP2_HUMAN
8063	STSAPQMSPGSSDNQSSSPQAQK2.00012002E+23	sp Q14157 UBP2L_HUMAN
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8065	SKLSPSPSLR20000000	sp Q5T200 ZC3HD_HUMAN
8066	SPLTATPTFVTDSESTR200000000000000000	sp P13056 NR2C1_HUMAN
8067	ATVTPSPVKGK20002000000	sp Q9H1E3 NUCKS_HUMAN
8068	KAEPSEVDMNSPK2000	sp Q9NR30 DDX21_HUMAN
8069	LSPDPVAGSAVSQELR2000000000000000	sp Q9BUR4 WAP53_HUMAN
8070	LNDPFQPPFGNDSPK2000	sp P42566 EPS15_HUMAN
8071	RPISDDDCPSASK20000000000	sp Q96PU4 UHRF2_HUMAN
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8073	ARLSYSDK200000	sp Q05655 KPCD_HUMAN
8074	QQDLHLESPQRQPEYSPESPR200000002000000	sp Q9H6F5 CCD86_HUMAN
8075	TSVATQCDPEEIIVLSDS2000000000000200	sp Q9UER7 DAXX_HUMAN
8076	DSLGTYSR200000000	sp Q9C0C2 TB182_HUMAN
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8078	SLHSAHSLASR200200200000	sp O15027 SC16A_HUMAN
8079	WGQPPSPTPVPRPPDADPNTSPKPLEGRPER2E+27	sp Q14839 CHD4_HUMAN
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8082	RRTSPPPR20200000	sp Q8IYB3 SRRM1_HUMAN
8083	NDLQDTEISPR2000	sp Q9NS91 RAD18_HUMAN
8084	AGSPLCNSNLQDSEEDIEPSYSLFR2E+23	sp Q9NRY4 RHG35_HUMAN
8085	YGHKSPHFR20200000	sp Q86U06 RBM23_HUMAN
8086	QPLLLSEDEEDTK200000000	sp B5ME19 EIFCL_HUMAN
8087	TRDDGDEEGLLTHSEEELEHSQDTDADDGALQ2002000000000	sp Q99613 EIF3C_HUMAN
8088	SSSKDRPSQAAGDNQDEAK2.2E+20	sp Q9UKM9 RALY_HUMAN
8089	FHSPSTWSPNKDTPQEK20002000000000000	sp Q9Y2W1 TR150_HUMAN
8090	AQSSPAAPASLSAPEPASQAR2E+19	sp Q99590 SCAFB_HUMAN
8091	SPSVSSPEPAEK20220000000	sp Q8WUI4 HDAC7_HUMAN
8092	AGLGSPPERPKTSPGSPR200000020002000	sp Q9UQ35 SRRM2_HUMAN
		sp Q9H6F5 CCD86_HUMAN

8093	LSDGDMTSALR2000000000	sp Q8WXE1 ATRIP_HUMAN
8094	SEGFMDR200000000	sp Q7Z4V5 HDGR2_HUMAN
8095	FQSQADQDQASGLQSPSR200200	sp Q9P2B4 CT2NL_HUMAN
8096	RPACDPEPGESGSSSDEGCTVVRPEK2022000000000000	sp O15541 R113A_HUMAN
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8101	SLSDSEDDSK2000200000	sp Q13185 CBX3_HUMAN
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8103	SPLAQMEER20000000000	sp Q16181 SEPT7_HUMAN
8104	DGDSYDPYDFSDTEEEMPQVHTPK300200000100002000	sp P25205 MCM3_HUMAN
8105	HGGSPQPLATTPLSQEPVNPPEASPTR2002000000020000000	sp Q9UQ35 SRRM2_HUMAN
8106	ILDNTGEPAPVLSSPPADVSTFLAFPSPEK20000	sp P46060 RAGP1_HUMAN
8107	GPEENYSRPEAPNEFYDGDNDKESDVEI200000	sp O15379 HDAC3_HUMAN
8108	DYGHSSSRDDYPSR200000000	sp P38159 RBMX_HUMAN
8109	QPYPSPPFNDQHSQDLDSR20000000	sp Q96E39 RMXL1_HUMAN
8110	AQSPSSGDEPEPSGK2022000000000000	sp Q07157 ZO1_HUMAN
8111	ETESAPGSPR2000002000	sp Q9HCM7 FBSL_HUMAN
8112	LQTTDNLPMSPPEEFDEVSRI20000000000	sp Q6PKG0 LARP1_HUMAN
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8114	SLSPILPGR2020000000	sp P49790 NU153_HUMAN
8115	SIEDLHR20000000	sp Q6ZSZ5 ARHG1_HUMAN
8116	AMVDSQQKSPVK20000	sp Q96PY5 FMNL2_HUMAN
8117	SESPKEPEQLR202000000000	sp Q03001 DYST_HUMAN
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8121	TEMDKSPFNPSQDSR2000000000020000000	sp Q9UPN6 SCAF8_HUMAN
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8127	ASESSSEKDDYEIFVK22000000000000	sp Q5VZL5 ZMYM4_HUMAN
8128	ESGVVAVSPEK20000	sp P18583 SON_HUMAN
8129	DKDDQEWESPPKPTVFISGVIAR202000000000000000	sp Q96T58 MINT_HUMAN
8130	RPPTPEAQSEER20000200000	sp P51397 DAP1_HUMAN
8131	QADPQPQAWPPDSPVR20000	sp Q9BTK6 PAGR1_HUMAN
8132	RPSQGPPAR20000000	sp Q63ZY3 KANK2_HUMAN
8133	SLHSAHSLASR200200200200	sp Q9COC2 TB182_HUMAN
8134	RQSVSPPYKEPSAYSSTR2000030000000000000	sp O15027 SC16A_HUMAN
8135	RSPSPAPPPR20000000	sp Q9NYV4 CDK12_HUMAN
8136	IEIIQPLLDMAAGTSNAAPVAENVTNNEGSPPPPVK20000000	sp Q8IYB3 SRRM1_HUMAN
8137	AGASIPSSGAASPR20000000	sp O95104 SFR15_HUMAN
8138	SSPEQPIGQGR20000000000	sp Q9P270 SLAI2_HUMAN
8139	KGGYSQAASSDSAQSDVSLTACK20002000000000	sp Q8N201 INT1_HUMAN
8140	RLSSSESPQRDPPPPPPPLLR2.202E+20	sp P05534 1A24_HUMAN
		sp Q09160 1A80_HUMAN
		sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
		sp Q9H2G4 TSYL2_HUMAN

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8142	ASHTLLPSHR2000000000	sp Q9UKV3 ACINU_HUMAN
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8144	ELSVQDQPSLSPTSLQNSSHTTTAK2000000000000000	sp Q95628 CNOT4_HUMAN
8145	EVEEDESEDEEMSEDEEDSSGEEVVIPQKK2.00000200000002E+25	sp P19338 NUCL_HUMAN
8146	SAMDSPVPASMFAPESPSPAAR10000002000000	sp Q00178 GTPB1_HUMAN
8147	SSTPSSPTGTSSSDSGHHIGWGER2.02E+23	sp P46019 KPB2_HUMAN
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8150	SPPPVS20000000	sp Q01130 SRSF2_HUMAN
8151	IDTHSPSHSSTVK2000000000	sp Q8NI27 THOC2_HUMAN
8152	TLVHSSSDGHIDPQHAAGK20000000000000	sp Q8N5C8 TAB3_HUMAN
8153	ERSPSPLR2000000	sp Q9Y2V2 CHSP1_HUMAN
8154	SGSDAGEARPPTPASPR2000000000002000	sp Q96FS4 SIPA1_HUMAN
8155	QSPSPSEVSQSPGR20000	sp P55201 BRPF1_HUMAN
8156	RESESEDETTPAAPQLIKK202000000000000000	sp Q060832 DKC1_HUMAN
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8159	TPEPSSPVKEPPPVLAKPK2000000000000000	sp Q86TC9 MYPN_HUMAN
8160	YISQEEGPLSPFLGQLDEDYR20000000000000	sp Q5VUA4 ZN318_HUMAN
8161	SSLSSHSHQSQIYR2202000000000000	sp Q015027 SC16A_HUMAN
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8163	DCNESTDNDELEDEPEEPFYR220000000000000000	sp Q9Y2K1 ZBTB1_HUMAN
8164	AGLSPAPELLSR2000000000	sp Q095402 MED26_HUMAN
8165	ISQVSSGETEYNPTEAR200000000000000000	sp Q6ZJ1 NBEL2_HUMAN
8166	LGGDEDEGEEEEEDENSRSQSEDDSETEKPEADDPK20200000000000	sp Q9UHR5 S30BP_HUMAN
8167	ASPVTSAAAAPTASPANK20000000000000000000	sp Q9UIF9 BAZ2A_HUMAN
8168	DLVLPQALPASPALK200000	sp Q15554 TERF2_HUMAN
8169	SRSYSPAHNR20202000000	sp Q9Y2W1 TR150_HUMAN
8170	MQAGQISVQSSEPSSPEPGK200002000000	sp P37275 ZEB1_HUMAN
8171	SRDESASETSTPSEHSAAPSPQVEVR200000000000000000	sp Q92614 MY18A_HUMAN
8172	SLDGAEFRPASVSENHDAGPDGDKR2.0000000002E+26	sp Q9H4G0 E41L1_HUMAN
8173	SLSPPHKR202000000	sp Q5VV67 PPRC1_HUMAN
8174	KRPACDPEPGESGSSSDEGCTVVRPEK20022000020000000	sp Q015541 R113A_HUMAN
8175	KESKEETSIDVAGKPNEVTK2.00002E+19	sp P53985 MOT1_HUMAN
8176	RREEGPPPPSPDGASSDAEPEPPSGR200002200000000000	sp Q9NTJ3 SMC4_HUMAN
8177	FFTTGSDSESESSLGEELVTKPVGGNYGK2.02022E+23	sp B5ME19 EIFCL_HUMAN
8178	LSNSDSQDIQGR200000000000	sp Q99613 EIF3C_HUMAN
8179	HSSSPLSPGGPHTLTKPTTSSSER2.002E+22	sp Q9Y618 NCOR2_HUMAN
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8181	VPVNLLNSPDCDVK20000000	sp P33981 TTK_HUMAN
8182	RPESPSEISPIKGSVR20200000000000	sp Q7Z5K2 WAPL_HUMAN
8183	RGEGDAPFSEP GTTSTQRPSSPETATK200000000	sp Q9UQ35 SRRM2_HUMAN
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8186	EAGSPAQEFK20000000	sp Q5JSZ5 PRC2B_HUMAN
8187	SSDYQFPSSPFTDTLK2000000000000000	sp Q9UK61 F208A_HUMAN
8188	AAPEASSPPASPLQHLLPGK2200000000000000	sp Q96TA1 NIBL1_HUMAN
8189	IEEVLSPGSPSKSPSK2000200000200	sp Q9UEY8 ADDG_HUMAN
8190	TASGSSVTSLDGTR22022000000	sp Q92597 NDRG1_HUMAN
8191	AGPTLTDENGDGLPPSPGDTSYQDQVDDFHEAR2000000000000000	sp Q9NQZ2 SAS10_HUMAN
8192	SYLEGSSDNQLK20000000	sp Q9UHI6 DDX20_HUMAN
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8195	GSSDGRGSDSEDLPHR20000000200000000	sp Q9UPQ0 LIMC1_HUMAN
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8198	SDKSPDLAPTPAQSTPR200000000000000000	sp Q9BY44 EIF2A_HUMAN
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8201	GEIKDSSPSSSPK20000202000	sp Q6WKZ4 RFIP1_HUMAN
8202	SYLEGSSDNQLKDESETPVDDR2.000022E+22	sp Q9UHI6 DDX20_HUMAN
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8204	HFKDEDEDEDVASPDGLGR20000000	sp 095365 ZBT7A_HUMAN
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8211	SPTEDFR20200000	sp Q8IXT5 RB12B_HUMAN
8212	SSSSAAASTPPPGPPADPLGYLPLHGGYQYK2.22E+32	sp Q8NDT2 RB15B_HUMAN
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8214	MPLLELGGETTPPLSTERSPEAVGSECP SR2E+21	sp Q9Y4K1 AIM1_HUMAN
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8216	AAKPGPAEAPSPTASPSGDASPPATAPYDPR20000000000	sp Q9UPT8 ZC3H4_HUMAN
8217	SPTPKSPPSR20000200000	sp Q9NVD7 PARVA_HUMAN
8218	SPQRSPLQAESSPTAGK200020020000000	sp Q8IWB9 TEX2_HUMAN
8219	HRSSDSDLSPPR22020020000	sp Q9BRD0 BUD13_HUMAN
8220	SPSPDGSPAATPEIR2000000000	sp 000499 BIN1_HUMAN
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8231	SLGSASPGPGQPPLSSPTR2E+19	sp Q9C0B5 ZDHC5_HUMAN
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8238	MVSQSQPGSR2000200	sp 075122 CLAP2_HUMAN
8239	EVTPESGPPK200000000	sp P46087 NOP2_HUMAN
8240	SSPELTR2000000	sp Q9UQ35 SRRM2_HUMAN
8241	LPPPPGSPGLHSPTASPPPTAR20000200000000000	sp 095785 WIZ_HUMAN
8242	SRTASGSSVTSLDGTR20202020000000000	sp Q92597 NDRG1_HUMAN
8243	AVSPPHLDGPPSPRSPVIGSEVFLPNSNHV ASGAGEAER2.000000002002E+38	sp P29590 PML_HUMAN
8244	MANSSPVLSK2000000	sp Q9NXE8 CWC25_HUMAN
8245	RSPSVSSPEPAEK2020200000000	sp Q9UQ35 SRRM2_HUMAN
8246	APPAPLAASEVTNSNSAER20000	sp Q9Y2U8 MAN1_HUMAN

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8255	GRASSHSSQTQGGGSVTK2200000000000000	sp P02545 LMNA_HUMAN
8256	GRSPSQYHDFAEK2000000000000000	sp Q9UGU0 TCF20_HUMAN
8257	SCFESSPDPELK20002200000000	sp Q9UQ35 SRRM2_HUMAN
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8263	ELSNSPLRENSFGSPLEFR2000000000002000000	sp Q9UQ35 SRRM2_HUMAN
8264	IPLKPSPSK200	sp Q99755 PI51A_HUMAN
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8267	HDSPDPSPPRR2000200000	sp P46379 BAG6_HUMAN
8268	GKYSPTVQTR20000000	sp Q9BRD0 BUD13_HUMAN
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8271	VSTVTSPSR200	sp Q9UHJ3 SMBT1_HUMAN
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8273	NNSNTCNIENELEDNRK2000000000002000	sp P78332 RBM6_HUMAN
8274	GGSYSQAASSDSAQGSVDVSLTACK3E+21	sp Q6PL18 ATAD2_HUMAN
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8276	VAESAAAAPQGGPELPPSPAPPEQPAPAPER2E+29	sp P30447 1A23_HUMAN
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8278	TRHSPTPQQSNR2000000000	sp P01892 1A02_HUMAN
8279	SPSRPHPGGHPHSSPLAPHPISGDR2.0000000002E+23	sp Q09160 1A80_HUMAN
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8284	LSSLRASTKSESSQK2202000000	sp Q8IYB3 SRRM1_HUMAN
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8286	KNSITEISDNEDDLLEYHR200200000000000000	sp Q6PL18 ATAD2_HUMAN
8287	YSISLSPPEQQK20000000	sp Q8IZP0 ABI1_HUMAN
8288	DLHSDNNEEESDEVEDNDNDSEMERPVNR2.0000002E+26	sp P49674 KC1E_HUMAN
8289	KSSVSDAPVHITASGEPVPISEESEELDQK2E+28	sp Q13459 MYO9B_HUMAN
8290	GPPSPAPVMHSPSR2000001000000	sp P62753 RS6_HUMAN
8291	LEQTSVRDPSPEADAPVLGSPEK2000000000000000	sp Q684P5 RPGP2_HUMAN
8292	LPEPSVLSEVTKPPPCLENSPPSR2000000	sp Q86UU1 PHLB1_HUMAN
8293	SPSPPLPHIPPEPRTPPFPAK2.02E+23	sp Q15057 ACAP2_HUMAN
8294	DSTSQHDDNISTTSGFSSRASDKDITVSK22200220000000000000	sp P42568 AF9_HUMAN
		sp O14646 CHD1_HUMAN
		sp Q13573 SNW1_HUMAN
		sp O15234 CASC3_HUMAN
		sp Q9BZL4 PP12C_HUMAN
		sp Q8IZ21 PHAR4_HUMAN
		sp P49790 NU153_HUMAN

8295	QPGVDSLSPVASLPK20000000000	sp Q9UBW7 ZMYM2_HUMAN
8296	KTSPASLDFPESQK200000000000	sp Q96JM3 CHAP1_HUMAN
8297	HSTPSNSSNPSGPPSPNSPHR200020020000	sp Q9Y5S2 MRCKB_HUMAN
8298	GTGGSESSRASGLVGR2002002000000	sp Q9UH99 SUN2_HUMAN
8299	RASPSPPPK20200000	sp Q8IYB3 SRRM1_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P05534 1A24_HUMAN
8300	GGSYSQAASSDSAQGSVDVSLTACK2000000000	sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
		sp P30447 1A23_HUMAN
8301	KAPSASDSKADSDGAKPEPVAMAR2.0200020002E+23	sp Q7Z4V5 HDGR2_HUMAN
8302	VSPSTSYTPSR20000000000	sp Q9BTE3 MCMBP_HUMAN
8303	SGSIVELIAGGSSCSPVLSR2000000	sp P42858 HD_HUMAN
8304	GPIHSPVELK2000000	sp Q16649 NFIL3_HUMAN
8305	VQTTPPPAVQGQK20000000000	sp Q2M2I8 AAK1_HUMAN
8306	LRSPSNDSAHR20000000	sp Q5T200 ZC3HD_HUMAN
		sp P20020 AT2B1_HUMAN
8307	SSLYEGLEKPESR2000000000000	sp Q01814 AT2B2_HUMAN
		sp Q8TAD8 SNIP1_HUMAN
8308	RPDHSGGSPSPTSEPAR200000000000000	sp Q13459 MY09B_HUMAN
8309	VSPAPGSAPETPEDK2000002000000000	sp Q6GYQ0 RGPA1_HUMAN
8310	SQTPSPSTLNIDHMEQK202020000000010000	sp Q9Y2W1 TR150_HUMAN
8311	SPPSTGTYGSSQKEESAASGGAAYTK2.002E+27	sp P50402 EMD_HUMAN
8312	RLSPPSSAASSYSFSDLNSTR2.0000000003E+20	sp Q15345 LRC41_HUMAN
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8314	SRTHSTSSSLGSGESPF222220000000000000	sp Q6VMQ6 MCAF1_HUMAN
8315	QDDDLNCEPLSPHNITPEPVSK20000000000000	sp Q9UHB7 AFF4_HUMAN
8316	SRSSSPGKQAVSSLNSSH22000000000000000000	sp 075420 PERQ1_HUMAN
8317	SLSVPDSGR20000000	sp 060832 DKC1_HUMAN
8318	KRESESEDETPPAAPQLIK2000200200000000000	sp Q8NFA0 UBP32_HUMAN
8319	SPSSLSANI ISSPK200000000020000	sp Q9UQ35 SRRM2_HUMAN
8320	SGSSQELDVKPSASPQER2022000000020000000	sp Q8WYA6 CTBL1_HUMAN
8321	EYAENIGDGRSPEFR200000	sp Q5VTR2 BRE1A_HUMAN
8322	SGSALLQSSTEDPKDEPAELKPDSEDLSSQSSASK2.2E+27	sp 060583 CCNT2_HUMAN
8323	QGQSQAAASSSVTSPIK200000000	sp Q96D71 REPS1_HUMAN
8324	SSSLDMNR2000000	sp Q92597 NDRG1_HUMAN
8325	SRTASGSSVTSLDGTR22000000000	sp P49810 PSN2_HUMAN
8326	TSLMSAESPTPR200200200000	sp Q9BXS6 NUSAP_HUMAN
8327	GRLSVASTPISQR20002000000	sp 094842 TOX4_HUMAN
8328	LSTTPSPTSSLHEDGVEDFR20220000000000000	sp 075367 H2AY_HUMAN
8329	AASADSTTEGTPADGFTVLSTK20020000000000000	sp Q9NUW8 TYDP1_HUMAN
8330	WTISSSESEEEKPKPKPSTSLLCAR2.0222E+27	sp Q8WYP5 ELYS_HUMAN
8331	STPLASPSPPGRSPQR200020000	sp Q92609 TBCD5_HUMAN
8332	EFTGSPSSATK20000000000	sp Q6ZRV2 FA83H_HUMAN
8333	RSQSSSQGDNPLAPGYLPPHYK2.22E+19	sp Q86VM9 ZCH18_HUMAN
8334	DSGSDQDLGAGVR2000000000000	sp P11388 TOP2A_HUMAN
8335	FTMDLSDSEDFDFDEKTDDEDFVPSDASPPK1.000200002E+30	sp Q96I25 SPF45_HUMAN
8336	SPTGPSNSFLANMGTVAHK2000000000100000000	sp 076021 RL1D1_HUMAN
8337	AAESETPGKSPEKKPK20000000	sp P62753 RS6_HUMAN
8338	LSSLRASSTKSESSQK22202000000	
8339	LSSDENSNPDLSGDENDDAVSIESGTNTERPDTPTNPNAPGR2.000000002E+32	sp 075530 EED_HUMAN
8340	LSPQMQRVHDK20000000000	sp Q12830 BPTF_HUMAN
8341	SAGGSSPEGGEDSDREDGNYCPPVK2.2E+21	sp Q9H6Z4 RANB3_HUMAN

8342	APSASDSKADSDGAKPEPVAMAR2.0202000002E+23	sp Q7Z4V5 HDGR2_HUMAN
8343	SFSKEVEER2020000000	sp Q04637 IF4G1_HUMAN
8344	NSSTPGLQVPVSPTVPIQNQK2.2E+19	sp Q03164 KMT2A_HUMAN
8345	SSSVGSSSSYPISPAVSR220002000000	sp Q15149 PLEC_HUMAN
8346	SLVHEVGKPPQDVTDDSPSK200000000	sp Q96T58 MINT_HUMAN
8347	KRESESEDETPPAAPQLIK200000000000000000	sp 060832 DKC1_HUMAN
8348	LVKPSSSSSGSR20000	sp Q9H6B4 CLMP_HUMAN
8349	CQTAEADSESDHEVPEPESEM2020000000000100	sp Q9BPX3 CND3_HUMAN
8350	DEPPPASQSTSQDCSQALKQSP200	sp Q5JRA6 MIA3_HUMAN
8351	RFSFSQR200000	sp Q92614 MY18A_HUMAN
8352	ADTEGKSPSHSSSNRPFTPPTSTGGSK2.02E+21	sp Q15648 MED1_HUMAN
8353	LSPPHSPR20002000	sp Q92974 ARHG2_HUMAN
8354	LPHSSSENK2000000	sp Q76I76 SSH2_HUMAN
8355	SPQGSPVLAEDSEGEG2000000200000	sp Q99638 RAD9A_HUMAN
8356	SFSSSAEEHS200000000	sp Q96EQ0 SGTB_HUMAN
8357	DNTFFRESPVGR200000	sp Q8NEY8 PPHLN_HUMAN
8358	RSSGFISELPSEEGK20000000000000	sp Q5VZK9 LR16A_HUMAN
8359	IEEAMDGSETPQLFTVLPEK200000000000	sp Q13435 SF3B2_HUMAN
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8361	NSSLGSPNLCGSPPGSIR20000000	sp Q9C0B0 UNK_HUMAN
8362	ETSVSKEDTDHEEKASNEVDTK2.0000000000002E+21	sp 075475 PSIP1_HUMAN
8363	LTRPAASPAVGEK20000000	sp Q8N2M8 CLASR_HUMAN
8364	RLPSSPASPSPK2000200000	sp Q9H4G0 E41L1_HUMAN
8365	SPISELHSAPLTPVAR200200000000000000	sp Q7Z3B3 KANL1_HUMAN
8366	ATVTPSPVK2020000	sp Q9H1E3 NUCKS_HUMAN
8367	NGSLDSPGKQDTEDEEDEKDK200000200000000000	sp Q9Y266 NUDC_HUMAN
8368	SCDELSPVSPTQGGYPSEPTR2000000000000000	sp Q9HCD6 TANC2_HUMAN
8369	SPQRSPLQSAESSPTAGK200020000000000000	sp Q8IWB9 TEX2_HUMAN
8370	TEEARPSPAPGPGTPTGTPTTR200020000	sp Q4KMP7 TB10B_HUMAN
8371	SSPINPAQSGSPNHVDSAYFPGSSTSSSSDNDEGSGGATNHISGNK2E+45	sp Q9NZB2 F120A_HUMAN
8372	NTFTAWSDEESDYEIDDRDVK2.0000002E+19	sp Q6PKG0 LARP1_HUMAN
8373	GDTPGHATPGHGATSSAR20000000000000	sp 075533 SF3B1_HUMAN
8374	RPSVASSVSEEFYFEVR2002202000000000	sp A6ND36 FA83G_HUMAN
8375	SRPLATGPSSQSHQEK20000000	sp Q96RL1 UIMC1_HUMAN
8376	NTPSQHSHSIQHSPER2000000200020000	sp Q9NYF8 BCLF1_HUMAN
8377	TSFDQDSVDVIFPSDFPTEPPSLPR2E+24	sp Q02880 TOP2B_HUMAN
8378	KVDEGAGDAAVASGGAQTLALAGSPAPSGHPK2000000000	sp Q9NZT2 OGFR_HUMAN
8379	SAPASPTHPLMSPR2000200000012000	sp P85037 FOXK1_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
8380	GGSYSQAASSDSAQGSVDVSLTACKV2000200000000000	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
8381	SASSNSRKSLSPGVSR202202002020000000	sp Q9UKV3 ACINU_HUMAN
8382	HADGEKEDQFNGSPPRPQPR2000000000	sp Q69YN4 VIR_HUMAN
8383	RSSIGLR200000	sp Q96GN5 CDA7L_HUMAN
8384	HSDSGTSEASLSPSSPPSRPR200200000000	sp Q7Z4S6 KI21A_HUMAN
8385	ERPTSPAPHRPPK2000000000	sp Q8N6T7 SIR6_HUMAN
8386	ETPHSPGVEDAPIAK2000000000000000	sp Q9UHB6 LIMA1_HUMAN
8387	DFLSGPDYDHVTGSDSEDEEVEGEP3.00000002E+20	sp Q6ZW31 SYDE1_HUMAN
8388	SSGHSSSELSPDAVEK220002200200000000	sp Q9UQ35 SRRM2_HUMAN
8389	ASPSPPPKR202000000	sp Q8IYB3 SRRM1_HUMAN
8390	AGLSPANCSDR2000000000	sp Q14692 BMS1_HUMAN

8391	VSDSESEGPQK2020000000	sp Q96ST2 IWS1_HUMAN
8392	KAPSASDSDSKADSDGAKPEPVAMAR2.0202020002E+23	sp Q7Z4V5 HDGR2_HUMAN
8393	SISLGDSEGPVATLAQPLR2E+20	sp O43379 WDR62_HUMAN
8394	GGPGKLSR2000	sp Q12888 TP53B_HUMAN
8395	TSPTTPEASATNSPCTSKPATPAPSEK2000000000000000	sp Q12873 CHD3_HUMAN
8396	HKSGSMEEDVDTSPGGDYITSPSSPTSSSR30000020000000	sp P08651 NFIC_HUMAN
8397	ILEGPQQSLSGSPLQPIPASPPPQIPPGLRPR2.00000002E+21	sp O15446 RPA34_HUMAN
8398	GSSDGRGSDSEDLPHR200000000	sp Q9UPQ0 LIMC1_HUMAN
8399	SDELLGSDSDSHDGESESNAK200200002000000	sp P38398 BRCA1_HUMAN
8400	DWDKESDGPDDSRPESASDSDT200000000020200000	sp Q9Y3B9 RRP15_HUMAN
8401	HQYSDYDYHSSSEK202000	sp Q9UDY2 ZO2_HUMAN
8402	ESPSPAPKPR2020000000	sp Q8IYB3 SRRM1_HUMAN
8403	GDAASSPAPAASVGSSQGGAR200000000000000000	sp Q8WVB6 CTF18_HUMAN
8404	LSSPIDMTSK2000000000	sp Q16649 NFIL3_HUMAN
8405	DDSHAEDSEDEKEDHK2020002000000000	sp Q9H1E3 NUCKS_HUMAN
8406	DGSDYDPYDFSDTEEMPQVHTPK2.0003E+21	sp P25205 MCM3_HUMAN
8407	SPQTLAPVGEDAMKTPSPAAEDAREPEAK102020000000000000	sp Q92797 SYMPK_HUMAN
8408	SPSVSPEPAEK20202000000000	sp Q9UQ35 SRRM2_HUMAN
8409	QQAAYYAQTSPQGMPPHPPAPQGQ2000100000000000	sp Q96AE4 FUBP1_HUMAN
8410	LSEDEEEDEAEDDQSEASGKK2.20000000000002E+21	sp Q8NEJ9 NGDN_HUMAN
8411	FSPGAPGSGSQPNQK2000000200000000	sp Q15942 ZYX_HUMAN
8412	LSPERSDSESNENEPHLPHGVSQR2.000000002E+24	sp Q9BQ52 RNZ2_HUMAN
8413	SSSKGSVEEIMSQPK2200200000000000	sp Q8IWB9 TEX2_HUMAN
8414	SRSPSPELNNK20202200000000	sp Q9UQ35 SRRM2_HUMAN
8415	ARSRTPPSAPSQSR20200000000000	sp Q9UQ35 SRRM2_HUMAN
8416	YTPTSPSYSPSSPEYTPTSPK3.00000000002E+21	sp P24928 RPB1_HUMAN
8417	KPPAPSPVQSQSPSTNWSPAVPVKK2.000202E+20	sp Q8IYB3 SRRM1_HUMAN
8418	DLESCDDDNQGSKSPK202000000002000	sp Q5T3I0 GPTC4_HUMAN
8419	SHLSQHTATSSK2000000000	sp P49790 NU153_HUMAN
8420	SPSESPQHIAHPER20000000000000	sp Q68DK2 ZFY26_HUMAN
8421	TIDLGAAHYTGDKASPDQNASTHTPQSSVK2000000000000000	sp Q14677 EPN4_HUMAN
8422	TNSDSALHQSTMTPTQPESFSSGSQDVHQK2.0000001E+26	sp Q6UUU9 CRTC1_HUMAN
8423	KADTEGKSPSHSSNRPFPTSTGGSK2.02E+21	sp Q15648 MED1_HUMAN
8424	LVEDERSDR2000	sp Q96G23 CERS2_HUMAN
8425	RRSDEDAAGEPK220000000000	sp Q9BUI4 RPC3_HUMAN
8426	ASPEENNAGGGSPSSGGNPTNTSGTTR200000000000000000	sp Q9NRH2 SNRK_HUMAN
8427	GPSPAAASPEGSPLR200020000	sp Q9P107 GMIP_HUMAN
8428	QLSSTSPLAYPYSQMVSSDR200000000000000000	sp Q6UUU7 CRTC3_HUMAN
8429	SDSYVELSQYR2000000000	sp P52298 NCBP2_HUMAN
8430	SQDATFSPGSEQAEKSPGPIVSR2000002000000000	sp Q86WB0 NIPA_HUMAN
8431	SPISPSELHSAPLTPVAR2000000000000200000	sp Q7Z3B3 KANL1_HUMAN
8432	SSPSARPPDVPGQPQAAK200000000000000000	sp Q96JP5 ZFP91_HUMAN
8433	GGNVFAALIQQSEEEEEEEKHPPKPAKPEK2E+19	sp Q8NE71 ABCF1_HUMAN
8434	HNFLFEDNMALPSESVSLLTDLKPPTGSNQASPAR2.02E+23	sp Q9BZQ8 NIBAN_HUMAN
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8436	YHGHSMSDPGVSYR2000000000	sp P08559 ODPA_HUMAN
8437	RPTETNPVTSNSDEECNETVK20020000000000	sp P29803 ODPAT_HUMAN
8438	DPNSATATAPPSPLKR200000	sp P46100 ATRX_HUMAN
8439	DLELPSQEAPSFQGTESPKPCK2000000	sp Q92766 RREB1_HUMAN
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8441	MQAGQISVQSSEPSSPEPGK2000000	sp Q92560 BAP1_HUMAN
8442	SPLQSAESSPTAGK200000002000000	sp P37275 ZEB1_HUMAN
8443	FHSSDSEEEHK2020000000	sp Q8IWB9 TEX2_HUMAN
		sp Q96ST2 IWS1_HUMAN



8444	ESSLPSK200000	sp 075477 ERLN1_HUMAN
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8448	SQSWEER200000	sp P05388 RLAO_HUMAN
8449	ALSGGGSVASPPSPAMPK20002000000	sp Q6Y7W6 PERQ2_HUMAN
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8454	SPTVLCQK2000000	sp Q8NFG4 FLCN_HUMAN
8455	ASAGTPSLSAGVSPK2000	sp Q5VUA4 ZN318_HUMAN
8456	QLPALDGLMGPESPPAQEEEEAPVSPHK1.0002E+19	sp Q8TEK3 DOT1L_HUMAN
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8458	TLENSPISSCDTSDTEGVPVNSAAVLK2.200220E+21	sp Q14684 RRP1B_HUMAN
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8466	SRSPYSPVLR20203000000	sp Q01082 SPTB2_HUMAN
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8468	SLVESVSSSPNK20000	sp Q9BQG0 MBB1A_HUMAN
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8472	SRYRSPYSR20000200	sp Q9BW71 HIRP3_HUMAN
8473	EPTSVSENEGISTIPSPVTSPLSR2E+23	sp Q9BRL6 SRSF8_HUMAN
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8475	NPSDSAVHSPFTK200000	sp Q08495 DEMA_HUMAN
8476	LSGEHSESSTPRPR2000000000	sp Q14157 UBP2L_HUMAN
8477	QSPGSTSPKPPHTLSR2000000000000000	sp Q7Z3K3 POGZ_HUMAN
8478	EEAPASPLRPLYPQISPLK300020000	sp 043159 RRP8_HUMAN
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8481	FELLTPPLSPSR200020000	sp Q9BY44 EIF2A_HUMAN
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8483	VELKSEANDAVNSSTK2000000020000	sp P04198 MYCN_HUMAN
8484	SESSDFEVVPK202000000000	sp Q8WXI9 P66B_HUMAN
8485	EGAASPAPETPQPTSPETSPK20002000	sp 015234 CASC3_HUMAN
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8487	TSPLKDNPSPEPQLDDIKR2.2E+19	sp Q9Y3X0 CCDC9_HUMAN
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		sp Q96JC9 EAF1_HUMAN
		sp Q9P1Y5 CAMP3_HUMAN

		sp P30504 1C04_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
8489	GGSCSQAASSNSAQGSDESLIACK2E+20	sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
8490	AKPSPAPPSTTTAPDASGPQK200000000000000000	sp P40855 PEX19_HUMAN
8491	SPAPGLASGDRPPSVSSVHSEGCNR2E+26	sp Q9Y618 NCOR2_HUMAN
8492	VSITSEVESTSNPSSSSLQK202000000000	sp Q04656 ATP7A_HUMAN
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8494	DDPSDVTDEDEGPAEPPPPKLPAPFR2.002E+25	sp Q3YEC7 RABL6_HUMAN
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8496	SSTPSSPTGTSSSDSGGHHIGWGER2E+25	sp P46019 KPB2_HUMAN
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8499	GNIETTSDEGQVFSFK2000	sp Q5T200 ZC3HD_HUMAN
8500	SPSESSPQHPTPPARPR200020000000000000	sp O14733 MP2K7_HUMAN
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8505	DLSTSPKPSPIPSPVLR2000000002000000	sp Q8NDI1 EHBP1_HUMAN
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8508	SITSTPLSGK20002000000	sp Q9P0K7 RAI14_HUMAN
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8510	LSSPPLR200000	sp Q96ET8 TV23C_HUMAN
8511	FAFSPLSEEEEEDEQKEPMLK20000000000000000000	sp Q8NBN3 TM87A_HUMAN
8512	ISLEQPPNGSDTPNPEK2000000	sp Q9UHI6 DDX20_HUMAN
8513	SSSKDSRPSQAAGDNQGDEAK2E+19	sp Q9Y2W1 TR150_HUMAN
8514	ICSIYTSQGENSLVQEGSEASPIGK200000	sp Q9Y4W2 LAS1L_HUMAN
8515	LASELSFDFR200000	sp Q9Y4K4 M4K5_HUMAN
8516	AKSPTSPSPPRNSDQEGGGK2.020202E+19	sp Q8IYB3 SRRM1_HUMAN
8517	THSTSSSLGSGESPFRR200000000000	sp Q9UGV2 NDRG3_HUMAN
8518	SLGGESSGTTVPVGSFHTEAAR200200000000000000	sp O15085 ARHGB_HUMAN
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8520	ALMDEEDMDDVVDADAYLIPQQGFFSSPSTSR2000000	sp P00533 EGFR_HUMAN
8521	TVGTPIASVPGSTNTGTVPGSEK2E+23	sp Q99460 PSMD1_HUMAN
8522	TPTSSPASSPLVAK20022000000	sp Q14684 RRP1B_HUMAN
8523	SRPTSEGSDIESTEPQK2200000000000000	sp Q4G0J3 LARP7_HUMAN
8524	KHSPSPPTPTESR2000000000000000	sp Q92922 SMRC1_HUMAN
8525	EAASSSGTQPAPPAPASPWDSSK20000000	sp Q9H7N4 SFR19_HUMAN
8526	GESAADSDGWSAPSDLR200000000000000000	sp Q9BUL5 PHF23_HUMAN
8527	TSAVSSPLLDQQR20000000000	sp Q5T200 ZC3HD_HUMAN
8528	SGAHSSASPPR20000	sp P18615 NELFE_HUMAN
8529	LDSPTMSR2000000	sp P19634 SL9A1_HUMAN
8530	TDESSPVLSPR20002000	sp Q8NA72 POC5_HUMAN
8531	SRSPTPPSSAGLGSNSAPPIDSR2.002200002E+20	sp Q8IWX8 CHERP_HUMAN

8532	SSTGPEPPAPTPLLAER2000000000000000	sp Q9UQ35 SRRM2_HUMAN
8533	ATHGSNSLPSSAR20000000	sp Q8TDM6 DLG5_HUMAN
8534	ESKEEETSIDVAGKPNEVTK2000000000000000	sp P53985 MOT1_HUMAN
8535	QNSDDEEQQLSDEEK20000000200000	sp Q8WVC0 LEO1_HUMAN
8536	DSSSTDSANEKPALIPR2200000000000000	sp Q9NUQ6 SPS2L_HUMAN
8537	GPQPPTVSPIR20000	sp P46087 NOP2_HUMAN
8538	SPSAISTGTLVSK20000000000000	sp Q9NR48 ASH1L_HUMAN
8539	LTQAEISEQPTMATVVPQVPTSPK10000000002000	sp Q70E73 RAPH1_HUMAN
8540	HLAQASQELQ200000	sp Q8NBP7 PCSK9_HUMAN
8541	SPVSPLAAGIPLPAQLTK2E+19	sp Q96SI1 KCD15_HUMAN
8542	VETVSPSESPK200002000	sp Q99590 SCAFB_HUMAN
8543	KAPSASDSKADSDGAKPEPVAMAR2.0002020002E+23	sp Q7Z4V5 HDGR2_HUMAN
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8545	KSDGACDPSSSDKENSSQIAQDHQK20000000020000000000	sp Q9UIG0 BAZ1B_HUMAN
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8547	GSLAEAVGSPPAATPTPTPTR2000002000000000	sp Q9Y6I3 EPN1_HUMAN
8548	RQSSGSATNVASTPDNR2000200000000000	sp Q7Z460 CLAP1_HUMAN
8549	DPLSSPGGPGSR2000000000	sp Q32P44 EMAL3_HUMAN
8550	SVSSNVASVSPIPAGSK2000000000	sp Q9Y6G9 DC1L1_HUMAN
8551	SNTPESIAETPPAR200000000200000	sp Q9Y4H2 IRS2_HUMAN
8552	KRTPSVGK2020000	sp P46013 KI67_HUMAN
8553	IQNSDDEER20000000	sp Q8WVC0 LEO1_HUMAN
8554	RRSFSISPSR202020000	sp P18583 SON_HUMAN
8555	GSGGSSGDELREDDEPVK2200000000000000	sp Q7Z4V5 HDGR2_HUMAN
8556	YTPTSPSYSPSSPEYTPSPK220030000000	sp P24928 RPB1_HUMAN
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8558	TNSGISSASGGSTEPTTPDSEPAQALLR2000000000000000	sp Q86X10 RLGPB_HUMAN
8559	LQEEQDGGSSDEDR2200000	sp Q14147 DHX34_HUMAN
8560	RKSEDGTPAEDGTPAATGGSQPPSMGR2000000000000000	sp Q9BQG0 MBB1A_HUMAN
8561	AVTTVTQSTPVGPSVPPPEELQVSPGPR2E+27	sp P51610 HCFC1_HUMAN
8562	IFDEVMGCFSDSPQSPTFPEAGHTSLYDEDKVR2.000002E+24	sp Q13085 ACACA_HUMAN
8563	SPPKSPPEEGAVSS2000000000000020	sp Q01130 SRSF2_HUMAN
8564	AVGMPSPVSPK2002000	sp Q9UHB7 AFF4_HUMAN
8565	GSQHSPTRPVAAAAASLGLPGGAAR2E+22	sp Q86VQ1 GLC11_HUMAN
8566	YSPTSPTYSPVYTPSPK3002000000000000	sp P24928 RPB1_HUMAN
8567	TPTPSTPGDTQNPAPVPPAEDGIK2E+24	sp Q14839 CHD4_HUMAN
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8569	LMLSTSEYSQSPK3002000	sp Q12888 TP53B_HUMAN
8570	KFDHESPGETDEKSG220020000200	sp Q13427 PIG_HUMAN
8571	SSSSDSIHSVR200000000000	sp Q8WYL5 SSH1_HUMAN
8572	SLTRSPPAIR20002000000	sp Q9UQ35 SRRM2_HUMAN
8573	DLFETGLEDENNSPLEDEESLESIR2E+21	sp Q9HAW4 CLSPN_HUMAN
8574	VQMTSPSSTGSPMFK10200000000000	sp P49790 NU153_HUMAN
8575	GLNSQSSDHLNKR2200000000	sp A8CG34 P121C_HUMAN
8576	GHVTPNLSR2000200	sp Q96HA1 P121A_HUMAN
8577	TESVSDKEDKPLAPSGGTEGPEQPPPCPSQTGSPPVGLIK2000000000	sp Q96CP6 GRM1A_HUMAN
8578	STGVSWFTQSDENEQEQQSDTEEGSNKK2.000000002E+19	sp Q8WWM7 ATX2L_HUMAN
8579	QLSALSSDGEQGGRPVPR20000000000000	sp O14974 MYPT1_HUMAN
8580	ESEDSETQPFDTHEAYGPCLSPPR2E+24	sp O15438 MRP3_HUMAN
8581	VEPSSQSPGRSPR20000000	sp Q14676 MDC1_HUMAN
8582	VVNITSPGHDAASSRPTTASVSATAAPR22000000000000000000	sp Q8IUD2 RB6I2_HUMAN
8583	SKSEEHAEDSVMDHHR200000000	sp Q15723 ELF2_HUMAN
8584	GDDDEESDEEAVK20000000	sp Q8NC51 PAIRB_HUMAN
		sp O43395 PRPF3_HUMAN

8585	AASLTEDRDR200000000	sp Q04637 IF4G1_HUMAN
8586	IQFKPDDGISPER20000	sp Q96I24 FUBP3_HUMAN
8587	SAGGSSPEGGEDSDREDGNYCPPVK2.000220000002E+25	sp Q9H6Z4 RANB3_HUMAN
8588	AYLGYQAVPSGSSGSLSTSSSSSPPGTPSPADAK2E+25	sp P17676 CEBPB_HUMAN
8589	EHAPLASPVENK2000000	sp Q9ULJ3 ZBT21_HUMAN
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8591	AYLGYQAVPSGSSGSLSTSSSSSPPGTPSPADAK200000000	sp P17676 CEBPB_HUMAN
8592	VESVEPSENEASK200020000000	sp Q9H307 PININ_HUMAN
8593	SDGACDSPSSDKENSSQIAQDHQK2000000002000000000	sp Q9UIG0 BAZ1B_HUMAN
8594	KASVVDPSTESSPAPQEGSEQPASPASPLSSR2000020000000000	sp Q66K14 TBC9B_HUMAN
8595	KEKTPELPEPSVK20000000000	sp Q8IYB3 SRRM1_HUMAN
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8597	RGSSAAASPGSPPPGR200000002000000	sp Q86UX6 ST32C_HUMAN
8598	SSSISEEKGSDDEKPR200000000020000000	sp Q9UKV3 ACINU_HUMAN
8599	SESARGSDSEDEVLR2020000020000000	sp Q8WVC0 LEO1_HUMAN
8600	SRSPSPELNNK20220000000	sp Q9UQ35 SRRM2_HUMAN
8601	EVEEDSEDEEMSEDEEDSSGEEVVIPQKK1.200000022E+20	sp P19338 NUCL_HUMAN
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8603	SVELCCASR2000000000	sp Q9Y4F5 C170B_HUMAN
8604	AASPHQVTPQADEK2000000000000	sp Q96L73 NSD1_HUMAN
8605	KGSSGNASEVSVACLTER200000000000	sp Q69YQ0 CYTSA_HUMAN
8606	SSSPTS YWK20000000	sp Q92614 MY18A_HUMAN
8607	KSSPSVKPAVDPAAAK200000000000000	sp Q6FI81 CPIN1_HUMAN
8608	TVSSPPTSPRPGSAATVSASTSNIIPPR20200000000000	sp 076094 SRP72_HUMAN
8609	ARVSPPPK20200000	sp Q13247 SRSF6_HUMAN
8610	RREEGPPPPSPDGASSDAEPEPPSGR200000000000	sp Q9NTJ3 SMC4_HUMAN
8611	TPLLDEEEENPDK200000000000000	sp P23634 AT2B4_HUMAN
8612	ASLVALPEQTASEEETPPPLLTK200000200000000	sp Q9UKV3 ACINU_HUMAN
8613	GLSDHVS LDGQELGTR20000000000	sp Q7L8J4 3BP5L_HUMAN
8614	SVSHGSNHTQKPDEQR200000000000000	sp Q9Y520 PRC2C_HUMAN
8615	VIYSQPSAR2000	sp Q9Y624 JAM1_HUMAN
8616	RESESEDETPPAAPQLIK2000200000000000000	sp 060832 DKC1_HUMAN
8617	YSPTSPTYSPTTPK200030000000	sp P24928 RPB1_HUMAN
8618	SNSFSDER2000000	sp Q09666 AHNK_HUMAN
8619	ATRPASTSQPDPSPDR20000020000	sp P20749 BCL3_HUMAN
8620	LITPPPPPPSPER20000	sp Q96HE9 PRR11_HUMAN
8621	GAAAADLLSSSPESQHGGTQSPGGGQPLLQPTK2E+23	sp Q9H3S7 PTN23_HUMAN
8622	VDKETNTETPAPSPTVVRPK2000200000000	sp Q8IX03 KIBRA_HUMAN
8623	VIPAKSPPPHSTQLGAPSR20000000000000000	sp Q14814 MEF2D_HUMAN
8624	APSASDSKADSDGAKPEPVAMAR2.0202000002E+23	sp Q7Z4V5 HDGR2_HUMAN
8625	VNLSSIR22000	sp Q96DX4 RSPRY_HUMAN
8626	DGLAPEKTSPDR20000	sp Q7L4I2 RSRC2_HUMAN
8627	GHYEVTGSDDDETGKLQSGVSLASK200000000000000	sp Q09666 AHNK_HUMAN
8628	SRTSPITR200200000	sp Q9UQ35 SRRM2_HUMAN
8629	SRSPAPPADK200000000	sp Q9Y618 NCOR2_HUMAN
8630	QRSLSPVAAPPLR202000000000	sp Q8NDT2 RB15B_HUMAN
8631	RLSSASTGKPLSVEDDFEK2200000000000000	sp 075152 ZC11A_HUMAN
8632	RDSFDNCSLGESSK2000020000000	sp Q5UIP0 RIF1_HUMAN
8633	SPTTVLFTSSPIK200000020000	sp Q2KHR2 RFX7_HUMAN
8634	LPQPPEGQCYSN200	sp Q13404 UB2V1_HUMAN
8635	STPLASPSPPGRSPQR20002000200020000	sp Q8WYP5 ELYS_HUMAN
8636	SPHSPSPATPPSQSSSCSDAPMLSTVHSAK222020000000000000	sp Q9NYF8 BCLF1_HUMAN
8637	VLSGNCNHQEGTSSDDELPSAEMIDFQK20200000000000000	sp P16383 GCFC2_HUMAN

8638	SPGFASPK200002000	sp P49790	NU153_HUMAN
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8640	TPGVLLPGAGGAAGFGMTSPPPPTSPSR200000020000	sp Q96L91	EP400_HUMAN
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8642	TSPQVLGSILK20000000000	sp Q8IY63	AMOL1_HUMAN
8643	QSVTSFPDADAFHHQVHDDLLSSSEEECKDR2200000000	sp Q7Z417	NUFP2_HUMAN
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8645	QGGSQPSSSPGQSQVTPQDQEK2E+20	sp Q76L83	ASXL2_HUMAN
8646	GGQGDPAVPTQQPADPSTPEQQNSPSGSEQFVR200000020000000000	sp Q9H0L4	CSTFT_HUMAN
8647	NTSLPPLWSPEAER2000002000000	sp O15231	ZN185_HUMAN
8648	RDEDMLYSPELAQR20000000	sp P49790	NU153_HUMAN
8649	KDRHDTDPSPR2000	sp Q9Y2W2	WBP11_HUMAN
8650	LSPERSSDSESNENEPHLPHGVSQR2.000200202E+24	sp Q9BRD0	BUD13_HUMAN
8651	SAGGSSPEGGEDSDREDGNYCPPVKR2.000220000002E+26	sp Q9BQ52	RN22_HUMAN
8652	VTSSVPLPSGSTSSPGR20000	sp Q9H6Z4	RANB3_HUMAN
8653	VPSTPPPK200000	sp Q15036	SNX17_HUMAN
8654	SSTVATLQGTPDHGDPR200000000	sp Q9H7N4	SFR19_HUMAN
8655	KRPSLPSSPSPGLPK2000002000000	sp Q8IVT2	MISP_HUMAN
8656	SLESSVIMDR202000001000	sp O43294	TGF11_HUMAN
8657	DGDSYDPYDFSDTEEEMPQVHTPK300200000000002000	sp Q9H2Y7	ZN106_HUMAN
8658	SRESSPSHGLLK2000200000000	sp P25205	MCM3_HUMAN
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8660	MPLAQKPALAPKPTSQTTPASPLSK2000200000	sp Q86VM9	ZCH18_HUMAN
8661	ATSNVFMFDQSQIQEFK2000000000000000000	sp Q6ZU35	K1211_HUMAN
8662	TSSPPSPQHSPALR20002000000000	sp O14950	ML12B_HUMAN
8663	TGSNISGASSDISLDEQYK2002002200000000000	sp P19105	ML12A_HUMAN
8664	KPSDSLVAASSR2020002200	sp P24844	MYL9_HUMAN
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8666	QESCSPHHPQVLAQQGSGSSPK2000	sp P22059	OSBP1_HUMAN
8667	EKEDNSSEEEEEIEPFPEER22000000000000000	sp Q5T0W9	FA83B_HUMAN
8668	STSAPQMSPGSSDNQSSSPQPAQQK200000000	sp P30510	1C14_HUMAN
8669	SSTLSSSSMSLSPK2000	sp P04222	1C03_HUMAN
8670	ISICSSDKR200000	sp Q07000	1C15_HUMAN
8671	HTSCSSAGNSKPVQEAPSVAR2.002E+20	sp Q29865	1C18_HUMAN
8672	AKSPTSPSPPRNSDQEGGK2.02020200002E+19	sp Q29960	1C16_HUMAN
8673	GVGSGPHPPDTQQPSPSK20000	sp Q29963	1C06_HUMAN
8674	SSSTEPPPPVVR2000000000	sp P30504	1C04_HUMAN
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8676	NFSSEIEDLEK2200000000	sp Q95604	1C17_HUMAN
8677	EVEEFSDEDDDDSDSSEAQKQSK2.00000002002E+20	sp Q9TNN7	1C05_HUMAN
8678	TDSV IADQTPPTR2000000000020000	sp P30505	1C08_HUMAN
		sp Q8N1G0	ZN687_HUMAN
		sp Q4LE39	ARI4B_HUMAN
		sp Q14157	UBP2L_HUMAN
		sp O15119	TBX3_HUMAN
		sp Q13547	HDAC1_HUMAN
		sp Q9P246	STIM2_HUMAN
		sp Q8IYB3	SRRM1_HUMAN
		sp Q68CZ2	TENS3_HUMAN
		sp Q8N684	CPSF7_HUMAN
		sp Q5T200	ZC3HD_HUMAN
		sp Q7Z3K6	MIER3_HUMAN
		sp Q9Y2W2	WBP11_HUMAN
		sp P17544	ATF7_HUMAN

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		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
		sp O14715 RGPD8_HUMAN
8680	YSLSPSK20000	sp Q99666 RGPD5_HUMAN
		sp A6NKT7 RGPD3_HUMAN
		sp P0DJ00 RGPD1_HUMAN
		sp P0DJ01 RGPD2_HUMAN
8681	SSTDLSLPGPISR20000000000	sp Q9HCD5 NCOA5_HUMAN
8682	HVTLPSSPR2000	sp Q5T4S7 UBR4_HUMAN
8683	AQSPVSGPNVAHLTDGATLNDR2E+20	sp Q5THJ4 VP13D_HUMAN
8684	LSLQHTQQNADGQEDGESERNSSSGDPGR2022000000	sp O75052 CAPON_HUMAN
8685	SQEEPKDTFEHDPSESIDFNK2E+22	sp Q9NYF8 BCLF1_HUMAN
8686	AALAPAKESPRK20000	sp Q13428 TCOF_HUMAN
8687	ANNSPALTGNSQPQHAAAAAAQQQQCGGGGATKPAVSGK2E+38	sp Q5VTL8 PR38B_HUMAN
8688	SLDSESI SPGDFHR200200020000000	sp Q52LW3 RHG29_HUMAN
8689	LPPPPPQAPPEEENESEPEEPSGVEGAAFQSR200000000000	sp O60341 KDM1A_HUMAN
8690	EDFDTIESKDEK2200200000000	sp Q9NRL2 BAZ1A_HUMAN
8691	KASSDSEDSSEEEEEVQGPAPK2. 22020022E+22	sp Q14978 NOLC1_HUMAN
8692	RDSLGTYSR200000000	sp Q9C0C2 TB182_HUMAN
8693	LMKPPSPEHQSLDTQQEGGEEVGPVAQQAIAVAEPSTSECLR2E+30	sp Q9Y6X9 MORC2_HUMAN
8694	SSSPLPTVQLHPQSPTAGK2000000000002000000	sp O60245 PCDH7_HUMAN
8695	TPQSPAPCVLLR2002000000000	sp Q9Y613 FHOD1_HUMAN
8696	DTYSDRSGSSPDSEITELK3000202200000000000	sp P17812 PYRG1_HUMAN
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8699	YVASYLLAALGGNSSPSAK200000	sp P05387 RLA2_HUMAN
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8704	GSISSTSEVHSPNVGLR2000020000000000000	sp Q8WWQ0 PHIP_HUMAN
8705	SSFHYSGLK2000000000	sp Q8NC51 PAIRB_HUMAN
8706	KPEKPLFSSASPQDSSPR20000000000	sp O00712 NF1B_HUMAN
8707	RNSSDSAIDNPKPNKLPK2002000000000000000	sp P46100 ATRX_HUMAN
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8712	QFTSSSIK20000	sp Q04695 K1C17_HUMAN
8713	SRTSPVTR202000000	sp Q9UQ35 SRRM2_HUMAN
8714	GYSSPEPDIQDSSGSEAQSVKPTR2. 00000002202E+22	sp Q9BZF1 OSBL8_HUMAN
8715	SSSPTSSLTQPIEMPTLSSPTEERPTVGGQDNPLK2. 02000000001E+37	sp Q7LBC6 KDM3B_HUMAN
8716	TVSASSTGDLPK22000000	sp P78362 SRPK2_HUMAN
8717	GRLDSSEMDHSENYDTMSSPLPGK2. 010020000001E+20	sp Q9NTI5 PDS5B_HUMAN
8718	SWVIGSPEILR2000000	sp Q6P0Q8 MAST2_HUMAN
8719	VTPHPSPR20002000	sp Q8IWQ3 BRSK2_HUMAN
8720	LHQASASSTSSLSTR200000	sp O15085 ARHGB_HUMAN
8721	SEDESETEDEEEKSQEDQEQK2000000000000000000	sp P25205 MCM3_HUMAN
8722	RPPEEDFRHSPEEDFRQSPQEHFR2000000020000000	sp Q8IXT5 RB12B_HUMAN
8723	RSLSGSSPCPK20200000000	sp Q9UQ35 SRRM2_HUMAN
8724	LLKPGEEPSEYTDDEEDTK20020000000	sp O15173 PGRC2_HUMAN
8725	LLASPSSR200000	sp Q15003 CND2_HUMAN

8726	SFGTRPLSSGFSPEEAQQQDEEFEK200000000000000000	sp Q9C0C2 TB182_HUMAN
8727	SKPNLPSESR20000	sp Q08170 SRSF4_HUMAN
8728	EVEEDSEDEEMSEDEEDDSSGEEVVIPQKK2.00001200000002E+25	sp P19338 NUCL_HUMAN
8729	SSRSSPELTR22002000000	sp Q9UQ35 SRRM2_HUMAN
8730	TSPVVAPTSEPSSPLHTQLLK22000000000	sp 075044 SRGP2_HUMAN
8731	ATTPPNQGRPDSPVYANLQELK2.0000000002E+21	sp Q8IWW6 RHG12_HUMAN
8732	LNASPAAR200000	sp Q9UGU0 TCF20_HUMAN
8733	TSPAPWK2000000	sp Q9UQ35 SRRM2_HUMAN
8734	DGQVINETSQHDDLE200000000	sp P08670 VIME_HUMAN
8735	SSTVTEAPIAVVTSR2200000000000000	sp Q8TD19 NEK9_HUMAN
8736	LSSSSPGATVSPSSSDAEYDKLPDADLAR2E+27	sp Q8NI08 NCOA7_HUMAN
		sp P30510 1C14_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
8737	GGSCSQAASSNSAQGSDESLIACK2000000000	sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
8738	SVSSPTSSNTPPTK200000000000000000	sp Q5M775 CYTSB_HUMAN
8739	ELSTSPCNPIVR2200000000	sp Q9H2Y7 ZN106_HUMAN
8740	GQSQTSPDHRSDTSSPEVR202000000002200000	sp Q9UQ35 SRRM2_HUMAN
8741	QEEEEAAQGPVVVSPASDYK20000000	sp 075391 SPAG7_HUMAN
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8743	SPAKSPNPSTPR2000000022000	sp 076021 RL1D1_HUMAN
8744	RLSESSALK20000000	sp Q96S55 WRIP1_HUMAN
8745	TDALTSSPGR20000220000	sp P49736 MCM2_HUMAN
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8747	GLLPGGTQVLDGTSGFSPAPK200000	sp Q96S94 CCNL2_HUMAN
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8751	LASDDRPSPPR2000000000	sp P98175 RBM10_HUMAN
8752	SRSPPPVSK2020000000	sp Q01130 SRSF2_HUMAN
8753	RFSRSPIR2020000	sp P18583 SON_HUMAN
8754	KLSEADNR2000000	sp Q9UNL2 SSRG_HUMAN
8755	LSSSSPGATVSPSSSDAEYDKLPDADLAR2.00000003E+19	sp Q8NI08 NCOA7_HUMAN
8756	SREDSPELNPPPIEDNR20002000000000000000	sp Q03164 KMT2A_HUMAN
8757	ALEAASLSQHPPSLCISDSEEEEEER2000200000000000	sp 043159 RRP8_HUMAN
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8759	TESGEETDLISPPQEEVK20000000000000000000	sp 043493 TGON2_HUMAN
8760	SMVSPVPSPTGTISVPNSCPASPR2.102E+24	sp P85037 FOXK1_HUMAN
8761	LVEDERSDREETESSEGEAAAAGGAK2.00002002E+21	sp Q96G23 CERS2_HUMAN
8762	TSPLKDNPSPEPQLDDIKR200000000000	sp Q96JC9 EAF1_HUMAN
8763	GSVILDSGHLSTASSDDLKGEESFR2200000000000000	sp Q92614 MY18A_HUMAN
8764	SGSEDEDDGETLLEVAGTQGK2.002000002E+20	sp Q9P1Y6 PHRF1_HUMAN
8765	SGSSPDSEITELKFPSINH20000000000000000000	sp P17812 PYRG1_HUMAN
8766	MVSQSQPGSR200	sp 075122 CLAP2_HUMAN
8767	SLGEKSPAASGAR2000000000	sp 075683 SURF6_HUMAN
8768	EEGSLSDTEADAVSGQLPDPTTNPAGK2000000000000000	sp Q8N1G4 LRC47_HUMAN
8769	RKPSTSDSDSNFEK22000200000000	sp P11388 TOP2A_HUMAN

8770	GLAPNTPGK20000	sp 043663 PRC1_HUMAN
8771	DLSTSPKPSPIPSVLGR200000002000000	sp Q8NDI1 EHBP1_HUMAN
8772	RVPSPTAPK20000000	sp Q9UQ35 SRRM2_HUMAN
8773	SRSYTPEYR2023000000	sp Q13595 TRA2A_HUMAN
8774	LLKPGEPESEYTDDEEDTKDHNKQD20000000000000000	sp O15173 PGRC2_HUMAN
8775	LPSSPASPSPKGTPEK200020000	sp Q9H4G0 E41L1_HUMAN
8776	ALEQQEAESDSSDTEEKDDDDDEEDVGK2.02202E+21	sp Q9BXY0 MAK16_HUMAN
8777	ERSKDASPINR2000200000	sp Q13523 PRP4B_HUMAN
8778	RPSSTSVPLGDK2000000000	sp Q9NXR1 NDE1_HUMAN
8779	ISSPMHPSAGGQR200000000000	sp O00267 SPT5H_HUMAN
8780	VASPSQGQVGS SSPK2000	sp Q99569 PKP4_HUMAN
8781	DTIIDVVGAPLTPNSR200000	sp Q9NPF5 DMAP1_HUMAN
8782	KPVNYSQFDHSDSDDDFVSATVPLNK2.0000002E+21	sp Q96B01 R51A1_HUMAN
8783	LKTSLGK20000	sp P46013 KI67_HUMAN
8784	EESDDEDKESEEEPPK2200000020000000	sp P35659 DEK_HUMAN
8785	SSSTEPPPPVR200000000	sp Q8N684 CPSF7_HUMAN
8786	SPPSQKSPHDSKSPLDHR2000002000002000000	sp O14647 CHD2_HUMAN
8787	SSGHSSSELSPDAVEK22002020020000000	sp Q9UQ35 SRRM2_HUMAN
8788	RSSPPPPSGSSSR2000000000000	sp Q5VUA4 ZN318_HUMAN
8789	TVCGSPLAAQSPSAELPQAAPPPGSPSLPGR2E+29	sp Q969V6 MKL1_HUMAN
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		sp Q99880 H2B1L_HUMAN
		sp P33778 H2B1B_HUMAN
		sp Q99877 H2B1N_HUMAN
		sp Q8N257 H2B3B_HUMAN
		sp Q96A08 H2B1A_HUMAN
		sp Q5QNW6 H2B2F_HUMAN
		sp P06899 H2B1J_HUMAN
8791	QVHPDTGISSK2000	sp Q16778 H2B2E_HUMAN
		sp P23527 H2B1O_HUMAN
		sp P57053 H2BFS_HUMAN
		sp P58876 H2B1D_HUMAN
		sp O60814 H2B1K_HUMAN
		sp P62807 H2B1C_HUMAN
		sp Q93079 H2B1H_HUMAN
		sp Q99879 H2B1M_HUMAN
8792	RRDEDMLYSPELAQR20000000	sp Q9Y2W2 WBP11_HUMAN
8793	VSEEQTQPPSPAGAGMSTAMGR20000010000000	sp Q16666 IF16_HUMAN
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		sp P30447 1A23_HUMAN
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8798	SRNSPLLER2000000	sp Q7KZI7 MARK2_HUMAN
8799	VSGAQEMVSSAK200001000000	sp O60664 PLIN3_HUMAN
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8801	DASDDLDDLNFFNQK200000000000000	sp P20042 IF2B_HUMAN
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8803	KISPPSYAR20000000	sp Q8N2M8 CLASR_HUMAN
8804	ALEEGDGSVSGSSPR20000	sp Q8N3X1 FNBP4_HUMAN



		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
8805	EDALDDSVSSSSVHASPLASSPVRK200000	sp O14715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
		sp A6NKT7 RGPD3_HUMAN
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8808	ERPSGPESPVQGPR20000000	sp Q9UMN6 KMT2B_HUMAN
8809	SPQPDPVKTPTSSK200000002002000	sp P46013 KI67_HUMAN
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8811	SSGVVPQSAPPVPTASSR200000000000	sp P41162 ETV3_HUMAN
8812	TPTPSTPGDTQNPNTAPVPPAEDGIK2.00000002E+21	sp Q14839 CHD4_HUMAN
8813	SSSFSELCHR2000000000	sp Q5T8I3 F102B_HUMAN
8814	RRSFSISPVR202020000	sp P18583 SON_HUMAN
8815	AAESETPGKSPEKKPK200000000000	sp O76021 RL1D1_HUMAN
8816	SGLDLNTSLSPSLQPGK2000000000	sp Q99708 COM1_HUMAN
8817	SPQRSPLQSAESSPTAGK2000200000000020000	sp Q8IWB9 TEX2_HUMAN
8818	SPSPAVPLR2000000000	sp Q6P4R8 NFRKB_HUMAN
8819	STSAPQMSPGSSDNQSSSPQPAQQK1.2002E+19	sp Q14157 UBP2L_HUMAN
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8821	QLVYVEQAGSSPK20000	sp Q9C0H5 RHG39_HUMAN
8822	GGHSSVSTESESSSFHSS2020000000000000	sp P61073 CXCR4_HUMAN
8823	LDSSPVSSPR20020000	sp O60293 ZC3H1_HUMAN
8824	SQSKSPTGTPAR20200020000	sp Q13595 TRA2A_HUMAN
8825	VTQGAASPGHGIQEK2000000000	sp Q9HB58 SP110_HUMAN
8826	SSSVLSLEGSEK200020002000	sp A1L390 PKHG3_HUMAN
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8829	IPRPSVSGCSR2000200	sp O75122 CLAP2_HUMAN
8830	SLPSSQLKGSPPQAIISR20000000	sp Q8WYP5 ELYS_HUMAN
8831	MLPHAPGVQMQAIPEDAVHEDSGDEDGEDPDKR1.00000000002E+24	sp Q92769 HDAC2_HUMAN
8832	AGPVSPGCVK2000000	sp Q53GL7 PAR10_HUMAN
8833	AQQELVPPQQQASPPQLPK20000000	sp O15027 SC16A_HUMAN
8834	TAVPPLSEGDGYSSENTSRSR22000000	sp Q9BYW2 SETD2_HUMAN
8835	RESESEDETPPAAPQLIK200020020000000000	sp O60832 DKC1_HUMAN
8836	SHMSGSPGPGSNTAPSTPVIIGSDKPGMEEK2.02E+29	sp A0FGR8 ESYT2_HUMAN
8837	DDGDEEGLLTHSEEELEHSQDTDADDGALQ2.000000002E+21	sp Q9UKM9 RALY_HUMAN
8838	QKSDAEEDGGTVSQEEEDRKPK2.0000000202E+20	sp P27824 CALX_HUMAN
8839	HNSPTLSCR20000000	sp Q5T5U3 RHG21_HUMAN
8840	LPPQSSGVDTSPCPNPFR200000	sp O75815 BCAR3_HUMAN
8841	EAASSPAGEPLR2200000000	sp Q96EB6 SIR1_HUMAN
8842	SSSLGSTPHEELER22000000000000	sp Q9NRA8 4ET_HUMAN
8843	ALFKPPEDSQDDESDDAEEEQTTKR2000020000000000000	sp Q13769 THOC5_HUMAN
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8845	IKDTCIQSPSK20000	sp Q03188 CENPC_HUMAN
8846	TLSSSSMDLSR20000000	sp Q9H0B6 KLC2_HUMAN
8847	DSHSSEDEASSQTDLSQTISK202000000000	sp Q5JTV8 TOIP1_HUMAN
8848	HGGSPQPLATTPLSQEPVNPPEASPTR2.200200000002E+19	sp Q9UQ35 SRRM2_HUMAN
8849	GGNLPPVSPNDGAK2000000000	sp Q9UK61 F208A_HUMAN
8850	SPNSHDSEPWTLLR200000000	sp Q5T5Y3 CAMP1_HUMAN
8851	SPPSVQSLAMR200000000000	sp Q96KQ7 EHMT2_HUMAN
8852	SRSGEGEVSGLMR20000000000000	sp Q13263 TIF1B_HUMAN
8853	IYSFTDNAPSPSIGSSR2000000000	sp O15014 ZN609_HUMAN
8854	QGSPVSPNALDR20000000	sp Q5VWQ8 DAB2P_HUMAN

8855	SWDSSSPVDRPEPEAASPTTR220000000000200000	sp Q86WB0 NIPA_HUMAN
8856	ASSLESGVPSR200	sp P01598 KV106_HUMAN
8857	ASTTSLSGDSETEGK20200000000	sp Q8N1G2 CMTR1_HUMAN
8858	VRRPIPGSDQTTTMSGGEGEPGEGEGDGEDVSSDSAPDSAPGPAPK1E+33	sp Q06587 RING1_HUMAN
8859	TPLALAGSPTPK2000000200000	sp 095785 WIZ_HUMAN
8860	SQLSFLPTEAQAQVSPPPYPAPQELTQPLLQQPR2E+32	sp Q6UUU7 CRTC3_HUMAN
8861	GQTPLTEGSELDGHSDPEESFAR2000000000	sp Q86VR2 F134C_HUMAN
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		sp POCL81 GG12G_HUMAN
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		sp 076087 GAGE7_HUMAN
8863	SPMCNQPSINK200000000000	sp Q08499 PDE4D_HUMAN
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8869	SRSSSVGSSSYPIPAVSR2220000000000	sp Q15149 PLEC_HUMAN
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8871	GIISDGESN22000200	sp Q9BW71 HIRP3_HUMAN
8872	SFSISPVR200020000	sp P18583 SON_HUMAN
8873	DSLSPVLHPSDLILTR2000000000000000	sp Q9HCN4 GPN1_HUMAN
8874	GSLAEAVGSPPPAATPTPTPTR2.0000002E+22	sp Q9Y6I3 EPN1_HUMAN
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8877	GHPAQRPEPSPPEPEQAPEGSQAGAEGSSPEASRSPAR20000020000	sp Q6ZW31 SYDE1_HUMAN
8878	FIQELSGSSPK20000	sp Q01664 TFAP4_HUMAN
8879	AFLAELEQNSPK2000	sp Q9UPN3 MACF1_HUMAN
8880	EIQNGNLHESDSESVPR2000000	sp Q86UP2 KTN1_HUMAN
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		sp P49792 RBP2_HUMAN
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		sp A6NKT7 RGPD3_HUMAN
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		sp PODJD0 RGPD1_HUMAN
		sp PODJD1 RGPD2_HUMAN
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8891	KGSVSHDTVQPR200000	sp Q96D71 REPS1_HUMAN
8892	IYHPSCYEDYQNTSSFDCTPSPSK2020000	sp 094913 PCF11_HUMAN
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8895	KKSSSEAKPTSLGLAGGHK202000002000000000	sp Q96RK0 CIC_HUMAN
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8897	LSSLRASTSK2200000000	sp P62753 RS6_HUMAN
8898	YQPLASTASDNDFVTPEPR200000200000	sp Q15021 CND1_HUMAN
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8901	AGPDLPSPSLSHALEDEGWAAAAPSPGSAR2000000	sp Q9H3Q1 BORG4_HUMAN
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8906	IHLPLNYPPGSPDLGR2000000	sp Q08174 PCDH1_HUMAN
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8919	SFSISPSR202000000	sp Q92794 KAT6A_HUMAN
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8922	AELGMDSTSQSPPIK1000000200000	sp P35659 DEK_HUMAN
8923	QLSSSAAPAAQPPGSR20000000000000000	sp Q9NR19 ACSA_HUMAN
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		sp Q09160 1A80_HUMAN
		sp P05534 1A24_HUMAN
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		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
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8928	LLPYPTLASPASD200	sp P0C1Z6 TFPT_HUMAN
8929	ETPAATEAPSSPK200000	sp P80723 BASP1_HUMAN
8930	SPQRPSDWSK20000000000	sp Q5UIP0 RIF1_HUMAN
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8947	FGSTGSTPPVSPTPSER200020200000	sp Q969V6 MKL1_HUMAN
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8956	LSSGPEQITLEASSTEGHPGAPSPQHTDQTEAFQK2E+33	sp Q6P996 PDXD1_HUMAN
8957	SLDGAEF SRPASVSENHDAGPDGDKR2E+26	sp Q9H4G0 E41L1_HUMAN
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		sp Q7Z3J3 RGPD4_HUMAN
		sp A6NKT7 RGPD3_HUMAN
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		sp Q99666 RGPD5_HUMAN
		sp 014715 RGPD8_HUMAN
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8980	DLESTSRDSLALDK22000000000	sp Q5T200 ZC3HD_HUMAN

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		sp O14715 RGPD8_HUMAN
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		sp PODJD0 RGPD1_HUMAN
		sp PODJD1 RGPD2_HUMAN
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8983	SKDVNGPEPLNSR200	sp Q99653 CHP1_HUMAN
8984	QASASYDSEEEEEGLPMSYDEK2000000000000000	sp Q15059 BRD3_HUMAN
8985	SFSISPSR2020000	sp P18583 SON_HUMAN
8986	IDAGTMAEPSASPSKR200000	sp Q8IXQ3 CI040_HUMAN
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8992	KSPSGPVKSPPLSPVGTTPVK20000000200000	sp Q9BVC5 ASHWN_HUMAN
8993	VEPSSQSPGRSPR2000	sp Q8IUD2 RB612_HUMAN
8994	ADRDQSPFSK200000	sp Q13427 PPIG_HUMAN
8995	KKSSSEAKPTSLGLAGGHK2000000020000000000	sp Q96RK0 CIC_HUMAN
8996	RSPSVSPEPAEK20202200000000	sp Q9UQ35 SRRM2_HUMAN
8997	DHSPTPSVFNSEER200020000000000	sp Q6UN15 FIP1_HUMAN
8998	RSSPSARPPDVPGQPQAAK2. 2E+19	sp Q96JP5 ZFP91_HUMAN
8999	QSPSPSTRPIR20200000000	sp Q8IYB3 SRRM1_HUMAN
9000	SYQNSPSSDDGIRPLPEYSTEK2. 000202E+22	sp Q15648 MED1_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
9001	EDALDDSVSSSVHASPLASSPVR2000200000	sp A6NKT7 RGPD3_HUMAN
		sp Q99666 RGPD5_HUMAN
		sp O14715 RGPD8_HUMAN
9002	LSGSDLGGHSSLER20000000000000	sp O95613 PCNT_HUMAN
9003	QPSWDPSVPSSTVPAPSPLSAAAVSPSK2000020000000000000	sp Q9NQC3 RTN4_HUMAN
9004	LQEEGGGSEEEETGSPSEDMQSAR20000000020000000000	sp Q13610 PWP1_HUMAN
9005	DNSNSDSEGEDRRPAGPPR20202000000000000000	sp Q8NCF5 NF2IP_HUMAN
9006	TPTPSTPGDTQNPNTAPVPPAEDGIK2. 00020002E+21	sp Q14839 CHD4_HUMAN
9007	TRTSSSETEEEKKPEKPLILQSLGPK2. 2E+22	sp Q9Y5K6 CD2AP_HUMAN
9008	DTCYSPKPSVYLSTPSSASK2000000000000000000	sp Q9Y5K6 CD2AP_HUMAN
9009	NYQQNYQNSESGEKNEGSESAPEGQAQQR3. 00202E+24	sp P67809 YBOX1_HUMAN
9010	SSPGSVAASPSGGGR20000000000000	sp Q9BXB5 OSB10_HUMAN
9011	AAFTTPDHAPLSPQSSVASSGSEQTEEQGSSR2E+21	sp Q5PRF9 SMAG2_HUMAN
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9013	QDDSPRP I I GPALPPGFIK20000000000000000000	sp Q8IXQ4 GPAM1_HUMAN
9014	VLEDEEGSKDIELSDDPYDCIR20000000000000000000	sp Q15024 EXOS7_HUMAN
9015	GSPHYFSPFRPY200000000000000	sp Q13242 SRSF9_HUMAN
9016	SSTPSSPTGTSSSDSGHHIGWGER2. 2E+24	sp P46019 KPB2_HUMAN
9017	TSISDPPEPPLLPPR2000000000	sp Q07889 SOS1_HUMAN
9018	LPDLSPVENK2000000	sp Q9Y520 PRC2C_HUMAN
9019	SAASPVVSSMPERASESSSEEKDDYEIFVK2. 00000000100002E+30	sp P18583 SON_HUMAN
9020	GSVILDSGHLSTASSDDLKGEESFR22022000000000000000	sp Q92614 MY18A_HUMAN
9021	HNSWSSSR20000000	sp Q96F86 EDC3_HUMAN
9022	LRSFASQSTDR20200000000	sp Q9ULJ3 ZBT21_HUMAN
9023	NSNFSVQHPSSTSPTEK20000000	sp Q03468 ERCC6_HUMAN
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9025	SASVSSISLTK2002000000	sp Q07889 SOS1_HUMAN
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9028	SPSFGGASEASAR200000000000	sp Q8WVT3 TPC12_HUMAN
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9030	SPTPKSPPSR200200000	sp Q9NVD7 PARVA_HUMAN
9031	DGQSSPQPSPR20002000	sp Q5VZK9 LR16A_HUMAN
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9033	LQSGVSLASK200200	sp Q09666 AHNK_HUMAN
9034	HQASDSENEEPPKPR2000000000000000	sp Q96ST2 IWS1_HUMAN
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9037	RKTLDAEVVEKPAK2000000000000000	sp Q9H3P2 NELFA_HUMAN
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9048	TPHPSPAHSSR2000200000000	sp Q6ZS17 FA65A_HUMAN
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9050	EVDGLLTSEPMGSPVSSK102000000	sp Q8WWM7 ATX2L_HUMAN
9051	YSPPIQR2000000	sp Q8IYB3 SRRM1_HUMAN
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9054	RQSSDSCSEEPDSTTPPAK22020200000000000000	sp Q8N1G0 ZN687_HUMAN
9055	QSVTSFPDADAFHHQVHDDLLSSSEEECK222000000	sp P50402 EMD_HUMAN
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9057	SNVSSPATPTASSSSTPTR20020000000000000000	sp Q12888 TP53B_HUMAN
9058	DRQSPLHGNHITISHTQATGSR2E+19	sp Q86VQ1 GLC11_HUMAN
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9067	TSSKESPISPTSDRK20022000000000000000	sp Q01082 SPTB2_HUMAN
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9069	LSGEHSESSTPRPR2000200000	sp Q7Z3K3 POGZ_HUMAN
9070	SLKESEQESEEEILAQK20000000200000000000	sp Q9BW71 HIRP3_HUMAN
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9072	LFSADPFLEAQAK2000000000000000	sp Q5TDH0 DDI2_HUMAN
9073	SLTRSPPAIR20202000000	sp Q9UQ35 SRRM2_HUMAN
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9076	KISENSYSLDDLEIGPGQLSSSTFDSEK2E+23	sp Q9UHB6 LIMA1_HUMAN
9077	HTDDEMTGYVATR20000000	sp Q16539 MK14_HUMAN
9078	TPSSPPIPPASET22000000000000000000	sp Q9NZJ0 DTL_HUMAN

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9081	KPVTVSPTTPTSPTGEAS200000000000	sp Q9Y6G9 DC1L1_HUMAN
9082	RPNKQEESESPVERPLKENPPNK2020000000000000	sp Q96SB4 SRPK1_HUMAN
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9084	TESVPSDINNVPVDR20000000000000	sp Q8IVT5 KSR1_HUMAN
9085	SRESSPSHGLLK2200000000	sp Q96JH7 VCIP1_HUMAN
9086	KKTVSFSSMPTEK202000000000	sp Q9UPRO PLCL2_HUMAN
9087	TASRPDDIPDPSPPK200000000020000	sp Q5VZK9 LR16A_HUMAN
9088	SPQVLGSSLSVR20000000000000	sp Q86UU0 BCL9L_HUMAN
9089	NEGSSIEDAQTSQHATLKPER200000000000	sp Q9UKL3 C8AP2_HUMAN
9090	ARNSGTQSDGEEKQPLVMER20002000000000000000	sp Q6P1M3 L2GL2_HUMAN
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9092	TPLSSLSATSNNSTVACIDR2000000000000000	sp P00533 EGFR_HUMAN
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9096	HHQEEDAGPTQPSAKPQLK2000000000	sp Q9C086 IN80B_HUMAN
9097	SSSSSGVPYSPAIPNK2000000000000000	sp Q9UPT5 EXOC7_HUMAN
9098	MSPGVAGSPR2000000000	sp Q15596 NCOA2_HUMAN
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9100	SPVDSGTILR200000000000	sp P49454 CENPF_HUMAN
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9106	THSTSSSLGSGESPFGR2200000000000000	sp Q9UGV2 NDRG3_HUMAN
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9114	LSVPTSDEEDEVPAKPR20000000000000000000	sp Q8NE71 ABCF1_HUMAN
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9126	SLSPVPR20200000	sp A2AJT9 CX023_HUMAN
9127	SQDSYPGSPSLSPR2000000000002000	sp Q6VN20 RBP10_HUMAN
9128	LPSGSPSPK2000	sp Q9P275 UBP36_HUMAN
9129	SPEPDTLPLAAGSDHPLPR2000000000000000	sp Q6PCT2 FXL19_HUMAN
9130	RNSNSPPSPSSMNQR20200000200000	sp Q7Z5L9 I2BP2_HUMAN
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9133	VSKSPSPASTSTVPNMTDAPTAPK22000000000000	sp Q659A1 ICE2_HUMAN
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9137	SDDSKSSPELVTHLK20002000000000	sp Q07960 RHG01_HUMAN
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9146	YSDGDKSDDLVDVSNEDPATPR3E+24	sp Q9UHR4 BI2L1_HUMAN
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9160	SPGSPVGEETGSPPK2002000000000000	sp Q92576 PHF3_HUMAN
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9167	AQEPPAGGGSIHDLPLFTETSPTSAPWDPLPGPPVLPSPHSHL2E+22	sp Q13595 TRA2A_HUMAN
9168	TGSLSDYSR20000000	sp P62995 TRA2B_HUMAN
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		sp Q9NUQ3 TXLNG_HUMAN



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9212	GPTSSPCEEEGDEGEEDRTSDLR2.2E+20	sp Q86VM9 ZCH18_HUMAN
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9225	QNGSNDSDRYSDNEEDSKIELK2. 000003E+19	sp	P42167	LAP2B_HUMAN
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9229	ISSVPNSLSK200000000	sp	P10636	TAU_HUMAN
9230	SPTLSQVHSPLVTSPSANLK2E+20	sp	Q5VZ89	DEN4C_HUMAN
9231	SVSASSGPCSPAAPHPPPPR2. 002E+19	sp	Q86UU0	BCL9L_HUMAN
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9239	EASAESDVGKSESEFTQYTTTHILK2. 00200002E+24	sp	Q15751	HERC1_HUMAN
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9242	IACKSPPPEMDTPTSTR2000000000000000	sp	P24928	RPB1_HUMAN
9243	KPVTVSPPTPTSPTEGEAS202200000000000	sp	P46013	KI67_HUMAN
9244	EGSTSSLEPEPDTEK222000000000000	sp	Q9Y6G9	DC1L1_HUMAN
9245	LKSSTSFANIQENSN202000000000000	sp	Q9COH5	RHG39_HUMAN
9246	SSGDPEQIKEDSLSEESADAR20000200000	sp	Q86WC4	OSTM1_HUMAN
9247	AGEPNSPDAEANS PDV TAGCDPAGVHPPR2. 000000000002E+25	sp	Q9BVS4	RIOK2_HUMAN
9248	STPPSPTHR2002000000	sp	Q08J23	NSUN2_HUMAN
9249	GSVPLSSPPATHFPDETEITNPVK2. 0002E+19	sp	A8CG34	P121C_HUMAN
9250	SYEDMIGEEVPSDQYYWAPLAQHER2E+25	sp	P11388	TOP2A_HUMAN
9251	DAEPQPGSPA AESLEEPDAAAGLSSTK200000000000000	sp	060716	CTND1_HUMAN
9252	LNSVQSSERPLFLVHPIEGSTTVFHSLSR2E+24	sp	Q9H9J4	UBP42_HUMAN
9253	RPSASSPNNNTAAK2000000000	sp	P49327	FAS_HUMAN
9254	MGQAPSQSLPPAQDQPRSPVPSAFSDQSR2E+23	sp	075151	PHF2_HUMAN
9255	RSSLSPGR2020000	sp	Q9UQ35	SRRM2_HUMAN
9256	RNSDRSEDGYHSDGDYGEHDYR200030200000000000	sp	Q9BUQ8	DDX23_HUMAN
9257	EGEDGDQPTTPKPLK200000000	sp	P52756	RBM5_HUMAN
9258	HSDTSPRR200200000	sp	P18858	DNL11_HUMAN
9259	HFEASCGQLSPAR2000000000	sp	Q9BRD0	BUD13_HUMAN
9260	FDSSLLSSDDETK2000000	sp	Q96N21	AP4AT_HUMAN
9261	TLSSSSMDLSR200000000	sp	P28715	ERCC5_HUMAN
9262	AEGIEGETLTASPQAPGSPEDSEGVLISLPR2. 000002E+21	sp	Q9H0B6	KLC2_HUMAN
9263	SPSEESAPTTSPESVSGVSPSSGSSGR200200000000000000	sp	Q5T011	SZT2_HUMAN
9264	LNFDMTASPK2000	sp	P54725	RD23A_HUMAN
9265	TSASDVTNIYPGDAGK200000000000000000	sp	Q9ULU4	PKCB1_HUMAN
9266	HCGLSLSSTPPGK2002000000	sp	Q15042	RB3GP_HUMAN
9267	TEAPGTPEGPEPERPSPGDGNPR200000000000000000	sp	Q71F23	CENPU_HUMAN
9268	SEESD TDTSERQDDSYIEPEVPLK2. 00002022000002E+26	sp	Q9Y666	S12A7_HUMAN
9269	SQTTTERDSDTDVEEELPVENR2. 20002E+20	sp	Q9BZF1	OSBL8_HUMAN
9270	NEMGIPQTTSPENAGPQNTK200000000000	sp	Q14676	MDC1_HUMAN
9271	SYSPDGKESPSDKK202000002000000	sp	Q15398	DLGP5_HUMAN
9272	HASPILPITEFS DIPR2000000000000000	sp	P43243	MATR3_HUMAN
		sp	P42167	LAP2B_HUMAN

9273	QQSTSSDRVSTPESLDFLK220000000000000000	sp	Q12965	MYO1E_HUMAN
9274	SPVGSAGPQAAAPAPAAHVAGNPGGDAAPAATGTAASLATAAGSEDAEK2E+51	sp	P16989	YBOX3_HUMAN
9275	GGSEGGPESSTASNSVPPPPQGGSGR2E+24	sp	Q9P206	K1522_HUMAN
9276	GGAAGGALPTSPGPALGAK2000000000	sp	Q9UPR0	PLCL2_HUMAN
9277	QSVTSFPDADAFHHQVHDDLLSSSEEECK2E+29	sp	P50402	EMD_HUMAN
9278	SQDATFSPGSEQAEKSPGPIVSR200200000000000000	sp	Q86WB0	NIPA_HUMAN
9279	KSSTSDWPTDNQDGDGK2000000000000000	sp	Q92794	KAT6A_HUMAN
9280	RHSSDINHVLVTQGRESPEGSYTDANQEV2. 00002E+20	sp	Q5TON5	FBP1L_HUMAN
		sp	P49792	RBP2_HUMAN
		sp	Q7Z3J3	RGPD4_HUMAN
9281	LNQSGTSSVGTDEESDVTQEEER20020002000000000	sp	A6NKT7	RGPD3_HUMAN
		sp	P0DJD0	RGPD1_HUMAN
		sp	P0DJD1	RGPD2_HUMAN
9282	EAKPGAAPEVGVPSLSPSSPSSWTETDVEER2000000000000000	sp	Q6L8Q7	PDE12_HUMAN
9283	SKSPSPR202000000	sp	Q9UKV3	ACINU_HUMAN
9284	DGSDYDPYDFSDTEEMPQVHTPK200000000002000	sp	P25205	MCM3_HUMAN
9285	KKTVSASESEDR20200000000	sp	Q9ULG6	CCPG1_HUMAN
9286	FSGFSAKPNSGEAPSSPTPK2000000	sp	P49916	DNLI3_HUMAN
9287	GAGATSGSPAGR200000000	sp	P16455	MGMT_HUMAN
9288	YSPSQNSPIHHIPSR2020000000000000	sp	Q9NYF8	BCLF1_HUMAN
9289	SPGSGSQSSGWHEVEPGMPSPTLKK2E+26	sp	Q12857	NF1A_HUMAN
9290	SAASPVVSSMPERASESSSEKDDYEIFVK2. 000002022E+22	sp	P18583	SON_HUMAN
9291	RGSLEMSDGEPLSR2000000000000000	sp	Q6ZN18	AEBP2_HUMAN
9292	LPSSPASPSK202000	sp	Q9H4G0	E41L1_HUMAN
9293	ARHDPDPSPLR200020000	sp	Q9BRD0	BUD13_HUMAN
9294	DEPPPLSPAPLTPATPSSLDPFPSR2. 00002E+19	sp	Q7Z6Z7	HUWE1_HUMAN
9295	KRSEGFMDR200001000	sp	Q7Z4V5	HDGR2_HUMAN
9296	SIQPQVSPR2000	sp	Q13322	GRB10_HUMAN
9297	ASAPERTPSPAPK20200000	sp	Q63ZY6	NSN5C_HUMAN
9298	GYSSPEPDIQDSSGSEAQSVKPSTR3. 020000002202E+24	sp	Q9BZF1	OSBL8_HUMAN
9299	RRSPSPYYSR200000000	sp	Q13595	TRA2A_HUMAN
		sp	P62995	TRA2B_HUMAN
9300	SPVSPLAAQGIPLPAQLTK2. 002E+19	sp	Q96SI1	KCD15_HUMAN
9301	MSDSESEELPKPQVSDSESEPPR2. 020000000002E+23	sp	Q96ST2	IWS1_HUMAN
9302	RPASPSPEHLPATPAESPAQR20000002000000000	sp	Q9H7L9	SDS3_HUMAN
9303	VEGSPTEANLPTQLDNSLSPAASEAGEK2000002020000000000	sp	P50851	LRBA_HUMAN
9304	TDSTSDGRPAWMR200000000100	sp	Q14204	DYHC1_HUMAN
9305	SPCETISSPSTLESK2000000000	sp	Q9C0D5	TANC1_HUMAN
9306	HTTENKPHSPKTPEPR200200000	sp	P54278	PMS2_HUMAN
		sp	Q68D20	PMS2L_HUMAN
9307	SYTSDLQK200000000	sp	Q8WWI1	LMO7_HUMAN
9308	FHSSDSEEEHKK20200000000	sp	Q96ST2	IWS1_HUMAN
9309	SPTSPSPPRNSDQEGGGK2. 02020200002E+19	sp	Q8IYB3	SRRM1_HUMAN
9310	ALEAASLSQHPPSLCISDSEEEEEER2. 00000000002E+19	sp	Q43159	RRP8_HUMAN
9311	IRHSSDTSPPR220200000	sp	Q9BRD0	BUD13_HUMAN

		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
		sp Q29960 1C16_HUMAN
9312	GGSCSQAASSNSAQGSDESLIACK20002002000000	sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
9313	EVEEDESEDEEMSEDEEDDSSGEEVVIPQK2.00000000000022E+24	sp P19338 NUCL_HUMAN
9314	MCESSPPESADTPTSTR20000000000000	sp P46013 KI67_HUMAN
9315	DVLLSSSSR200200	sp Q9Y2X9 ZN281_HUMAN
9316	TASQLENKPVK2002000000	sp P30414 NKTR_HUMAN
9317	AAATGNASPGKLEHSK2000000000	sp Q8NB78 KDM1B_HUMAN
9318	EWYQSTIPQSPSPAPDDPEEGR3.000000002E+20	sp Q08499 PDE4D_HUMAN
9319	ATSPSSSVSGDFDDGHHSVSTPGPSR2.022E+24	sp Q86U86 PB1_HUMAN
9320	GSPEPSALPPQR200000000000	sp 000562 PITM1_HUMAN
9321	SGAVQGAGSLGPGSPVR20000	sp Q9P270 SLAI2_HUMAN
9322	TEFLHSQNSLSR200000	sp Q96FS4 SIPA1_HUMAN
9323	QQSTSSDRVSQTPESLDFLK202000000000	sp Q12965 MYO1E_HUMAN
9324	SSGSPYGGGYGSGGGSGGYGSR30000000000000	sp P51991 ROA3_HUMAN
9325	SSPPASPTIIPR200020000000	sp P80192 M3K9_HUMAN
		sp P10636 TAU_HUMAN
9326	TPPKSPSSAK20002000000	sp TAU_HUMAN
9327	KRPEPSSDYDLSPAK2000020000	sp Q6W2J9 BCOR_HUMAN
9328	SPPPPPPPPR200000000000	sp Q5T5P2 SKT_HUMAN
9329	RRSSTVAPAQPDAESEWTDVETR2.2E+22	sp Q02241 KIF23_HUMAN
9330	HSQSPEPHSPTPEPPAWGSSIVK2.00000002E+22	sp Q5VZ89 DEN4C_HUMAN
9331	TDDEFVPSDASPPK2000000000020000	sp P11388 TOP2A_HUMAN
9332	LTNLSSNSDV2002000	sp Q13641 TPBG_HUMAN
9333	TAPSSPLTSPSDTR22000000	sp Q8IWB9 TEX2_HUMAN
9334	SLSLGDKEISR2000000000	sp Q9UMZ2 SYNRG_HUMAN
9335	ERFSPPR20000	sp Q9BXP5 SRRT_HUMAN
9336	FLMPEAYPSSPR2000	sp 043166 SIIL1_HUMAN
9337	SSLSSHSHQSQIYR200000000000	sp 015027 SC16A_HUMAN
9338	DAPTSPASVASSSSTPSSK2002000000000000	sp Q04726 TLE3_HUMAN
9339	SGPKPFSAPKPKQTSPSPKR20020000	sp Q01518 CAP1_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
9340	EDALDDSVSSSVHASPLASSPVRK200000	sp A6NKT7 RGPD3_HUMAN
		sp 014715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
9341	SPYKFPSSPLR200000200000	sp P06400 RB_HUMAN
9342	VTGTEGSSSTLVDYTSTSTGGSPVRK200000	sp Q7Z6E9 RBBP6_HUMAN
9343	GLWSTDAEEDKETK2000000000	sp Q99549 MPP8_HUMAN
9344	GRPSSPRTPLYLQPDAYGSLDR200200000000000000	sp Q8WUF5 IASPP_HUMAN
9345	RPGNSTVTSSPPVTPK20002000	sp Q7Z2Z1 TICRR_HUMAN
9346	SNSVSLVPPGSSHTLPR2000000000000000	sp Q6R327 RICTR_HUMAN
9347	EANPTPLTPGASSLSQLGAYLSDSDSNGSN200200200	sp Q9BW85 CCD94_HUMAN

		sp Q562R1 ACTBL_HUMAN
		sp Q6S8J3 POTEE_HUMAN
		sp A5A3E0 POTEF_HUMAN
		sp Q9BYX7 ACTBM_HUMAN
9348	SYELPDGQVITIGNER2000000000000000	sp P60709 ACTB_HUMAN
		sp P62736 ACTA_HUMAN
		sp P63261 ACTG_HUMAN
		sp P63267 ACTH_HUMAN
		sp P68032 ACTC_HUMAN
		sp P68133 ACTS_HUMAN
9349	TTCMSSQGSDDQIKR200000000000	sp Q9P0V9 SEP10_HUMAN
9350	ATSEVPGSQASPNPVPDGLHR200000000000	sp Q96GM8 TOE1_HUMAN
9351	HSSDINHLYTQGRESPEGSYDDANQEVNR200000000000	sp Q5TON5 FBP1L_HUMAN
9352	SRGSGHSQEQPAPQPSGGDPSPPQER2E+23	sp Q8WXE0 CSKI2_HUMAN
9353	ATSPSSSVSGDFDDGHHSVSTPGPSR2E+25	sp Q86U86 PB1_HUMAN
9354	HVTLPSSPR22000	sp Q5T4S7 UBR4_HUMAN
9355	AEQPSPNSDSGQDAHPDPDANPDAAR2.000002E+23	sp Q9BUA3 CK084_HUMAN
		sp P22612 KAPCG_HUMAN
9356	TWTLCGTPEYLAPEIILSK2E+19	sp P22694 KAPCB_HUMAN
		sp P17612 KAPCA_HUMAN
9357	LGMLSPEGTCK1020000000	sp P49327 FAS_HUMAN
9358	AVAITQSPSSVR2000000	sp P15822 ZEP1_HUMAN
9359	SFELPPYTPPPILSPVR300000020000	sp Q6PJG2 EMSA1_HUMAN
9360	FICVPTTCSNTIDLPMSPR2000	sp P40763 STAT3_HUMAN
9361	CALSSPSLAFTPPIK200000200000	sp Q8NFH5 NUP53_HUMAN
9362	SFSISPVR2020000	sp P18583 SON_HUMAN
9363	NKPGPNIESGNEDDDASFK200000002000	sp Q060841 IF2P_HUMAN
9364	SHTSEGAHLDITPNSGAAGNSAGPK2.02E+25	sp Q92597 NDRG1_HUMAN
9365	GGGAGGSSVGTGGGGTGGVGGGAGSEDSGDR20000000	sp Q06587 RING1_HUMAN
9366	LSSSPGATVSPSSSDAEYDKLPDADLAR2.0202E+21	sp Q8NI08 NCOA7_HUMAN
9367	NFKPSPSK20000	sp Q9BX63 FANCI_HUMAN
9368	GPSPSPVGPASVAQSR2000002000000000	sp Q014497 ARI1A_HUMAN
9369	SLSPLGGRDDSPVSHR20200000002000000	sp Q14004 CDK13_HUMAN
9370	SETSSNPSSPEICPNK2200000000	sp Q684P5 RPGP2_HUMAN
9371	GQEVETSVTYR20000000	sp Q043169 CYB5B_HUMAN
9372	RLTSPAPR2200000	sp Q6A6NMS7 L37A1_HUMAN
		sp P10636 TAU_HUMAN
9373	VAVVRTPPKSPSSAK20002000000	sp TAU_HUMAN
9374	NSVLSDPGLDSPRTSPVIMAR200000000020000000	sp Q015085 ARHGB_HUMAN
9375	SPGSTPTTSSQAPQK20000000000000	sp P35658 NU214_HUMAN
9376	SPQPDPVDTPASTK200000002000000	sp P46013 KI67_HUMAN
9377	GRLSVASTPISQR20002002000	sp Q9BXS6 NUSAP_HUMAN
9378	ASPTLDFTER2000000000	sp Q043426 SYNJ1_HUMAN
9379	EEGAAPSPHVAELATMEVESAR2000000000000000	sp Q9NRS6 SNX15_HUMAN
9380	KPGPPLSPEIRSPAGSPELR200002000200000	sp Q96JM3 CHAP1_HUMAN
9381	SPQTLAPVGEDAMKTPSPAEDAREPEAK2.0000000000102E+29	sp Q92797 SYMPK_HUMAN
9382	EGDPVSLSTPLETEFGSPSELSR2E+19	sp Q9BUR4 WAP53_HUMAN
9383	DDISEIQLSLASDHSGRSHDRPPR20020020000000	sp Q07157 ZO1_HUMAN
		sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
9384	GGSYSQAASSDSAQGSVSLTACK3.00000002E+21	sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN

9385	RNSNSPPSPSSMNQR200200000000	sp Q7Z5L9 I2BP2_HUMAN
9386	QESCLGNSPPFEK2000000	sp Q86W56 PARG_HUMAN
9387	KKSSSEAKPTSLGLAGGHK2000002000000000	sp Q96RK0 CIC_HUMAN
9388	KVSWSGR200000	sp P51798 CLCN7_HUMAN
9389	SGGIVISPF20000	sp Q16204 CCDC6_HUMAN
9390	SQSAAVTPSSTTSSTR200000000000000000	sp Q16186 ADRM1_HUMAN
9391	MNEEISSDSESESLAPR1000022000000000000	sp 043818 U3IP2_HUMAN
9392	ITPPAAKPGSPQAK200000	sp P35658 NU214_HUMAN
9393	KRPESSDYDLSPAK20000020000	sp Q6W2J9 BCOR_HUMAN
9394	KSPSDTEGLVK200000000	sp 095297 MPZL1_HUMAN
9395	SLQTGVGELHGETR200000000000000000	sp Q13393 PLD1_HUMAN
9396	VSPSPTYR200000000	sp P11274 BCR_HUMAN
9397	NATLAWDSSHSSIQNSPK200002000	sp 015440 MRP5_HUMAN
9398	SRSVSPCSNVESR2020000000000	sp Q9UQ35 SRRM2_HUMAN
9399	SKSPPKSPEEEGAVSS2000002000000000000	sp Q01130 SRSF2_HUMAN
9400	NEENTEPGAESSENADDPNKDTSENADGQSDENKDDYTIPDEYR2E+33	sp P43243 MATR3_HUMAN
9401	APLKPYVPS SDK200000	sp P41252 SYIC_HUMAN
9402	TVDSQGPTPVCTPTFLER2000000000000	sp Q9BYG3 MK67I_HUMAN
9403	AAPSTAPAEATPPKPGAEAPPK2000000000000000	sp 095817 BAG3_HUMAN
9404	SPPASPESWK200000000000	sp Q96JM3 CHAP1_HUMAN
9405	KVPCSPDFQR2000000	sp Q68DQ2 CRBG3_HUMAN
9406	EKEEPPSPIEATPPQSLLEK2000020000000000	sp 075151 PHF2_HUMAN
9407	FIQELSGSSPK2000	sp Q01664 TFAP4_HUMAN
9408	SLDGAEF SRPASVSENHDAGPDGDK20002000000000000000	sp Q9H4G0 E41L1_HUMAN
9409	QPATPTAAESSEGE GEGDDGGETESR2. 020002E+24	sp Q96S55 WRIP1_HUMAN
9410	HQSESQGVLS DK2000	sp P38398 BRCA1_HUMAN
9411	LQQQHSEQPPLQSPVMTR200000002000000	sp Q5JTV8 TOIP1_HUMAN
9412	YSPPAISPLVSEK2000020000000	sp Q14694 UBP10_HUMAN
9413	LGAVGSGGFK20000	sp 095163 ELP1_HUMAN
9414	IPLKPSPSK20000	sp Q99755 PI51A_HUMAN
9415	SAPASPNHAGVLSAHSSGAQTPELSR2E+27	sp A2A3N6 PIPSL_HUMAN
9416	GQSQTSPDHRSDTSSPEVR202000000000200000	sp Q01167 FOXK2_HUMAN
9417	YHGHSMSDPGVSYR200002000	sp Q9UQ35 SRRM2_HUMAN
9418	SPQPDPVK200000000	sp P29803 ODPAT_HUMAN
9419	AGGDLSPGR20000	sp P08559 ODPA_HUMAN
9420	AKSPISSGSGSHMSGTSSSSGMK2E+22	sp P46013 KI67_HUMAN
9421	RGNDPLTSSPGR2020000	sp Q9BUH8 BEGIN_HUMAN
9422	LSSLRASTKSSESSQK2000000002022000	sp Q15648 MED1_HUMAN
9423	RRSPSPYYSR202000000	sp P49736 MCM2_HUMAN
9424	ELPSQASGLANPPPVNLEDSLDEDLIR200000000	sp P62753 RS6_HUMAN
9425	KTSASDVTNIYPGDAGK2000000000000000000	sp Q13595 TRA2A_HUMAN
9426	SPSAIPEQNHSLNDQAK2000000000000000000	sp P62995 TRA2B_HUMAN
9427	TKPTQAAGPSSPQKPPTPEETK2000000000200	sp Q9NPG3 UBN1_HUMAN
9428	YQEDSDPERSDYEEQLQK3. 000000002E+19	sp Q15042 RB3GP_HUMAN
9429	GSSWLQDGSAR2000	sp Q13129 RLF_HUMAN
9430	GPAETAAADSEPDTDPEER20000000020000000	sp P18887 XRCC1_HUMAN
9431	RVSGDAAQDLDR200000000000	sp Q6P6C2 ALKB5_HUMAN
9432	LTQTSGQSTHTHKEPASGDEGIK200000000	sp 075676 KS6A4_HUMAN
9433	QEDSESESEESDSEEAASPAQVK2. 022000002E+21	sp Q5XUX0 FBX31_HUMAN
9434	EDLTDLEQPGSPGYSTATEPGSSELGVPEQPDLQEGTHVEK2E+31	sp Q9Y4H2 IRS2_HUMAN
9435	KRESESESDETPPAAPQLIK200000000000	sp P46013 KI67_HUMAN
		sp Q13428 TCOF_HUMAN
		sp Q92538 GBF1_HUMAN
		sp 060832 DKC1_HUMAN

9436	YSPTSPTYSPPTPK300020030000000	sp P24928 RPB1_HUMAN
9437	ESEESDDDMGFGLFD200200010000000	sp P05386 RLA1_HUMAN
		sp P05387 RLA2_HUMAN
9438	SRGQGYVGGQR200000000000	sp Q1KMD3 HNRL2_HUMAN
9439	ISLPLPNFSSLNLR2000000000000	sp P08670 VIME_HUMAN
9440	RHSHSHSPMSTR202010000	sp P62995 TRA2B_HUMAN
9441	TTPPPR20200000	sp Q8IYB3 SRRM1_HUMAN
9442	SETAAPAAAPPAEK20000000000000	sp P16403 H12_HUMAN
9443	TLSVAAAFNEDEDSEPEEMPPEAK200001000000	sp Q8WW12 PCNP_HUMAN
9444	FKAEAPLPSPK2000	sp Q09666 AHNK_HUMAN
9445	RPAEATSSPTSPERPR2000000000	sp Q9BU76 MMTA2_HUMAN
9446	KAENAEGQTPAIGPDGEPLDETSQMSDLPVK2000000	sp P51532 SMCA4_HUMAN
9447	NASVPNLR2000000	sp Q9H019 MFR1L_HUMAN
9448	SPSPYYSR200000000	sp P62995 TRA2B_HUMAN
		sp Q13595 TRA2A_HUMAN
9449	WTSQHSNTQTLGK2000000	sp Q6PKG0 LARP1_HUMAN
9450	SDNSDIRPSGPPPTLPASPSNHVSSLPPFIAPPGR2.00002E+27	sp Q8NEZ4 KMT2C_HUMAN
9451	ASTPDWVSEGPQPGLR2200000000000000	sp Q8IVT2 MISP_HUMAN
9452	SWDSSSPVDRPEPEEAAPTTR2220000000000200000	sp Q86WB0 NIPA_HUMAN
9453	SPQRSPLQSAESSPTAGK2000200020000000000	sp Q8IWB9 TEX2_HUMAN
9454	SPQPDPVDTASTK200000002000200	sp P46013 KI67_HUMAN
9455	GLSNAGDPEIVK2000000000000	sp O43847 NRDC_HUMAN
9456	SLSRPSSLIEQEK220000000	sp Q12802 AKP13_HUMAN
9457	TAHNSEADLEESFNEHELEPSSPK22000	sp Q8IWS0 PHF6_HUMAN
9458	HSQSPPEPHSPTEPPAWGSSIVK2.0000002E+19	sp Q5VZ89 DEN4C_HUMAN
9459	QQDLHLESPQRQPEYSPESPR200000030002000	sp Q9H6F5 CCD86_HUMAN
9460	SSRSVEDDKGHLVCR2200000000000000	sp P49761 CLK3_HUMAN
9461	KAASPSQSVR202002000	sp Q8IYB3 SRRM1_HUMAN
9462	RASHTLLPSHR2000000000	sp Q9UKV3 ACINU_HUMAN
9463	TSCRLSGGLGAGSCR200020000000000	sp Q04695 K1C17_HUMAN
9464	LSPCVPAKPPLAEFEGLDR2E+20	sp Q15742 NAB2_HUMAN
9465	SSPGQPEAGPEGAQERPSQAAPAVEAEGPGSSQAPR2E+19	sp P40222 TXLNA_HUMAN
9466	SDGAPASDSKPGSSEAAPSSK200000000	sp P80723 BASP1_HUMAN
9467	VPPAPVCPSPSPGSAVPSSPK200002000	sp O95817 BAG3_HUMAN
9468	SPDLEASLDLENNCHVGSIDFRPK2E+26	sp Q13111 CAF1A_HUMAN
9469	VESEESGDEEGKK200200000000	sp Q15435 PP1R7_HUMAN
9470	DRSPHRSSPSDTRPK2200000000	sp Q8IWS0 PHF6_HUMAN
9471	RESKPEEPPPK2000000000	sp O00255 MEN1_HUMAN
9472	GQEEISGALPVASSR2000000	sp Q9UBK8 MTRR_HUMAN
9473	SRTASGSSVTSLDGTR202000000000	sp Q92597 NDRG1_HUMAN
9474	SDASSPTISPASSP200220000000000	sp Q9Y2Y9 KLF13_HUMAN
9475	SGEQITSPVSPK20000002002000	sp O43314 VIP2_HUMAN
9476	TGQAGSLSGSPKPFSPQLSAPITTK2000020000000000	sp Q9ULU4 PKCB1_HUMAN
9477	DFLSGPDYDHTVGSDEDEEVEGEP2.000000002E+24	sp Q6ZW31 SYDE1_HUMAN
9478	NMVDLVNTHHLHSSDDEDDRLK1.00000000022E+22	sp Q9UPN7 PP6R1_HUMAN
9479	DLPSPDSR200000	sp Q9UGU0 TCF20_HUMAN
9480	SRSPVDLR202000000	sp Q13523 PRP4B_HUMAN
9481	SVSDPVEDK2000000000	sp Q8NFC6 BD1L1_HUMAN
9482	EQSEVSVSPR200000	sp Q7L4I2 RSRC2_HUMAN
9483	SEEEQSSSVK200000000000	sp P07910 HNRPC_HUMAN
9484	QQDLHLESPQRQPEYSPESPR20000002002000	sp Q9H6F5 CCD86_HUMAN
9485	TPLSFTNPLHSDSDSDERNSDGAVTQNK2.0020200002E+19	sp Q05209 PTN12_HUMAN
9486	NAVALSASPQLK200000	sp Q92547 TOPB1_HUMAN
9487	VISDESIDIGSDVEFKPDTK2000000000000000	sp P52701 MSH6_HUMAN

9488	MNEEISSDSESESLAPR100000202020000000	sp 043818 U3IP2_HUMAN
9489	SASQSSLDKLDQELK20000000000	sp 060271 JIP4_HUMAN
9490	TEDGGFEFEGASENNAKESPEK220000	sp 043719 HTSF1_HUMAN
9491	SAASPVVSSMPERASESSSEEKDDYEIFVK2.00000000000002E+30	sp P18583 SON_HUMAN
9492	VDINTPDVDVHGPDWHLK20000000000000	sp Q09666 AHNK_HUMAN
9493	DTHPGSPGSPGALRPLPAR2002000000000000	sp Q8N0Y2 ZN444_HUMAN
9494	STESVKSTVSETYLSKPSIAK2.020022E+20	sp Q7Z6B7 SRGP1_HUMAN
9495	SQRSNSPLPVPPSK2020202000000000	sp Q13247 SRSF6_HUMAN
9496	SLHSAHSLASR200200000200	sp 015027 SC16A_HUMAN
9497	INFSDLQQR2000000	sp Q15154 PCM1_HUMAN
9498	TAHNSEADLEESFNEHELEPSSPK2.0000002E+20	sp Q8IWS0 PHF6_HUMAN
9499	APSDSSLGTPSDGRPELR20000000000000	sp Q15642 CIP4_HUMAN
9500	SCDELSPVSPTQGGYPSEPTR200000000000	sp Q9HCD6 TANC2_HUMAN
9501	SDASSPTISPASSP20200000000	sp Q9Y2Y9 KLF13_HUMAN
9502	SKSPPKSPEEGAVSS200000000000020	sp Q01130 SRSF2_HUMAN
9503	AKSLDGVTNDR2000000000	sp Q9UBM7 DHCR7_HUMAN
9504	LSPSPTTEDPR200000000	sp Q9BTC8 MTA3_HUMAN
9505	TDHGAEIVYKSPVSGDTSR200000002000	sp P10636 TAU_HUMAN
9506	FSPPPVGQK20000000000	sp TAU_HUMAN
9507	QSGRESPLASR200000000000	sp Q9HCD6 TANC2_HUMAN
9508	APQTSSPPVVR200000000	sp Q8WWM7 ATX2L_HUMAN
9509	LGGLRPESPELTSVSR20000000200	sp Q8IYB3 SRRM1_HUMAN
9510	GLRDSHSSEDEASSQTDLSQTSK2.2000002E+19	sp Q9H6F5 CCD86_HUMAN
9511	SQDATFSPGSEQAEKSPGPIVSR2.00000000002E+19	sp Q5JTV8 TOIP1_HUMAN
9512	SPPEPPDFCCPK20000000000000	sp Q86WB0 NIPA_HUMAN
9513	SRDSGDENEPIQER200000000000	sp Q9Y6K9 NEMO_HUMAN
9514	RPDHSGGSPSPTSEPAR200202000000000	sp Q8WX93 PALLD_HUMAN
9515	LPSLGNKEPPGQEK20000000000000	sp Q8TAD8 SNIP1_HUMAN
9516	RPGFSPTSHR2000000	sp Q9UPY5 XCT_HUMAN
9517	QQAAYYAQTSPQGMPQHPPAPQGQ3E+19	sp Q8WUA4 TF3C2_HUMAN
9518	AQTPISSLPTSPSDEVGR20000200000000	sp Q96AE4 FUBP1_HUMAN
9519	SVSNGTAKPATSENFDEDLK200200000000000000	sp Q15751 HERC1_HUMAN
9520	KLSDINQEEASGTSLHQK2000000000000000	sp Q96K49 TM87B_HUMAN
9521	SAFTPATATGSSPSPVLGQGEK2.002E+19	sp Q4L235 ACSF4_HUMAN
9522	VDHGAEIITQSPGRSSVASPR20200022000000	sp Q15811 ITSN1_HUMAN
9523	AGGGPTLQCPPSSPEK200000	sp P11137 MTAP2_HUMAN
9524	SPSPSPTPEAK200000000000	sp Q96N66 MBOA7_HUMAN
9525	GEGDAPFSEPGTTSTQRPSPETATK20000000000000	sp Q8TAQ2 SMRC2_HUMAN
9526	SELCSPLGK200000	sp Q9UQ35 SRRM2_HUMAN
9527	VLSQSTPGTPSK200	sp Q9Y2F5 ICE1_HUMAN
9528	SSGHSSSELSPDAVEK2220000000000	sp Q6MZP7 LIN54_HUMAN
9529	KQSQIQNQQGEDSGDPEDTY2020000000	sp Q9UQ35 SRRM2_HUMAN
9530	SPQTLAPVGEDAMKTPSPAEDAR1020200000000	sp P18858 DNLI1_HUMAN
9531	TKPTQAAGPSSPKPPTPEETK20000000000000	sp Q92797 SYMPK_HUMAN
9532	NATQPPNAEEESGSSSASEEEDTKPKPTKR2.022202E+19	sp P18887 XRCC1_HUMAN
9533	SRSFTLDDSLK2020000000000	sp Q68E01 INT3_HUMAN
9534	DRSPPKSPEK200020000	sp Q86WR7 PRSR2_HUMAN
9535	DGPISPK2000	sp Q9UQ35 SRRM2_HUMAN
9536	EGEETEGSEDDKENDKTEEMPNDVLENK2.0000000000002E+26	sp Q8WWQ0 PHIP_HUMAN
9537	SHSDNDRFK2020000000	sp P49321 NASP_HUMAN
9538	TLLSSDDEF2200000	sp 075494 SRS10_HUMAN
9539	RLSSTSLASGHSVR202000000000	sp Q9Y519 T184B_HUMAN
9540	RLSSASTGKPLSVEDDFEK200000000020000000	sp Q9BZL6 KPCD2_HUMAN
		sp 075152 ZC11A_HUMAN



9541	DTLSNPQSPQSPYNSPKPQHK2. 0000000203E+19	sp	Q96PU5	NED4L_HUMAN
9542	RRTSPPPR200000	sp	Q8IYB3	SRRM1_HUMAN
9543	SISGTPAETPACK2000000000	sp	Q9Y4D8	HECD4_HUMAN
9544	SDSEESGSEEEEEEEEQPAAQPPTLPVEEK2. 00202E+31	sp	P51532	SMCA4_HUMAN
9545	RQSPSPSTRPIR20200000000	sp	Q8IYB3	SRRM1_HUMAN
9546	KTPSKPPAQLSPSVPK2000000002000000	sp	P18887	XRCC1_HUMAN
9547	NSVLSDPGLDSPRTSPVIMAR200200000000	sp	O15085	ARHGB_HUMAN
9548	DQPPFGSDDSVEADKSSPGIHLER2002000000200000000	sp	Q14676	MDC1_HUMAN
9549	RPACDPEPGESGSSSDEGCTVVRPEKK202200000000000000	sp	O15541	R113A_HUMAN
9550	HGSILNR200000	sp	Q9NZN5	ARHGC_HUMAN
9551	QRSHSGTSPDNTAPPPPPRPPQPSHSR2. 02E+25	sp	Q96D71	REPS1_HUMAN
9552	YPRPASVPPSPSLSR20002000200	sp	P13807	GYS1_HUMAN
9553	TPPVALNSSR20000002000	sp	Q9UQ35	SRRM2_HUMAN
9554	QRGSETGSETHESDLAPSDK200000000000	sp	P48634	PRC2A_HUMAN
9555	EWYQSTIPQSPSPAPDDPEEGR200000000000	sp	Q08499	PDE4D_HUMAN
9556	SATRPSPSPER200002020000	sp	Q9UQ35	SRRM2_HUMAN
9557	SLESSVIMDR22001000	sp	Q9H2Y7	ZN106_HUMAN
9558	GSLGISQEEQ200000	sp	Q9NZD8	SPG21_HUMAN
9559	SVSPESNDISEELNHFKPIVCSPTPK2. 02002E+29	sp	Q8NCN4	RN169_HUMAN
9560	LLLERPSPIR20000	sp	Q96T37	RBM15_HUMAN
9561	RPAEATSSPTSPERPR20000000	sp	Q9BU76	MTA2_HUMAN
9562	GSSGSPAHAESYSSGGGGQK2. 002E+20	sp	P15408	FOSL2_HUMAN
9563	EGEETEGSEEDDKENDKTEEMPNDVLENK2. 0000000002001E+23	sp	P49321	NASP_HUMAN
9564	QEKPAEKPAETPVATSPTATDSTSGDSSR2000000	sp	P54727	RD23B_HUMAN
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9566	SLSSDSTLLLFK20020000000	sp	Q9HAW4	CLSPN_HUMAN
9567	SVVSFDKVK2002000000	sp	Q15424	SAFB1_HUMAN
9568	RPAGGRPSPSAMGK20001000	sp	Q92615	LAR4B_HUMAN
9569	SASPEVSEGHENQHGESEAK2E+19	sp	O60930	RNH1_HUMAN
9570	GSSDGRGSDSEDLPHR2000000000000000	sp	Q9UPQ0	LIMC1_HUMAN
9571	MNEEISSDSESESLAPR100002000020000000	sp	O43818	U3IP2_HUMAN
9572	RPSVDAPVTDVGFRLR20000000000000	sp	Q9BUH8	BEGIN_HUMAN
9573	TDKSSASAPDVEDPEAFPALA2E+21	sp	Q8NC51	PAIRB_HUMAN
9574	EGNTTEDDFSSPGNGNK200000000	sp	Q15007	FL2D_HUMAN
9575	YLPLNTALYEPPLDPELPALDSDGSDGDEGRGDEK200020000000000000	sp	Q8WUB8	PHF10_HUMAN
9576	KHSDPHLLER200000000	sp	Q9Y6M7	S4A7_HUMAN
9577	RPAEATSSPTSPERPR20022000000	sp	Q9BU76	MTA2_HUMAN
9578	DVSGFSDPYCLLGIEQGVGVPGGSPGSR200	sp	Q70J99	UN13D_HUMAN
9579	SASEPSLHR2000000000	sp	P10398	ARAF_HUMAN
		sp	P04049	RAF1_HUMAN
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9581	LQQAALSPTTAPAVSSVSK200000000000	sp	Q8WXI9	P66B_HUMAN
9582	GPSQATSPIR200000	sp	Q7RTP6	MICA3_HUMAN
9583	SATSTSASPTLR2000000002000	sp	O75061	AUXI_HUMAN
9584	SVSHGSNHTQKPDEQR202000000000000000	sp	Q9Y520	PRC2C_HUMAN
9585	DMESPTKLDVTLAK2000000000	sp	P27816	MAP4_HUMAN
9586	HVSTSSDEGSPSASTPMINK200000000000	sp	Q7L1W4	LRC8D_HUMAN
9587	QNSPVTQDGIQESPWQDR20000000000000	sp	O75815	BCAR3_HUMAN
9588	GKLSAENPDDSEVPSSGINSTK20000000000000	sp	Q9Y5Q9	TF3C3_HUMAN
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9590	YGLQDSDEEEEHPSK300000000000000000	sp	P52948	NUP98_HUMAN
9591	QSPGHQSPLASPK2000020000000	sp	Q8N1G0	ZN687_HUMAN
9592	DEVFHTDQDDPSSDDEGMPYTRPVK220000100000000	sp	Q5JSH3	WDR44_HUMAN

9593	GSGSEQEGEDEEGGER2020000000000000	sp Q6PD62 CTR9_HUMAN
9594	TAPDTKPSPR2000	sp P22105 TENX_HUMAN
9595	DSPDSPEPPNKKPLVEMDETPQVEK2E+21	sp Q9NVM9 ASUN_HUMAN
9596	ENPPVEDSSDEDDKRNQGNLYDK2200000000000000	sp Q8TCJ2 STT3B_HUMAN
9597	TGPLSTSSPSR20000	sp Q8IY92 SLX4_HUMAN
9598	ARSVSPPPKR200000000	sp Q13247 SRSF6_HUMAN
9599	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2.00200000002E+32	sp Q8WUZ0 BCL7C_HUMAN
9600	NSSSPVSPASVPGQR22000000000000	sp Q9H1B7 I2BPL_HUMAN
9601	VQRPHSSPPR200000	sp Q9NUJ3 T11L1_HUMAN
9602	AGAVTSTPNRNSSK2000	sp Q9ULR3 PPM1H_HUMAN
9603	RVPSINQK200000	sp Q92834 RPGR_HUMAN
9604	YLCNSPQKPNDR200000000	sp O14975 S27A2_HUMAN
9605	HVLQTAVADSPR2000	sp Q7Z2Z1 TICRR_HUMAN
9606	GDAASSPAPAASVGSQGGAR20000000002000000	sp Q8WVB6 CTF18_HUMAN
9607	EEDCHSPTS KPPKPDQLK200000000000	sp P23588 IF4B_HUMAN
9608	EALNIIGDISTSTVSTVPPVDDTWLQSASSHSPTQR2000000000000000 0	sp P50570 DYN2_HUMAN
9609	RTSFSFQK2000000	sp Q9BPZ7 SIN1_HUMAN
9610	TIETSPSLSR2000000	sp Q9NVI1 FANCI_HUMAN
9611	RHSSETFSSTTTVTPVSPFAHNPK202002000000000	sp Q9Y4H2 IRS2_HUMAN
9612	QSPGHQSPLASPK2000020002000	sp Q8N1G0 ZN687_HUMAN
9613	SPMSTNSSVHTGSDVEQDAEKK2.000000000002E+22	sp Q9ULU4 PKCB1_HUMAN
9614	APPPISPTQLSDVSSPR20020002000	sp Q9P0K7 RAI14_HUMAN
9615	LAAAASPHSGGR20020000	sp O95613 PCNT_HUMAN
9616	TSPTPEASATNSPCTSKPATPAPSEK2.0002E+23	sp Q12873 CHD3_HUMAN
9617	ALSSDSILSPAPDAR20020000000	sp Q8IVT2 MISP_HUMAN
9618	LLEDGEDFNLGDALDSSNSMQTIQK21000000	sp P05783 K1C18_HUMAN
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9620	AGEPNSPDAEANS PDVTAGCDPAGVHPPR20000000000000	sp Q08J23 NSUN2_HUMAN
9621	LLRPNSLR2000	sp Q8WXG6 MADD_HUMAN
9622	AGGSPAPGPETPAISPSK2000000000000000	sp P33316 DUT_HUMAN
9623	KPSDSLVA SSSR200000002200	sp Q5T0W9 FA83B_HUMAN
9624	YSPTSPTYSPSPK220030000000	sp P24928 RPB1_HUMAN
9625	KLPGSDSENEELLNGHASDSENEVDVGK2.020000000002E+23	sp Q96ST2 IWS1_HUMAN
9626	SLSSPGGPSKPK20000000000	sp Q9BZE9 ASPC1_HUMAN
9627	RYSGSDSSASSAQSGPLGTR2000000000000000	sp Q99501 GA2L1_HUMAN
9628	TRSPSPTLGESLAPHK20000020000000000	sp Q86UU1 PHLB1_HUMAN
9629	DSSSLSSCTSGILEER22000000000	sp Q9H3Q1 BORG4_HUMAN
9630	LFDEEEDSSEK22000	sp O43719 HTSF1_HUMAN
9631	SGSALLQSSTEDPK2200000	sp Q5VTR2 BRE1A_HUMAN
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9633	SPQPPVEEEDHFDDTVVCLDTYNCDLHFK2000000000000000	sp Q00839 HNRPU_HUMAN
9634	EPGSGTESDTPDFHNQENEPSQEDPEDLDGSVQGVKPKK2.0202E+37	sp Q9HAF1 EAF6_HUMAN
9635	SRPTSEGS DIESTEPQK200020000000000	sp Q4G0J3 LARP7_HUMAN
9636	APSASDSKADSDGAKPEPVAMAR2.0200020002E+23	sp Q7Z4V5 HDGR2_HUMAN
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9638	SSWHQSSFHSTR200002000000	sp Q99959 PKP2_HUMAN
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9641	RSSQPSPTAVPASDSPPTK2200000000000000000	sp Q3KQU3 MA7D1_HUMAN
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9643	KEESP PPK2000000	sp P78332 RBM6_HUMAN
9644	SLNSVVTSEPDKHSLLVGDFR2.0000002000002E+21	sp Q04656 ATP7A_HUMAN
9645	EVEEDSEDEEMSEDEEDDSSGEEVVIPQKK2.0000120000002E+25	sp P19338 NUCL_HUMAN

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9647	TLSVAAAFNEDEDSEPEEMPPEAK2E+22	sp	Q8WW12	PCNP_HUMAN
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9649	STPFIVPSSPTEQEGR2000000200000000	sp	Q12888	TP53B_HUMAN
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9666	ASPATQPPPLLPPSATGPDATVGGPAPTLLPPSATASVK2E+39	sp	Q07666	KHDR1_HUMAN
9667	TGVPSTASVGK20020000	sp	P28290	SSFA2_HUMAN
9668	RNSSSPVSPASVPGQR2220000000000000	sp	Q9H1B7	I2BPL_HUMAN
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9670	SRTSVQTEDDQLIAGQSAR220020000000000000	sp	P35221	CTNA1_HUMAN
9671	STPHRTPIITPNTK220000000200000	sp	P26232	CTNA2_HUMAN
9672	YRNEGSPSPR202000	sp	Q9NQW6	ANLN_HUMAN
9673	TEPAAEAEASGPSESPSPAEEELPGSHAEPVPAQGEAPGEQAR2E+19	sp	Q5T200	ZC3HD_HUMAN
9674	TEDGGWEWSDEFDEESEEGK2E+21	sp	P55884	EIF3B_HUMAN
9675	VSPSPQESLSSSK202002000000	sp	O95747	OXSR1_HUMAN
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9677	GSPATSPHLGR22000000	sp	Q9NYV4	CDK12_HUMAN
9678	RPILQLSPPGPR2000000	sp	Q13671	RIN1_HUMAN
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9680	SILVSPTGPSR200020000000	sp	Q96S55	WRIP1_HUMAN
9681	TLENSPISSCDTSDTEGPVNVNSAAVLK2. 2E+21	sp	Q14684	RRP1B_HUMAN
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9683	DRPPCSPSR20000	sp	Q9NVU7	SDA1_HUMAN
9684	GGNLPPVSPNDGAK200020000	sp	O95758	PTBP3_HUMAN
9685	ILENSEDSSPECLF22000000	sp	Q9UK61	F208A_HUMAN
9686	SETSSNPSSPEICPNK2002200000000	sp	Q9NRZ9	HELLS_HUMAN
9687	SRDESASETSTPSEHSAAPSPQVEVR2. 0002020222E+26	sp	Q684P5	RPGP2_HUMAN
9688	SSSSLLASPGHISVK2200000000000000	sp	Q92614	MY18A_HUMAN
9689	NTPSQHSHSIQHSPER2000020000020000	sp	A0FGR8	ESYT2_HUMAN
9690	DGSGTPSRHSLSGSSPGMK20202000000	sp	Q9NYF8	BCLF1_HUMAN
9691	SPPPRSPSPPTHR2000000002000000	sp	Q9UQ35	SRRM2_HUMAN
9692	LQVYEGGGQPPQDLPGSPK2000	sp	Q96HA1	P121A_HUMAN
9693	DSPGIPPSANAHQLFR2000000000000000	sp	Q86Y91	KI18B_HUMAN
9694	DAPTSPASVASSSTPSSK20000000000000200	sp	P51812	KS6A3_HUMAN
9695	QQSLPPP2000000	sp	Q04726	TLE3_HUMAN
9696	GHTASESDEQQWPEEK2000000000000000	sp	Q9UPQ0	LIMC1_HUMAN
9697	TEVSPVPR200000	sp	Q9NTI5	PDS5B_HUMAN
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		sp	Q8IVH2	FOXP4_HUMAN

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9704	LNTSDFQK2000000	sp Q96B36 AKTS1_HUMAN
9705	IPIPETPPQTPPVLDSPHQ20002000000200000	sp Q969H4 CNKR1_HUMAN
9706	VHSPASLVPR2000000000	sp Q9BTE3 MCMBP_HUMAN
9707	ASPVTSPAAAFPTASPANK200000	sp Q9UIF9 BAZ2A_HUMAN
9708	SCFESSPDELK220000000	sp Q9UQ35 SRRM2_HUMAN
9709	EGTLTQVPLAPPPGAPPSPAPAR2000000	sp P48634 PRC2A_HUMAN
9710	EYIPGQPPLSQSSDSSPTR200020000	sp P07814 SYEP_HUMAN
9711	NSTELAPPLPVR2000000000000	sp Q9H7D0 DOCK5_HUMAN
9712	SPPSYSVLYPSSDPK2000000000000000	sp Q8N3V7 SYNPO_HUMAN
9713	EVIEIEDASPTKCPITTK20000000000	sp P46100 ATRX_HUMAN
9714	RSPERPTGDLR200000	sp Q5T200 ZC3HD_HUMAN
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9716	EPAPLLSAQYETLSDSD20000000	sp 075376 NCOR1_HUMAN
9717	AASDPLSSVSPAVSK200000002000000	sp P19532 TFE3_HUMAN
9718	ISICSSDKR20000	sp Q13547 HDAC1_HUMAN
9719	RPHGSTLSVSASPEK20000	sp Q92738 US6NL_HUMAN
9720	EEASLLSHSPGTSNQSQPCSPKPIR2E+22	sp Q9H4Z3 PCIF1_HUMAN
9721	STPEDHRNSSTETDQPHSPDSSSSVHSIR20000020000000000000	sp Q92613 JADE3_HUMAN
9722	ADVQLFMDDSYSHHSGLEYADPEK1. 000030002E+19	sp P51636 CAV2_HUMAN
9723	SEVEGFSLPSR20000	sp Q08AD1 CAMP2_HUMAN
9724	ADGAPSAAPPDGLLASPDGLLK200000000	sp P17535 JUND_HUMAN
9725	SSSLKSAQGTGFELGQLQSIR2. 0000002E+19	sp Q9C0B5 ZDHC5_HUMAN
9726	SSPVNNHSPGQTPTLGQPEAR2. 00000020002E+21	sp Q5SW79 CE170_HUMAN
9727	AGLGSPPERPKTSPGSPR200000002002000	sp Q96L14 C170L_HUMAN
9728	STPACSPILR200000	sp Q9H6F5 CCD86_HUMAN
		sp Q3KR37 GRM1B_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
9729	EDALDSSVSSSSVHASPLASSPVRK2. 00000000000002E+19	sp 014715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
		sp A6NKT7 RGPD3_HUMAN
9730	SVVSFDK20000	sp Q15424 SAFB1_HUMAN
9731	QQLPQTPPSCLK20000000	sp Q8IY81 SPB1_HUMAN
9732	SNVSSPATPTASSSSSTPTR2000000000000000	sp Q12888 TP53B_HUMAN
9733	NHSPSPVPTPTGAAPSLASPK200020000000000000	sp Q5SYE7 NHSL1_HUMAN
9734	GTGSGGQLQDLDCSSDDEGAAQNSTKPSATK2E+31	sp P08240 SRPR_HUMAN
9735	SSPASSPELPWRPR2200020000000000	sp Q9NYF3 FA53C_HUMAN
9736	TRDDGDEEGLLTHSEEELEHSQDTDADDGALQ2. 0000002002E+19	sp Q9UKM9 RALY_HUMAN
9737	ILDELKDDSTHESLSQESESEEDGIHVDSQK2. 00202E+22	sp Q9H6T3 RPAP3_HUMAN
9738	SDSSQPMLLR20000000	sp P11532 DMD_HUMAN
9739	VVSSEGRPSVDLSFQPSQPLSK2E+20	sp P49815 TSC2_HUMAN
9740	SSGTNFQGLPSK2000000000	sp Q68EM7 RHG17_HUMAN
9741	SRTSPAPWK2002000000	sp Q9UQ35 SRRM2_HUMAN
9742	ALEAASLSQHPPSLCISDSEEEEEER2. 0000000000202E+21	sp 043159 RRP8_HUMAN
9743	AGGPATPLSPTR20020000	sp Q03252 LMNB2_HUMAN
9744	EDILENEDEQNSPPK20000	sp Q9NTI5 PDS5B_HUMAN
9745	AGEAPTENPAPPTQSSAE2000	sp P16989 YBOX3_HUMAN
9746	SGSALLQSQSTEDPKDEPAELKPDSEDLSSQSSASK2. 02E+30	sp Q5VTR2 BRE1A_HUMAN
9747	VAHEPVAPPEDKESESEAK20000	sp 095674 CDS2_HUMAN

9748	RGNDPLTSSPGR20000	sp P49736 MCM2_HUMAN
9749	RQSPEPSPVTLGR200000000000	sp Q9NRL2 BAZ1A_HUMAN
9750	KLPPPGSPLGHSPTASPPPTAR20000002000000000	sp O95785 WIZ_HUMAN
9751	SPSPPLPTHIPPEPRTPPFPAK2.00002000000002E+21	sp Q8IZ21 PHAR4_HUMAN
9752	SPPVQHPHTVITISLGTAPSLQNNQPVEFNHAINVYVK2E+37	sp Q96ST3 SIN3A_HUMAN
9753	SQSRNSPLPVPPSK20202000000000	sp Q13247 SRSF6_HUMAN
9754	SASPYPSHLSPPQR20300000020000	sp Q5T200 ZC3HD_HUMAN
9755	RPSASSPNNNTAAK2002000000000	sp O75151 PHF2_HUMAN
9756	EVAENQQNQSSDPEEEKGSQPPAAESQSSLR2E+22	sp O43491 E41L2_HUMAN
9757	SQSKSPTGTPAR2020200020000	sp Q13595 TRA2A_HUMAN
9758	GTEPSPGGTPQPSRPVSPAGPPEGVPEEAQPPR2.00000000002E+32	sp Q8WUZO BCL7C_HUMAN
9759	TPSSSSTLAYSPPR20000000000000	sp Q6Q0C0 TRAF7_HUMAN
9760	EFSGPSTPTGTLEFEGGEVSLEGGK2E+20	sp Q09666 AHNK_HUMAN
9761	SISSPNVNR20000000	sp Q9NQT8 KI13B_HUMAN
9762	QLSSTSPLAPYPTSQMVSSDR20000000000000000	sp Q6UUV7 CRTC3_HUMAN
9763	HPSGGSSPPPR2002000000	sp Q6IQ23 PKHA7_HUMAN
9764	VSPLEGESEGDGDTD20000200000200	sp Q9Y606 TRUA_HUMAN
9765	HQEVQDQDPVFGSDSSGYQSDHK20000	sp Q3B726 RPA43_HUMAN
9766	FSPVLGR2000000	sp Q9NRG9 AAAS_HUMAN
9767	TQSLPVTEK20000000	sp Q9NQW6 ANLN_HUMAN
9768	QLSALSSDGEQGGRPVPR20002000000000000	sp O15438 MRP3_HUMAN
9769	DGLAPEKTSPDRDK2000000	sp Q7L4I2 RSRC2_HUMAN
9770	VLANSNPSSPSAAK20000002000000	sp Q9NV56 MRGBP_HUMAN
9771	RGTGGSESSRASGLVGR2000022002000000	sp Q9UH99 SUN2_HUMAN
9772	STSYGYSR2000000	sp Q13242 SRSF9_HUMAN
9773	KKTVSFSSMPTEK200020100000	sp Q9UPR0 PLCL2_HUMAN
9774	SLSRTPSPPPFR2020202000000	sp Q7L4I2 RSRC2_HUMAN
9775	AISAPTSPTR200000200	sp Q9NQG6 MID51_HUMAN
9776	TPTPSTPGDTQPNTAPVPPAEDGIK2.000000002E+22	sp Q14839 CHD4_HUMAN
9777	MSGFIYQ GK1200000000	sp Q14155 ARHG7_HUMAN
9778	SCSSPAVSAVSQLPLSPK2000	sp Q15052 ARHG6_HUMAN
9779	NSASATPLSGNSSR2002000	sp Q9Y2F5 ICE1_HUMAN
9780	ETSVSKEDTDHEEK2000002000000	sp Q9Y608 LRRF2_HUMAN
9781	GSPRPQAPK200000000	sp O75475 PSIP1_HUMAN
9782	SPQQTVPYVVPLSPK2000	sp Q14978 NOLC1_HUMAN
9783	GPEVTSQGVQTSSPACK200000	sp Q9UQR0 SCML2_HUMAN
9784	SAEALGPGALVSPR200000000002000	sp Q99700 ATX2_HUMAN
9785	SRSPPVSKR20200000000	sp Q9P227 RHG23_HUMAN
9786	MAESPCSPSQPPSPSPDEL PANVK2.0000000002E+21	sp Q01130 SRSF2_HUMAN
9787	TSSKESPIPSPTSDR200220000000000	sp Q14160 SCRIB_HUMAN
9788	SSTPPRQSPSR2000020000	sp Q01082 SPTB2_HUMAN
9789	HPAGPPGEAQEGSAK2000	sp Q9UQ35 SRRM2_HUMAN
9790	EALNIIGDISTSTVSTVPPVDDTWLQSASSHSPTQR200000000	sp Q6P1M3 L2GL2_HUMAN
9791	GRLSVASTPISR20020000000	sp P50570 DYN2_HUMAN
9792	SVPGTTSSPLVGDISP K20000000002000	sp Q9BXS6 NUSAP_HUMAN
9793	SPACSSLTPSLCK20000000020000	sp Q9UQR0 SCML2_HUMAN
9794	SPTPNQDGD TMVEK200000000000000	sp Q96R06 SPAG5_HUMAN
9795	DGDSYDPYDFSDTEEMPQVHTPK2.0003000000001E+21	sp Q3KR37 GRM1B_HUMAN
9796	SSIAGSEDAEPLAPPIKPIKPVTR2E+19	sp P25205 MCM3_HUMAN
9797	AQSLVISPPAPSPR2000000002000	sp Q03164 KMT2A_HUMAN
9798	MAGNEALSPTSPFR220000	sp P46013 KI67_HUMAN
9799	TEMKSPFNPSPPQDSPR20002000000000	sp Q86WR7 PRSR2_HUMAN
9800	RNSNSPPSPSSMNQR20000002000000	sp P08651 NFIC_HUMAN
		sp Q7Z5L9 I2BP2_HUMAN

9801	SRSPIIHSPK20200002000	sp 094913 PCF11_HUMAN
9802	SPGSGSQSSGWHEVEPGMPSPTTLK200000000002000000	sp Q12857 NFIA_HUMAN
9803	EQAPDISPK2000	sp Q7Z333 SETX_HUMAN
9804	SLSFSEPQQPAPAMK20000000000000	sp Q9H8N7 ZN395_HUMAN
9805	SSSMSSLTGAYTSGIPSSSR200000000000000000	sp A8CG34 P121C_HUMAN
9806	SRTPPTSR202000000	sp Q9UQ35 SRRM2_HUMAN
9807	KDSSDRENPPHDSR22000000000000	sp Q92613 JADE3_HUMAN
9808	SSEYPISQNPGLSADK200000000000000000	sp P38398 BRCA1_HUMAN
9809	SPPSSSEIFTPAHEENVR20000000000000000000	sp C9JLW8 F195B_HUMAN
9810	SLDSEDEDEEDDYQQR20020000000000000000	sp Q13442 HAP28_HUMAN
9811	LNSSVLPYGAESSEDSDEESK3000020020000000	sp Q9H9J4 UBP42_HUMAN
9812	RPAEATSSPTSPERPR202000000000	sp Q9BU76 MMTA2_HUMAN
9813	SRSDIDVNAASAK2000000000000000	sp Q7Z460 CLAP1_HUMAN
9814	SEEVPAFGVASPPPLTDTPTDTANAEGDLPTTMGGPLPPHLALK2E+23	sp Q08211 DHX9_HUMAN
9815	GHYEVTGSDDDETGKLQSGVSLASK200	sp Q09666 AHNK_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
9816	EDALDDSVSSSVHASPLASSPVRK200020002000000	sp A6NKT7 RGPD3_HUMAN
		sp O14715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
9817	SPSPPPTRR2020000000	sp Q8IYB3 SRRM1_HUMAN
9818	TSPADHGGSVGSESGSAVDSVAGEHSVSGR2.000000200202E+30	sp Q5T4S7 UBR4_HUMAN
9819	KLPPPPGSPFLGHSPTASPPPTAR200002000000000000	sp O95785 WIZ_HUMAN
9820	KASSEGGTAAGAGLDSLHK20000000000000000000	sp O15143 ARC1B_HUMAN
9821	LGSSGLGSASSIQAAVR200000000000000000	sp Q7Z6Z7 HUWE1_HUMAN
9822	AAQQAASSSGGQQAQTPTGK200000	sp P48729 KC1A_HUMAN
9823	ERSDSGGSSSEPFDR2020020000000000	sp P48634 PRC2A_HUMAN
9824	SRTPLLPR200000000	sp Q9UQ35 SRRM2_HUMAN
9825	AHSSPASLQLGAVSPGTLTPTGVVSGPAATPTAQHLR2E+31	sp P46937 YAP1_HUMAN
9826	AQSTDSLGTSGSLQSK2200000000000000	sp Q15276 RABE1_HUMAN
9827	LPASPSGSEDLSSVSSPTSSPK20200000	sp Q9P2N6 KANL3_HUMAN
9828	STGSATSLASQGER220000000200000	sp Q5SW79 CE170_HUMAN
9829	QQPPLGPSSLLSLPGLK200002000000	sp Q4KMP7 TB10B_HUMAN
9830	SASPYPHSLSSPQR200020000000000	sp Q5T200 ZC3HD_HUMAN
9831	YIEIDSDEEPR2000000	sp Q9NVU7 SDA1_HUMAN
9832	ALEAASLQHPPLSLCISDSEEEEEER2.00000000002E+21	sp O43159 RRP8_HUMAN
9833	QLGDSSGEGPEFVEEEEEVALRSDSEGSYTPGK2000020300000	sp Q14839 CHD4_HUMAN
9834	GHHVTDSENDEPLNLNASDSESEELHR2.000000000002E+23	sp Q96ST2 IWS1_HUMAN
9835	ASPVSFQELNR200000000000	sp P15822 ZEP1_HUMAN
9836	SSPPLRTPDVLESSGPAVR2.2E+19	sp O75676 KS6A4_HUMAN
9837	SRSYSPAHR20230000000	sp Q9Y2W1 TR150_HUMAN
9838	LSGEHSESSTPRPR2002000000	sp Q7Z3K3 POGZ_HUMAN
9839	LQEEGGSDDEETGSPSEDGMQSAR200000000000	sp Q13610 PWP1_HUMAN
9840	YVTKPNSDDEDDGDEKEDDEEDKDDVPGPSTGGSLR2E+30	sp P51784 UBP11_HUMAN
9841	AASDPLSSVSPAVSK200000002000200	sp P19532 TFE3_HUMAN
9842	SRSGSSQELDVKPSASPQR2.02022E+20	sp Q9UQ35 SRRM2_HUMAN
9843	VLSTSSLTR2220000	sp P16144 ITB4_HUMAN
9844	QLSALSSDGEGQGRPVPVR200000000000000000	sp O15438 MRP3_HUMAN
9845	RRSPSLSSK20200000	sp Q8IYB3 SRRM1_HUMAN
9846	VNSNGKESPGSSEFFQEAVSHGK2.00002002E+21	sp Q8N108 MIER1_HUMAN
9847	SPPREGSQGELTPANSQSR2.000002E+19	sp Q13098 CSN1_HUMAN
9848	TLVITSTPASPVR20000	sp Q8TBN0 R3GEF_HUMAN
9849	ETDEAAFEFDYNESDSESNVSVK20200002000	sp Q7Z6E9 RBBP6_HUMAN
9850	APQTSSSPPVVR2002000000	sp Q8IYB3 SRRM1_HUMAN

9851	EPLIPASPK2000	sp Q9Y613 FHOD1_HUMAN
9852	KKSPIINESR200000200	sp Q13523 PRP4B_HUMAN
9853	VEEVEDVSPGPWGLVK2000000000	sp O15061 SYNEM_HUMAN
9854	TDSEKPFRRSQSPK2000000002000	sp Q9Y2W1 TR150_HUMAN
9855	TPQEWAPQTAR200000000000	sp Q5JTV8 TOIP1_HUMAN
9856	RASISEPSDTPDPEPR20000000000000	sp Q8N1F8 S11IP_HUMAN
9857	IPSIVSSPLNSPLDR20000200000	sp P49790 NU153_HUMAN
9858	SPSLLQSGAK200020000	sp Q9BQG0 MBB1A_HUMAN
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9860	KHSVSPPPYPGHTK2020000000000000	sp Q9BZD6 TMG4_HUMAN
9861	LSGEHSESSTPRPR2020000000	sp Q7Z3K3 POGZ_HUMAN
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9863	SYISPHSSPHTPTR2200000000	sp Q96BY7 ATG2B_HUMAN
9864	VNSNGKESPGSSEFFQEAVSHGK2.000000002E+21	sp Q8N108 MIER1_HUMAN
9865	EDSSDSENAEPDLDDNEDEEPAVEIEPEPEPQVPTAPPPAK2.202E+41	sp P49711 CTCF_HUMAN
9866	VSEVEEKEPVPQPLPSDDTR2E+20	sp O43290 SNUT1_HUMAN
9867	RDSFDDRGPSLNPVLDYDHGSR2E+20	sp P43243 MATR3_HUMAN
9868	RYSGSDSDSISEGK30000202000000	sp O14646 CHD1_HUMAN
9869	IISPGSSTPSSTR220000000	sp Q5VT52 RPRD2_HUMAN
9870	KGSGKRPAAGASASPR200000000000002000	sp Q96S55 WRIP1_HUMAN
9871	KNESSDEISDAEQMPQHTLK220002000000000000	sp Q3L8U1 CHD9_HUMAN
9872	YYSDDDELTVQR30020000000000	sp Q8NFC6 BD1L1_HUMAN
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9875	TPESTKPGPVCQPPVSQSR2E+19	sp Q6AI12 ANR40_HUMAN
9876	GYSPYSVSGSGSTAGSR20000000000000	sp Q15149 PLEC_HUMAN
9877	RESESEDETPPAAPQLIKK2020200000000000000	sp O60832 DKC1_HUMAN
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9879	KETPPSPFESDK20000000000	sp Q96BT7 ALKB8_HUMAN
9880	SLVMHTPPVLK200002000000	sp P46013 KI67_HUMAN
9881	YISENESPRPSQK20000000	sp O94782 UBP1_HUMAN
9882	ETINLDTSSSDFTILQEIEEPSLEPENEK2E+29	sp Q15424 SAFB1_HUMAN
9883	LDSSMDHSENYDTMSSPLPGK2.0010000000301E+21	sp Q9NTI5 PDS5B_HUMAN
9884	EHYPVSSPSSPPAQPGGVSR220000000000000000	sp O75179 ANR17_HUMAN
9885	SAASPVVSSMPERASESSSEEKDDYEIFVK202200000000000000	sp P18583 SON_HUMAN
9886	LSSASTGKPLSVEDDFEK22000000000000000000	sp O75152 ZC11A_HUMAN
9887	TTAPTEGKSPLK20000	sp Q96T37 RBM15_HUMAN
9888	APPQTHLPSGASSGTGSASATHGGSPGTR200000000000	sp P46379 BAG6_HUMAN
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9890	EAQQKVPDEEENEESDNEKETEK2000000000	sp P11388 TOP2A_HUMAN
9891	STPLASPSPSGRSPQR200000000	sp Q8WYP5 ELYS_HUMAN
9892	GPSQATSPIR200200000	sp Q7RTP6 MICA3_HUMAN
9893	YHTPPRSR2000200	sp P30414 NKTR_HUMAN
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9895	STSPIIGSPPVR20000200000	sp Q86TB9 PATL1_HUMAN
9896	VRHSDPDPSPR200020000	sp Q9BRD0 BUD13_HUMAN
9897	RGPSPPASPTR200000200	sp Q6P5Z2 PKN3_HUMAN
9898	SDGACDSPSSDKENSSQIAQDHQK2.00000000000002E+24	sp Q9UIG0 BAZ1B_HUMAN
9899	DLESCDDDNQGSKSPK2000000002000	sp Q5T3I0 GPTC4_HUMAN
9900	SGLQLSLEQGTGENSFR200002000000002000	sp P23497 SP100_HUMAN
9901	SLSSSLDDTEVKK202000000000	sp O95292 VAPB_HUMAN
9902	KSEAKQMPR200001000	sp Q15345 LRC41_HUMAN
9903	APPSPGSLAAPGR200000000000	sp Q9Y4B5 MTCL1_HUMAN

9904	TPTPSTPGDTQPNT PAPVPPAEDGIK2. 0000000020002E+26	sp Q14839 CHD4_HUMAN
9905	RPGNSTVTSSPPVTPK200000002000	sp Q7Z2Z1 TICRR_HUMAN
9906	VSDSESEGPQKGPASDSETEASR2. 0200000000202E+21	sp Q96ST2 IWS1_HUMAN
9907	NMTVEQLLTGSPTSPTVEPEKPTR2. 000002E+22	sp Q9C0C9 UBE20_HUMAN
9908	NPSSSVPPPSAGSVK20000000000000	sp Q8WWI1 LMO7_HUMAN
9909	AGIPQHHPMAQNLQYPDDSDDEKK2000000	sp Q15223 PVRL1_HUMAN
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9911	SSSSSQESLNRPLSAK20000000000000	sp Q7Z460 CLAP1_HUMAN
9912	SLDGAEFSPASVSENHDAGPDGDKR2. 00000020002E+26	sp Q9H4G0 E4IL1_HUMAN
9913	SSISSHESFSLDSGIYR202200000000000000	sp Q9UPZ3 HPS5_HUMAN
9914	LLRPLSPVTPPPNSGSK20020000000000	sp Q9C0A6 SETD5_HUMAN
9915	TASTPTPPQTGGGLEPQANGETPQVAVIVRPPDR2. 02E+31	sp Q04637 IF4G1_HUMAN
9916	SPSPIGVGSEMCR20200000000000	sp Q9Y4B5 MTCL1_HUMAN
9917	TAANKSPCETISSPSSTLESK22000000000	sp Q9C0D5 TANC1_HUMAN
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9919	SPPITPNWR2000200000	sp Q9UQ88 CD11A_HUMAN
9920	NIISPPWDFR20000000	sp Q9UKB5 AJAP1_HUMAN
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9922	KFQEQECPPSPEPTR2000000	sp Q13042 CDC16_HUMAN
9923	FGSTGSTPPVSPTSER202200020000000	sp 000267 SPT5H_HUMAN
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9925	SDSQASLTVPLSPQVVR200000000002000000	sp Q969V6 MKL1_HUMAN
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9927	ASQTPVPPGAPSPDKDPAKESPKPEEQK200000000200000000	sp Q8TEV9 SMCR8_HUMAN
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9929	DSL GAYASQDANEQGQDLGK2E+19	sp Q15911 ZFH X3_HUMAN
9930	ETPRPEGGSPAGTPPQPK2. 0000000000002E+19	sp A7E2V4 ZSWM8_HUMAN
9931	DEEPSGWEEPSQISIR20000000	sp Q9C0C2 TB182_HUMAN
9932	TDHGAEIVYKSPVVGDTSPR200000022000	sp Q9BR39 JPH2_HUMAN
9933	KGSVDQYLLR200000000	sp Q9UPQ9 TNR6B_HUMAN
9934	GRASSHSSQTQGGGSVTK2000000000000000	sp P10636 TAU_HUMAN
9935	IDEDGENTQIEDTEPMSPVLNSK20000000	sp TAU_HUMAN
9936	STSAPQMSPGSSDNQSSSPQPAQQK200020000000000	sp Q8NCN4 RN169_HUMAN
9937	LDSQPQETSPELPR2000002000000	sp P02545 LMNA_HUMAN
9938	AGSPDVLR2000000	sp Q12888 TP53B_HUMAN
9939	LSLEGDHSTPPSAYGSVK200000000000	sp Q14157 UBP2L_HUMAN
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9942	TSDANETEDHLESLICK200000000000	sp P07355 ANXA2_HUMAN
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9945	RRSPSPTPTGPSR2000200000000	sp Q9H8M5 CNNM2_HUMAN
9946	IWDPTPSHTPAGAATPGR20000	sp Q09161 NCBP1_HUMAN
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9948	RVSLEPHQPGTPESK200000000000200	sp P46937 YAP1_HUMAN
9949	EEASLLSHSPGTSNQSQPCSPKPIR20002000000	sp P49450 CENPA_HUMAN
9950	VQTTTPPAVQGQK200000000000	sp O75533 SF3B1_HUMAN
9951	TVLGPDTAEPATAEAAVAPPDAGLPLPLPAEGSPPPK200000	sp Q86XP3 DDX42_HUMAN
9952	KLSSSEPEYEEDEFNDDQSIK2200000000000000	sp Q14980 NUMA1_HUMAN
9953	ESGVVAVSPEKSESPQKEDGLSSQLK2. 0000000002E+25	sp Q9H4Z3 PCIF1_HUMAN
		sp Q2M2I8 AAK1_HUMAN
		sp Q92667 AKAP1_HUMAN
		sp Q9H4L7 SMRCD_HUMAN
		sp Q96T58 MINT_HUMAN



9954	DHNVGSPPKSPTHASQHTQTEK200000000000000000	sp A0JNW5 UH1BL_HUMAN
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9957	SPSKPPGPNR20000000000	sp Q92766 RREB1_HUMAN
9958	SQGSPEDESAPQASPGPSK2000000000000000	sp Q9P107 GMIP_HUMAN
9959	YRSPYSGPK20020000	sp Q14498 RBM39_HUMAN
9960	GEIKDSSPSSSPK2202000	sp Q6WKZ4 RFIP1_HUMAN
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		sp P10316 1A69_HUMAN
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		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
		sp P01892 1A02_HUMAN
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9963	QPASAQSTPSTTPHSSPK2000000000000000	sp Q96A73 P33MX_HUMAN
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		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
		sp Q29960 1C16_HUMAN
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		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
9966	YGSQEHPIGDK2000000000	sp Q14644 RASA3_HUMAN
9967	EGLSGDALSSPSAPEPVVGTDDK2E+21	sp Q92503 S14L1_HUMAN
9968	GPKPEPPGSGSPAPRR2020000000	sp Q9UKJ3 GPTC8_HUMAN
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9971	SQSSHSYDDSTLPLIDR2002000000000000	sp 060716 CTND1_HUMAN
9972	IQPAGNTSPR2000	sp P49674 KC1E_HUMAN
9973	RESESEDETPPAAPQLIK2000020000000000	sp 060832 DKC1_HUMAN
9974	SMGSQEDDSGNKPSSYS20	sp Q9UNS2 CSN3_HUMAN
9975	SRSPQWR200000	sp Q9UQ35 SRRM2_HUMAN
9976	RGEGDAPFSEPGTTSTQRPSSPETATK200000000000	sp Q9UQ35 SRRM2_HUMAN
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9978	TAHNSEADLEESFNEHELEPSSPK2E+20	sp Q8IWS0 PHF6_HUMAN
9979	SPSPAHLPDDPK2020000000000	sp Q92615 LAR4B_HUMAN
9980	TSPASLDFPESQK2000000000000	sp Q96JM3 CHAP1_HUMAN
9981	HQEVQDQDPVFQGSDDSGYQSDHKK20000000000	sp Q3B726 RPA43_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
9982	EDALDDSVSSSVHASPLASSPVR200000000000020000	sp A6NKT7 RGPD3_HUMAN
		sp 014715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
9983	TYSLTTPAR200000	sp Q9Y4H2 IRS2_HUMAN
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9993	RPSESDKEDELKVK2020000000000	sp Q02952 AKA12_HUMAN
9994	NKPLSPIK20000	sp P35251 RFC1_HUMAN
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9996	SPHQLLSPSSFSPSATPSQK2202000000000	sp Q14181 DPOA2_HUMAN
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10005	LSKPPFQTNPSEMVS20000000000	sp Q6PKG0 LARP1_HUMAN
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10035	MVSTSATDEPR2000000000	sp Q8N1G0 ZN687_HUMAN
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		sp P10636 TAU_HUMAN
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		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
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		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
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		sp Q99666 RGPD5_HUMAN
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10053	TTSSANNPNLMYQDECRR20000000000000000	sp Q92841 DDX17_HUMAN
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10056	VKPPPQISPSK20000	sp Q9H6U6 BCAS3_HUMAN
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10210	ESAAPASPAPSPAPSPPTAPPQK2.000000000002E+22	sp Q9NZJ0 DTL_HUMAN
10211	ALEAASLSQHPPSLCISDSEEEEEER2.0000002E+21	sp Q96T37 RBM15_HUMAN
10212	ATDIPASASPPPVAGVPFFK2E+19	sp Q9UFC0 LRWD1_HUMAN
10213	VQMTSPSSTGSPMFK2000000200000	sp Q9GZY8 MFF_HUMAN
10214	SPSHSSNRPFTPPTSTGGSK2.02E+19	sp Q3KQU3 MA7D1_HUMAN
10215	TQTPVSPAPQPTTEER200000200000000000	sp 043159 RRP8_HUMAN
10216	LEPVYSPPGSPPGDPR30000200000000	sp Q8N1G0 ZN687_HUMAN
10217	TAVQYIESSDSEIETSELPPK200000000000	sp P49790 NU153_HUMAN
10218	SSQSPATAVPASDSPPTK220000000000000000	sp Q15648 MED1_HUMAN
10219	NPWSDDESKSESLEETEPVVIPR202020000200000000	sp Q14247 SRC8_HUMAN
10220	WDESNSERDVTK202020000000	sp 075676 KS6A4_HUMAN
10221	HHHSDSDSNSPCK2020200000	sp Q14527 HLTF_HUMAN
10222	SEDDSAKFDSEEDSASVFSFGLK20000000000000	sp Q3KQU3 MA7D1_HUMAN
10223	LNSSNSASPHR202000000	sp Q02880 TOP2B_HUMAN
10224	NEETSEESQEDEKQDTWEQQMR2.2002E+20	sp P30414 NKTR_HUMAN
10225	SQSSGSSATHPIVPGAR2000000000000000	sp Q9BU76 MMTA2_HUMAN
10226	GQSQTSPDHRSDTSSPEVR200000000022200000	sp Q02880 TOP2B_HUMAN
10227	SRSPTPPSSAGLGSNSAPPIDSR2.02E+22	sp P49792 RBP2_HUMAN
		sp P16383 GCFC2_HUMAN
		sp Q9Y4H2 IRS2_HUMAN
		sp Q9UQ35 SRRM2_HUMAN
		sp Q8IWX8 CHERP_HUMAN

10228	EIKLSDDFDSPVKGPLCK200002000000000	sp Q96T23 RSF1_HUMAN
10229	SGASEANLIVAK200000000000000	sp P46013 KI67_HUMAN
10230	SILVSPTGPSR20000000	sp Q14684 RRP1B_HUMAN
10231	QQSSSSKGDSPKPR220000200000000	sp Q5VWQ8 DAB2P_HUMAN
10232	TSAQHALTSVSRGSSLK2020020000	sp P61073 CXCR4_HUMAN
10233	REPGSGTESDTSPDFHNQENEPSQEDPEDLDGSVQGVKPK2. 02E+33	sp Q9HAF1 EAF6_HUMAN
10234	RRSPSPYYSR2000000	sp P62995 TRA2B_HUMAN
10235	QPPSEGPSNYDSYK20000000	sp Q13595 TRA2A_HUMAN
10236	SYSPDGKESPSDKK202000000000000	sp Q8TEW0 PARD3_HUMAN
10237	SQSESSDEVTELDLSHGKK2002000000000000000	sp P43243 MATR3_HUMAN
10238	TSADAQEPASPVVSPQQSPPTSPHTWR200020020000000	sp Q86YS7 C2CD5_HUMAN
10239	RPSSSEIITEGK200000000	sp Q96D71 REPS1_HUMAN
10240	DEVSHLQSGSHLANNSDPESTFR200000000	sp Q9BQF6 SENP7_HUMAN
10241	IQSSLSVNSDISK200000200	sp O15027 SC16A_HUMAN
10242	TSAQHALTSVSRGSSLK200000000000022000	sp Q9P1T7 MDFIC_HUMAN
10243	SSLAETLDSTGSLDQR20002000202000000	sp P61073 CXCR4_HUMAN
10244	TNSMGSATGPLPGTK210000000000000	sp Q9UGH3 S23A2_HUMAN
10245	STDFVFIPPEGK2000000000000	sp O15014 ZN609_HUMAN
10246	SRTPSASNDQQE2000000000	sp Q5UIP0 RIF1_HUMAN
10247	GLNPDGTPALSTLGGFSPASKPSSPR2. 0002E+20	sp O43765 SGTA_HUMAN
10248	ESPHFYR2000000	sp Q9UK58 CCNL1_HUMAN
10249	GLNPDGTPALSTLGGFSPASKPSSPR20020000	sp Q9HDC5 JPH1_HUMAN
10250	YSPTSPTYSPTPK300000200000000	sp Q9UK58 CCNL1_HUMAN
10251	SNPESIAETPPAR200000	sp P24928 RPB1_HUMAN
10252	ANEAGGQVGPEAPRPPEMSPEMR200000	sp Q9Y4H2 IRS2_HUMAN
10253	TPNNVSTPAPSPDASQLASSLSSQK20000000000000000	sp Q8N3D4 EH1L1_HUMAN
10254	NQSPTAEKPASSSLPSSPPPQLLTR20000000000	sp Q8WWI1 LM07_HUMAN
10255	IESDVQEPTPEDDLDIMLGNK2E+20	sp Q99567 NUP88_HUMAN
10256	TTLSEPLSSPAASPK20000000020002000	sp P20042 IF2B_HUMAN
10257	TAANKSPCETISSPSSTLESK2020000000000	sp Q8IVL1 NAV2_HUMAN
10258	DGTAPPPQSPGSPGTGQDEEWSDEESPR20002000	sp Q9COD5 TANC1_HUMAN
10259	TKSDESGEKNGDEDCQR2002000000000000000	sp Q9UKS6 PACN3_HUMAN
10260	SQKSPTGTPAR2020000020000	sp Q6PKG0 LARP1_HUMAN
10261	STTPPPAEPVSLPQEPK2000000000000000000	sp Q13595 TRA2A_HUMAN
10262	SPQPVSPPRQSQPPHSSPSPR2. 000002E+22	sp Q9UN86 G3BP2_HUMAN
10263	DTHEDHDTSTENTDESNHDPQFEPVSLPEQEIK2E+25	sp Q09472 EP300_HUMAN
10264	SSPASSPELPWRPR200200000000000	sp P43487 RANG_HUMAN
10265	STPLASPSPPGR2000200020000	sp Q9NYF3 FA53C_HUMAN
10266	ELSPRPR2000000	sp Q8WYP5 ELYS_HUMAN
10267	EGSVLDILKSPGFASPK200000000	sp Q99956 DUS9_HUMAN
10268	GPTPPAMGPR200000000	sp P49790 NU153_HUMAN
10269	ASVSGPNSPSETR2000000	sp Q3KQU3 MA7D1_HUMAN
10270	KFELLTPPLSPSR200020000	sp O15344 TRI18_HUMAN
10271	STLTTLPPFGAEANSPGK20000	sp P04198 MYCN_HUMAN
10272	SSIETKPDASQLPK2000000000000	sp P01106 MYC_HUMAN
10273	TRDDGDEGLLTHSEEELEHSQDTDADDGALQ2. 020000002002E+21	sp Q96GV9 CE030_HUMAN
10274	SYKVSTSGPR2000000	sp Q3KR37 GRM1B_HUMAN
10275	ELASPVPELR200000	sp Q9UKM9 RALY_HUMAN
10276	RGSQDEEDSCSLHSQTLSEDER202000000000000	sp P05787 K2C8_HUMAN
10277	QSPGSTSPKPPHTLSR20000000000	sp Q5T4S7 UBR4_HUMAN
10278	SPSPGLFR2000000	sp Q9UPZ3 HPS5_HUMAN
10279	VMHSSPSSSDGK2000000000000	sp O43159 RRP8_HUMAN
		sp Q9HAW4 CLSPN_HUMAN
		sp Q969R2 OSBP2_HUMAN

10280	LSGEHSESSTPRPR2200000	sp Q7Z3K3	POGZ_HUMAN
10281	EFGYDSPHDLSD200000200	sp Q96CW6	S7A60_HUMAN
10282	SRSSSVGSSSYPIPAVSR2.0222E+20	sp Q15149	PLEC_HUMAN
10283	LLKPGEEPSEYTDDEEDTKDHNKQD320000000000000	sp O15173	PGRC2_HUMAN
10284	NTSLPPLWSPEAER2000000	sp P49790	NU153_HUMAN
10285	YSPTSPTYSPVYTPSPK20000000000000000000	sp P24928	RPB1_HUMAN
		sp P30510	1C14_HUMAN
		sp Q29865	1C18_HUMAN
		sp P30504	1C04_HUMAN
		sp Q9TNN7	1C05_HUMAN
		sp P04222	1C03_HUMAN
10286	GGSCSQAASSNSAQGSDESLIACK2.000000000002E+22	sp Q07000	1C15_HUMAN
		sp Q29960	1C16_HUMAN
		sp Q29963	1C06_HUMAN
		sp P30505	1C08_HUMAN
		sp P30508	1C12_HUMAN
		sp Q95604	1C17_HUMAN
10287	DGSGTPSRHSLSGSSPGMK200000000002200000	sp Q9UQ35	SRRM2_HUMAN
10288	GSQAWVLS CSPSSQGQHK2000000000	sp Q6PCD5	RFWD3_HUMAN
10289	SRTPPSAPSQR2000020000000	sp Q9UQ35	SRRM2_HUMAN
10290	NEENTEPGAESSENADDPNKDTS ENADGQSDENKDDYTIPDEYR2000000000000000	sp P43243	MATR3_HUMAN
10291	SGAASMPSSPK2000	sp Q86SQ0	PHLB2_HUMAN
10292	WSPAYSFSSD SPLDSSPK200000000000	sp Q8NCG7	DGLB_HUMAN
10293	SLSRPSSLIEQEK2000000000000	sp Q12802	AKP13_HUMAN
10294	LSEEAECNPSTPSK20000	sp O94804	STK10_HUMAN
10295	SHMSGSPGPGSNTAPSTPVIGGSDKPGMEEK1.202E+30	sp A0FGR8	ESYT2_HUMAN
10296	SPLYLVSR200000000	sp Q8WYP5	ELYS_HUMAN
10297	DSSTSPGDYVLSVSENSOR2000000000000000	sp P46108	CRK_HUMAN
10298	RSPSLSSK20200000	sp Q8IYB3	SRRM1_HUMAN
10299	RTSPQVLGSILK200000000000	sp Q7Z417	NUFP2_HUMAN
10300	SPVGS GAPQAAAPAPAAHVAGNPGGDAAPAATG TAAAA SLATAAGSEDAEKK2000000	sp P16989	YBOX3_HUMAN
10301	EQGTESRSSTPLPTISSAENTR2.02E+20	sp P42166	LAP2A_HUMAN
		sp P42167	LAP2B_HUMAN
10302	QRGSETDTDSEIHESASDKDSLK2.0202E+19	sp Q9Y520	PRC2C_HUMAN
10303	TSQPPVPQGEAEEDSQGK20000	sp Q12789	TF3C1_HUMAN
10304	SLGGESSGTTVPVGSFHTEAAR2200000000000000000	sp O15085	ARHGB_HUMAN
10305	ENLQPQSSGVQGVPI SPEPLQRPEMLK2000000000000	sp Q9H467	CUED2_HUMAN
10306	LVPRPGSGSSGSSNSGSQPGSHPGSQSGSER2.0222E+27	sp O95819	M4K4_HUMAN
10307	SAPSSPTLDCEK2000200000000	sp Q9GZM8	NDEL1_HUMAN
10308	NSATCHSEDSLEID2002000000	sp Q13371	PHLP_HUMAN
10309	SSSSQESLNRPFSSK2200000000000	sp O75122	CLAP2_HUMAN
10310	FSGGFGAR20000000	sp O00571	DDX3X_HUMAN
		sp O15523	DDX3Y_HUMAN
10311	GSVSQPSTPSPKPTGIFQTSANSSFEPVK2.02E+23	sp O15027	SC16A_HUMAN
10312	TSAQHALTSVSRGSSLK20022000	sp P61073	CXCR4_HUMAN
10313	SLSMDSQVPVYSPSIDLK2000000000000000000	sp Q9ULJ3	ZBT21_HUMAN
10314	YLATSQPRPDSSGSH200200	sp Q9C0H2	TTYH3_HUMAN
10315	VDSPTVNTTLR2000000000	sp Q9H0B6	KLC2_HUMAN
10316	INPDNYGMDLNSDDSTDDEAHPR3000002000200000000	sp Q9NQS7	INCE_HUMAN
10317	EDSGTFLGK2000000	sp O75569	PRKRA_HUMAN
10318	SPTLSQVHSPLVTSPANLK2.000000000002E+20	sp Q86UU0	BCL9L_HUMAN
10319	NATLAWDSSHSSIQNSPK20002000	sp O15440	MRP5_HUMAN



10320	TLNAETPKSSPLPAK22000000	sp Q7L2J0 MEPCE_HUMAN
10321	YTPTSPSYSPSSPEYTPSPK200000000000000000	sp P24928 RPB1_HUMAN
10322	RPAEATSSPTSPERPR200020000000	sp Q9BU76 MMTA2_HUMAN
10323	ELSNSPLRENSFGSPLEFR2000002002000000	sp Q9UQ35 SRRM2_HUMAN
10324	TVSFAIEVVPAPPEPPPATPPQAK2.02E+22	sp 015047 SET1A_HUMAN
10325	DKDSGSSSPLPK2200000	sp Q86VQ1 GLC11_HUMAN
10326	SPTVPSQNPISR2000000000	sp Q03164 KMT2A_HUMAN
10327	ASPITNDGEDEFVPSDGLDKDEYTFSPGK2000000000000000	sp Q02880 TOP2B_HUMAN
10328	SATRPSPSERSSTGPEPPAPTPLLAER222000000000000000	sp Q9UQ35 SRRM2_HUMAN
10329	RKDSEEEAASPEGK200000200000	sp Q9HB96 FANCE_HUMAN
10330	LSTTPSPTSSLHEDGVEDFR22020200000000000000	sp 094842 TOX4_HUMAN
10331	RLSESLHVVDENKNESK2000000000000000	sp Q8NFC6 BD1L1_HUMAN
10332	GFGSEEGSR2000000	sp Q9Y5S9 RBM8A_HUMAN
10333	SPYAGLQLGAAK20000000000000	sp Q14134 TRI29_HUMAN
10334	DINTFVGPVEK200000	sp P46013 KI67_HUMAN
10335	RSPALHGTPTAGCGSR2000000200	sp Q8IY92 SLX4_HUMAN
10336	YSTPTYTGGQSESHVQSASEDTVTER200000000000	sp Q7Z2T5 TRM1L_HUMAN
10337	STCGDSEVEEESPGKR220000000000000000	sp 095696 BRD1_HUMAN
10338	SSITEPEGPNQNIQK2000000000000000	sp Q13625 ASPP2_HUMAN
10339	KGTTPPRSPEASPK200020002000	sp Q9HDC5 JPH1_HUMAN
10340	SKSDATASISLSSNLK200000000000	sp Q9UBF8 PI4KB_HUMAN
10341	SPSKPPGPNR200000000	sp Q92766 RREB1_HUMAN
10342	VPEPPSSHSQSGPSSGSPER20000	sp Q8N137 CNTRB_HUMAN
10343	RIACDEEFSDESEDEGEGGR202000000000	sp Q92769 HDAC2_HUMAN
10344	YPSSISSSPQK2000000	sp Q14157 UBP2L_HUMAN
10345	DGTAPPPQSPGSPGTGQDEEWSDEESPRK2E+27	sp Q9UKS6 PACN3_HUMAN
10346	TLLPWSDESPEASGPPGPR2.0000000020002E+21	sp Q9UMN6 KMT2B_HUMAN
10347	ASPDSSSCGERSPPPPPPPSDEALLHCVLEGK2.00002E+27	sp Q9BRK4 LZTS2_HUMAN
10348	SGSALLQSQSSTEDPKDEPAELKPDSEDLSSQSSASK2E+35	sp Q5VTR2 BRE1A_HUMAN
10349	STFAGSPER20000	sp Q2TBEO C19L2_HUMAN
10350	MNSQDSNTSVLEVVEKPLSER2E+20	sp Q9UJA3 MCM8_HUMAN
10351	KTSASDVTNIYPGDAGK2000000000000000	sp Q15042 RB3GP_HUMAN
10352	TPTSPLSPSSQK22002000000	sp Q5VT06 CE350_HUMAN
10353	GLQDGLRSSNSR2220000	sp P36915 GNL1_HUMAN
10354	SSPSVKPAVDPAAAK2000000000000000	sp Q6FI81 CPIN1_HUMAN
10355	EPLATLVDQSPESLK2000000	sp Q76L83 ASXL2_HUMAN
10356	TPTSSPASSPLVAK220020000000	sp Q14684 RRP1B_HUMAN
10357	ASPITNDGEDEFVPSDGLDKDEYTFSPGK20000	sp Q02880 TOP2B_HUMAN
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10359	SSSLDGEHHDGYHR2000000000000000	sp Q9C0J8 WDR33_HUMAN
10360	SRSGSSQELDVKPSASPQER2200000000000000	sp Q9UQ35 SRRM2_HUMAN
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		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
		sp Q29960 1C16_HUMAN
10361	GGSCSQAASSNSAQGSDESLIACK2000000	sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
10362	GPTTGEGALDLSVDHSPPKSPEGK2.000000000002E+21	sp 095684 FR10P_HUMAN
10363	YSPTSPTYSPVYTPSPK20030000000	sp P24928 RPB1_HUMAN

10364	HDYDDSSSEEQSAEDSYEASPGSETQR2300000000000	sp P98175 RBM10_HUMAN
10365	YTPTSPSYSPSSPEYTPK2.0000000000003E+20	sp P24928 RPB1_HUMAN
10366	ASSSGPRSPLDQR2000000	sp O14646 CHD1_HUMAN
10367	SVSASSGPCSPAAPHPPPPR2.0002E+21	sp Q86UY5 FA83A_HUMAN
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10369	VDSSTNSSPSPQQSESLSPAHTSDFR2.02E+22	sp Q9BQE9 BCL7B_HUMAN
10370	IPESETESTASAPNSPR2000002000	sp Q07889 SOS1_HUMAN
10371	SPLQSAESSPTAGK20022000000	sp Q8IWB9 TEX2_HUMAN
10372	ENFSPER20000	sp Q7Z6E9 RBBP6_HUMAN
10373	VGDSTPVSEKPVSAAVDANASESP2E+21	sp Q9H8Y8 GORS2_HUMAN
10374	AELKLSR200	sp Q92625 ANS1A_HUMAN
10375	SNCLGTDEDSQDSSDGIPSPR2.000000000002E+22	sp Q92993 KAT5_HUMAN
10376	ASASRPQPAPADGADPPPAEPEARPDGEGSPGK20000	sp P11274 BCR_HUMAN
10377	RPAEATSSPTSPERPR200002000000	sp Q9BU76 MMTA2_HUMAN
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10379	LFGQEELDPDSLSDASGSDGGRGPEPGVEPQDSR2.0200202E+24	sp Q9Y4F5 C170B_HUMAN
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10382	GGTSHPSVPVPPASSPSPLSGSALCGGK2200000000000000	sp P49848 TAF6_HUMAN
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10384	YQQITPVNQRS200	sp Q9NYL2 MLTK_HUMAN
10385	SWSYNGYSDLSTAR23000020000000	sp P30414 NKTR_HUMAN
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10387	LSPLTGQACR2000000000	sp Q9UKA4 AKA11_HUMAN
10388	SEESDTSERQDDSYIEPEPVEPLK2.00002022E+26	sp Q9BZF1 OSBL8_HUMAN
10389	NTVDLVTCHIHSDDDEIDFK20200000000	sp Q5H9R7 PP6R3_HUMAN
10390	NSNFSVQHPSSTSPTEK200000000	sp Q03468 ERCC6_HUMAN
10391	TSPSGKATLK200000000000	sp P27695 APEX1_HUMAN
10392	KASSPSPLTIGTPESQR202000000000000	sp Q9NPI6 DCP1A_HUMAN
10393	SRSPLSPR202002000	sp Q9NXX6 NSE4A_HUMAN
10394	GPTSSPCEEGDEGEEDRTSDLR2E+20	sp Q86VM9 ZCH18_HUMAN
10395	SESPPLSDPK2000000000	sp Q15059 BRD3_HUMAN
10396	SSSSQESLNRPFSSK220000000000000	sp O75122 CLAP2_HUMAN
10397	KCVTMSSAVVQLYAADR10200000000000	sp O00401 WASL_HUMAN
10398	SLSPGVSRSSTSYTETK2020002000000000000	sp Q9UKV3 ACINU_HUMAN
10399	SYLEGSSDNQLK2000002000000	sp Q9UHI6 DDX20_HUMAN
10400	SGYSSPGSPGTPGSR200200000000	sp TAU_HUMAN
10401	KAEAAAAPTVPAPGPAQPGHVSPTPATTSPGK200000	sp P10636 TAU_HUMAN
10402	STAQQELDGKASPTPVIVASHTANK2E+25	sp Q9Y6R0 NUMBL_HUMAN
10403	SFSISPSRR2020200000	sp P35606 COPB2_HUMAN
10404	VGDTEKPEPERSPPNR20000000000000	sp P18583 SON_HUMAN
10405	MSDSESEELPKPVSDSESEPPR2.0202000000002E+23	sp Q9NZ63 CI078_HUMAN
10406	DSHSSEDEASSQTDLSQTISKK2.02E+22	sp Q96ST2 IWS1_HUMAN
10407	RESGGRPIKPPK20000000000	sp Q5JTV8 TOIP1_HUMAN
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10409	DEVVSPLPSALQGPSGLSAPPAASVISAPSSSSR2000000000000	sp Q9UEY8 ADDG_HUMAN
10410	SPVVSAGDTSR200020000000	sp Q7L2J0 MEPCE_HUMAN
10411	HKSDSPESDAER20000200000	sp TAU_HUMAN
10412	ELLSPLSEPDDRYPLIVK2002000000000000	sp P10636 TAU_HUMAN
10413	RASSPFRR2200000	sp O75400 PR40A_HUMAN
10414	DHNSSEDEDEDKYADDIDMPGQNFDSK2.0000000000001E+23	sp Q9UHB7 AFF4_HUMAN
10415	HIKEEPLSEEPCTSTAIASPEKK200000200000	sp Q14978 NOLC1_HUMAN
		sp O95391 SLU7_HUMAN
		sp Q9Y2X3 NOP58_HUMAN

10416	LMQLMESEQK20000	sp Q86UP2 KTN1_HUMAN
10417	SSKSFSLDEPPLFIPDNIATIR2.02E+19	sp Q92576 PHF3_HUMAN
		sp P01891 1A68_HUMAN
		sp P10316 1A69_HUMAN
10418	KGGSYSQAASSDSAQGSVDVSLTACK20002000000000	sp P05534 1A24_HUMAN
		sp P30447 1A23_HUMAN
		sp Q09160 1A80_HUMAN
		sp P01892 1A02_HUMAN
10419	RGSLEMSSDGEPLSR200002000000000	sp Q6ZN18 AEBP2_HUMAN
10420	TVTPASSAKTSPAK200020000	sp Q16555 DPYL2_HUMAN
10421	RPESPSEISPIKGSVR200002000000000	sp Q7Z5K2 WAPL_HUMAN
10422	APVSGSRSPDR2020000	sp Q8N5F7 NKAP_HUMAN
10423	SDSDSSTLPR20000000000	sp Q9ULE0 WWC3_HUMAN
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10425	IQPQPPDEDGDHSDKEDEQPQVVVLK200000000000000	sp Q96AT1 K1143_HUMAN
10426	HNGSLSPGLEAR20000000	sp Q9C0C2 TB182_HUMAN
10427	LIDSSGSASVLTHSSSGNSLK2000000	sp Q96SU4 OSBL9_HUMAN
10428	YGGSHYSSSGYSNSR20000000	sp Q9BRL6 SRSF8_HUMAN
10429	LSPQQAFPR200000000	sp Q9BUH8 BEGIN_HUMAN
10430	AYSPTCSPTLGFK200000200000	sp Q5TGY3 AHDC1_HUMAN
10431	RHSSVSDSQCEPPSVGTEYSQGASPPQHQLK2.02E+30	sp P39880 CUX1_HUMAN
10432	TPSPKEEDEEPESPPEKK2000000	sp Q9H1E3 NUCKS_HUMAN
10433	WTKPSSFSDSER220000000	sp Q8NC56 LEMD2_HUMAN
10434	RSQSSSQGDNPLAPGYLPPHYK2.0022E+21	sp Q6ZRV2 FA83H_HUMAN
10435	KGAANASGSSPDAPAK20000000	sp O75962 TRIO_HUMAN
10436	IAVHHNSVEDVPEEK2000000000	sp Q9UKY1 ZHX1_HUMAN
10437	RNSSDSAIDNPKPNK2020000000000	sp P46100 ATRX_HUMAN
10438	SMGTGDTPGLEVPSSPLRK1000000000000200000	sp Q86WB0 NIPA_HUMAN
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10440	HSSISPVRPLNSSLGAELSR2.002E+20	sp Q9NYV4 CDK12_HUMAN
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10442	EPDATPPSPR2002000	sp P28715 ERCC5_HUMAN
10443	SSGHSSSELSPDAVEK20000000000	sp Q9UQ35 SRRM2_HUMAN
10444	IQALGLHLPDPPELESEDEDEEGATALNNHSSIPMDPEHVELVK200000000000000	sp Q16626 MEA1_HUMAN
10445	STSHTSDFNPNSGSDQR200000000000000000	sp Q99700 ATX2_HUMAN
10446	SSSPGKPAVSSLNSSHSR22000000000000000000	sp Q9UHB7 AFF4_HUMAN
10447	RQSLPASPR20002000	sp Q3KQU3 MA7D1_HUMAN
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10453	REPGSGTESDTSPDFHNQENEPSQEDPEDLDGSVQGVKPK2.200000000002E+31	sp Q9HAF1 EAF6_HUMAN
10454	YTGEEDGAGGHSPAPPQTEECLR2E+22	sp O94762 RECQ5_HUMAN
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10456	EAGELPTSPLHLLSPGTTPR20000000002000	sp Q969R5 LMBL2_HUMAN
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10458	SPSPAPPPRR20200000000	sp Q8IYB3 SRRM1_HUMAN
10459	KSRTLAQTV200	sp Q9H1N7 S35B3_HUMAN
10460	KPSSETDIENWASK2000000000	sp Q9C0H5 RHG39_HUMAN
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10463	ASPLYASYK200000000	sp Q9BUH8 BEGIN_HUMAN
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10477	KPTSPVKPIQNTTEYR2000000000000000	sp P25054 APC_HUMAN
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10494	TPHYGSQTPLHDGSR20200000000	sp O00267 SPT5H_HUMAN
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10526	HSPPMVR2000000	sp Q060256 KPRB_HUMAN
10527	SRSGSSQELDVKPSASPQER2020000000000000000	sp Q9UQ35 SRRM2_HUMAN
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10529	NAIASDSEADSDTEVPK20200000200000	sp Q8WVC0 LEO1_HUMAN
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10545	MFVSSSGLPPSPVPSPR200000000002000	sp Q7Z309 F122B_HUMAN
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10565	TTSISPALAR2000000	sp Q9UKE5 TNIK_HUMAN
10566	MKSLEQDALR200000000	sp Q14160 SCRIB_HUMAN
10567	MLPHAPGVQMQAIPEDAVHEDSGDEDGEDPDK200000000000	sp Q92769 HDAC2_HUMAN

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10579	SCVEEPEPEPEAAEGDGDKK2E+20	sp Q9UH99 SUN2_HUMAN
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		sp Q2M2I8 AAK1_HUMAN
		sp P35659 DEK_HUMAN

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10672	SRSYSR20200000	sp Q13595 TRA2A_HUMAN
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		sp Q95604 1C17_HUMAN
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		sp Q9BTX1 NDC1_HUMAN
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10691	SRSPVLR2000000	sp Q13523 PRP4B_HUMAN
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10695	GNKSYLEGSSDNQLK2000022000000	sp Q9UHI6 DDX20_HUMAN
10696	EIPSATQSPISK2000200000	sp Q9BQG0 MBB1A_HUMAN
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10698	AATSPGVGQSAPGTR20000000000000	sp Q6IE81 JADE1_HUMAN
10699	LSQSSQDSSPVR2000020000	sp O95251 KAT7_HUMAN
10700	LPSDTDDSDASEDPGGAER2000020020000000000	sp Q9UJK0 TSR3_HUMAN
10701	SEPPAPVR200000000	sp P35398 RORA_HUMAN
10702	VKSPQRPSDWSK20000000000	sp Q5UIP0 RIF1_HUMAN
10703	IERPGEGLMVDNPMR2000000000	sp Q96T88 UHRF1_HUMAN
10704	RPESPPSILTPPVPTADK2000000000000000000	sp P55198 AF17_HUMAN
10705	RPASPSPEHLPATPAESPAQR20000000000000000	sp Q9H7L9 SDS3_HUMAN



10706	TLENSPISSCDTSDTEGPVPVNSAAVLK2. 0022E+24	sp Q8N556 AFAP1_HUMAN
10707	LPALSHSEGEDEDEDEDEDEGKGSSEK2. 02E+23	sp P55201 BRPF1_HUMAN
10708	NSPGVPTGAK2000000000	sp Q08379 GOGA2_HUMAN
10709	RRPSPQSPR20002000	sp Q9UQ35 SRRM2_HUMAN
10710	QSVTSFPDADAFHHQVHDDLLSSSEEECKDR2220000000	sp P50402 EMD_HUMAN
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10712	GQSQTSPDHRSDTSSPEVR200002022000000	sp Q9UQ35 SRRM2_HUMAN
10713	GRPGSGTSGVDSLNFIESVK2000000000000000	sp Q8IXT5 RB12B_HUMAN
10714	RPPSQSLTSSSLSPDIFDPSPEEDKEEHISLAHR2000000000000000	sp Q96B97 SH3K1_HUMAN
10715	DPQTPVLQTK20000000	sp Q9ULW0 TPX2_HUMAN
10716	LAALKDEPQTPDVPVSPFGESPLSPIDMDTQER2000000020001000000	sp P17535 JUND_HUMAN
10717	METVSNASSSSNPSSPGR1000000000200200000	sp Q9NTI5 PDS5B_HUMAN
10718	NQTYSFSPSK20000	sp Q86YS7 C2CD5_HUMAN
10719	LEDVKNSTPFK200000	sp P55327 TPD52_HUMAN
10720	EWYQSTIPQSPSPAPDDPEEGR2000000000000000000	sp Q08499 PDE4D_HUMAN
10721	GPSPEGSSSTESSPEHPPK200020000220000000	sp Q9UQ35 SRRM2_HUMAN
10722	SNSQSDSHDEEVSTPPNPVVK2020000020000000000	sp P31321 KAP1_HUMAN
10723	TLTDEVNSPDSR2000000	sp Q8TAQ2 SMRC2_HUMAN
10724	YSPTSPTYSPK20000030002000	sp P24928 RPB1_HUMAN
10725	GFINDDDEDEGEDEGSDSGSEDDVGHK20200200000000	sp Q7KZ85 SPT6H_HUMAN
10726	HNSESEVPSSMFILEDDR2000000000000000	sp Q6R327 RICTR_HUMAN
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10728	AESPSPAPPGLR2000000000000	sp Q9BUA3 CK084_HUMAN
10729	LTTEATQTPVSNK2000000	sp Q9BXS6 NUSAP_HUMAN
10730	TVAISDAAQLPHDYCTTPGGTLFSTTPGGTR3000000000200000000	sp Q13542 4EBP2_HUMAN
10731	SRSQRSNSPLPVPPSK2020202020000000000	sp Q13247 SRSF6_HUMAN
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10733	VSSPSPELGTTPASIFR2002000000000000000	sp Q99501 GA2L1_HUMAN
10734	RPDSPTRPTLR2000000000	sp Q7RTP6 MICA3_HUMAN
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10736	SPNPSPERPR20002000000	sp Q96ST8 CEP89_HUMAN
10737	VTRSPSPVQEEHSDPEMTEEEK2. 020200000002E+22	sp Q8TF01 PNISR_HUMAN
10738	TPSSEEISPTK2000000000	sp Q15311 RBP1_HUMAN
10739	AGLSPERPPKTPGSPR200000020000000	sp Q9H6F5 CCD86_HUMAN
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10741	RLSSTSLASGHSVR202000002000	sp Q9BZL6 KPCD2_HUMAN
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10746	EAYSGCSGPVDECPSPVHK2000000	sp Q8N556 AFAP1_HUMAN
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10749	REVLVYSEGLSGEER2000200000	sp Q9H7N4 SFR19_HUMAN
10750	MSQPGSPSK202000	sp Q14839 CHD4_HUMAN
10751	QGSITSPQANEQSVTPQRR2000000000200000	sp Q9UQ35 SRRM2_HUMAN
10752	VDGPRSPSYGR20000	sp Q07955 SRSF1_HUMAN
10753	ETKAPPPASPAR2000000000200	sp Q5T8A7 PPR26_HUMAN
10754	EVAENQQNQSSDPEEEKGSQPPAAESQSSLR2E+23	sp Q43491 E41L2_HUMAN
10755	SRSPGRPPK20000000	sp Q14739 LBR_HUMAN
10756	EYSPPYAPSHQQYPPSHATPLPK2000000000000000	sp Q9UPT8 ZC3H4_HUMAN
10757	GLNPDGTPALSTLGGFSPASKPSSPR20020000000	sp Q9UK58 CCNL1_HUMAN

10758	SASSESEAEENLEAQPQSTVRPEEIPPIPENR2. 022E+31	sp Q13427 PPIG_HUMAN
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10760	RRSPSPPTTR2000000	sp Q8IYB3 SRRM1_HUMAN
10761	VSMPDVELNLKSPK2000	sp Q09666 AHNK_HUMAN
10762	SQQQPTPVTPK2000	sp 095251 KAT7_HUMAN
10763	KGQQSGTVSPQKQSAWK2000020000	sp A6QL64 AN36A_HUMAN
10764	AGAASPLNSPLSSAVPSLGK2000000000000000	sp 075962 TRIO_HUMAN
10765	YGPADVEDTTGSGATDSKDDDDIDLFGSDDEESEEAKR2. 2E+31	sp P24534 EF1B_HUMAN
10766	LSPEVQSVAESK200000000000	sp Q5T5U3 RHG21_HUMAN
10767	KHSSDDYYGDISSLESSQK2000000000000000	sp Q9UGU5 HMGX4_HUMAN
10768	NSVQSGESDSDEEESKEPKLKP2. 00202E+21	sp 075592 MYCB2_HUMAN
10769	SSGHSSSELSPDAVEK2000220020000000	sp Q9UQ35 SRRM2_HUMAN
10770	QFTSSSSIK22000	sp Q04695 K1C17_HUMAN
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10774	ERSPSPLRGNVPSPLPTR202000000002000000	sp Q9Y2V2 CHSP1_HUMAN
10775	DYTGCTSESLSPVK200020000	sp 095297 MPZL1_HUMAN
10776	ETSVSKEDTDHEEKASNEVDTK2. 202E+21	sp 075475 PSIP1_HUMAN
10777	SPQNQYPAELMR2000000000100	sp P33993 MCM7_HUMAN
10778	HLDGEEDGSSDQSQASGTTGGR20020000000000	sp Q9UNF1 MAGD2_HUMAN
10779	SPVVSVDTSR200000002000	sp TAU_HUMAN
		sp P10636 TAU_HUMAN
10780	SRGSSPGLR20202200000	sp Q9UQ35 SRRM2_HUMAN
10781	TTPSATSLPQTVMVMTSPVLTSTTK2000000000000	sp P18846 ATF1_HUMAN
10782	EILSHLQGDCTSPGR20000	sp Q9UQ49 NEUR3_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
10783	KGGSYSQAASSDSAQGSVDVSLTACK20002000000000	sp P05534 1A24_HUMAN
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		sp P10316 1A69_HUMAN
		sp P30447 1A23_HUMAN
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10786	LPASPSGSEDLSSVSSSPTSSPK22000	sp Q9P2N6 KANL3_HUMAN
10787	SPSSEWLVR20000000	sp Q9NWQ4 GPT2L_HUMAN
10788	RREEGPPPPSPDGASSDAEPEPPSGR2200000000000	sp Q9NTJ3 SMC4_HUMAN
10789	SEAPAEVTHFSPK2000	sp Q6PID6 TTC33_HUMAN
10790	WDCESICSTYSNLYNHPQLIK2200000000000000	sp Q96GA3 LTV1_HUMAN
10791	LSEQLSFR2000000	sp Q96PK6 RBM14_HUMAN
10792	TEPSSPLSDPSDIIR200000000000	sp Q9ULJ3 ZBT21_HUMAN
10793	RPLSGPDVGTQPAGLASGAK2000000000000	sp 096013 PAK4_HUMAN
10794	EDSGLFSPIR20000	sp Q9UKA4 AKA11_HUMAN
10795	SFVPEEEKHEER2000000000000	sp 094880 PHF14_HUMAN
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10798	STSAPQMSPGSSDNQSSSPQPAQK200022000000000	sp Q14157 UBP2L_HUMAN
10799	TRESHRSSSPTAGPSTEGAEGPEEK2. 0020222E+24	sp Q8NHM5 KDM2B_HUMAN
10800	HLSESSGKPLSTK200200000000	sp Q8IVT2 MISP_HUMAN
10801	KPPAPSPVQSQSPSTNWSPAVPVKK2. 000000022002E+20	sp Q8IYB3 SRRM1_HUMAN
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10803	RSSGISPYFSSR2200	sp P10070 GLI2_HUMAN
10804	SATSTSASPTLR200000	sp 075061 AUXI_HUMAN

10805	KGSSTDISEDWEK2000000000	sp Q9NW68	BSDC1_HUMAN
10806	HSHKPHPVETSQPSDK2000000	sp Q03188	CENPC_HUMAN
10807	SPSPSPTPEAK202000200000	sp Q8TAQ2	SMRC2_HUMAN
10808	GLPSPYNMSSAPGSR3010000200	sp P15924	DESP_HUMAN
10809	NRSPSDSDMEDYSPPPSLSEVAR2.0202010002E+21	sp Q6KC79	NIPBL_HUMAN
10810	WGQPPSPTPVPRPPDADPNTSPKPLEGRPER2.00000000000002E+25	sp Q14839	CHD4_HUMAN
10811	SNSPLPSIQLQPQSPSASKK2000000000000200000	sp Q08174	PCDH1_HUMAN
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10813	RFSHLNK200000	sp Q15928	ZN141_HUMAN
10814	AATAARPPAPPAPQPPSPTSPPRPTLAR20200000000000	sp Q96B36	AKTS1_HUMAN
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10822	SEVQQPVHPKPLSPDSRASSLESSPDK20022020220000	sp Q9UHB6	LIMA1_HUMAN
10823	GSRPPLILQSQSLPCSSPR20000	sp Q6KC79	NIPBL_HUMAN
10824	TDTGIVTVEQSPSSSK200	sp Q8NFC6	BD1L1_HUMAN
10825	FVPAENDSILMNPADQGEVQLSQNDK2000000	sp Q12888	TP53B_HUMAN
10826	SVSPESNDISSEELNHFKPIVCSPTPPK2.002002E+27	sp Q8NCN4	RN169_HUMAN
10827	SVASSQPAKPTK2200000000	sp Q07157	ZO1_HUMAN
10828	RNSDRSEDGYHSDGDYGEHDYR2.002000002E+20	sp P52756	RBM5_HUMAN
10829	TASRPDDIPDPSPPK20000000000022000	sp Q5VZK9	LR16A_HUMAN
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10832	GGHSSVSTESESSSFHSS200000	sp P61073	CXCR4_HUMAN
10833	SPSPSSPAAVNHSSSDISPVSNSTSSSPGK200000	sp Q075381	PEX14_HUMAN
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10835	SPFEHSVEHK200000	sp Q014646	CHD1_HUMAN
10836	SPHQLLSPSSFSPSATPSQK2200000000000	sp Q14181	DPOA2_HUMAN
10837	EQLSETEGSGEDEPGNDPSETTQKK2.00002E+22	sp Q13107	UBP4_HUMAN
10838	NSPLDCGSASPDK2000000	sp Q075179	ANR17_HUMAN
10839	SAGGSSPEGGEDSDREDGNYCPPVK2.00000002E+21	sp Q9H6Z4	RANB3_HUMAN
10840	STPACSPILR2000200000	sp Q3KR37	GRM1B_HUMAN
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10846	ERGSQSSDSSSLSSHR2000000000000	sp Q075376	NCOR1_HUMAN
10847	SPRPASPASNVLVPSGSTVYVK2002000000000000000	sp Q7Z589	EMSY_HUMAN
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10849	GNDPLTSSPGRSSR2020002200	sp P49736	MCM2_HUMAN
10850	GSGGSSGDELREDDEPVK2000200000000000000	sp Q7Z4V5	HDGR2_HUMAN
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		sp Q7Z3J3 RGPD4_HUMAN
		sp A6NKT7 RGPD3_HUMAN
		sp O14715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
		sp P0DJD0 RGPD1_HUMAN
		sp P0DJD1 RGPD2_HUMAN
		sp P54278 PMS2_HUMAN
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10869	SLSRTPSPPPFR2020002000000	sp Q7L4I2 RSRC2_HUMAN
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10888	SRVSVSPGRTSGK20200020000	sp Q8IYB3 SRRM1_HUMAN
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10910	QSPSPSEVSQSPGR2000000	sp P55201 BRPF1_HUMAN
10911	DTFEHDPSESIDEFNKSSATSGDIWPGLSAYDNSPR2.000002E+35	sp Q9NYF8 BCLF1_HUMAN
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10925	SSDSWEVWGSASTNR2000000000000000	sp P27816 MAP4_HUMAN
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10934	SRSRTPPVTR202000000	sp Q53F19 CQ085_HUMAN
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10939	NCPPSPLPLISK200000000	sp Q9B XK5 B2L13_HUMAN
10940	DVDDGSGSPHSPHQLSSK20200000000000	sp Q8IWC1 MA7D3_HUMAN
10941	DVDDGSGSPHSPHQLSSK200200002000	sp Q9H4A3 WNK1_HUMAN
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10943	ATTPPNQGRPDSPVYANLQELK2.2E+21	sp Q96G74 OTUD5_HUMAN
10944	TAPSSPLTSPSDTR220000000000	sp Q8IWW6 RHG12_HUMAN
10945	ESSANNSVSPSESLR2000002000	sp Q8IWB9 TEX2_HUMAN
10946	QNSPAGNKENIKNETSPSFSK2.00000000000002E+20	sp Q04726 TLE3_HUMAN
10947	SESPKEPEQLRK2020000000000	sp Q99700 ATX2_HUMAN
10948	LSSSSPGATVSPSSSDAEYDKLPDADLAR2E+19	sp P09651 ROA1_HUMAN
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10950	EDKSPSEESAPTTSPESVSGVPSSGSSGR200200000000000000	sp Q5QP82 DCA10_HUMAN
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10952	HSPTSEPTPPGDALPPVSSPHTHR2E+23	sp P78337 PITX1_HUMAN
10953	YSPTSPTYSPPTPK200000000000	sp Q9H6S3 ES8L2_HUMAN
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10956	YMAENPTAGVVQEEEEEDNLEYSDGNPIAPTK1.00002E+31	sp Q86XP3 DDX42_HUMAN
10957	RGSQNSSEHRPPASSTSEDEVKASPSSANK20000020000000	sp O15014 ZN609_HUMAN
10958	DYTGCSSTSESLSPVK2020000	sp O95297 MPZL1_HUMAN
10959	NPEAALSPTFR200000	sp Q86U44 MTA70_HUMAN
10960	SGAQASSTPLSPTR200	sp P02545 LMNA_HUMAN
10961	SASPYPHSLSSPQR20000000020000	sp Q5T200 ZC3HD_HUMAN
10962	RPMSDPSWNR20000000	sp Q14C86 GAPD1_HUMAN
10963	GFNGCDSPEPDGDSLEQSPILLEDK2.000000020002E+19	sp Q14814 MEF2D_HUMAN
10964	APTKPPR200000	sp Q9BRG2 SH23A_HUMAN
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		sp POCL81 GG12G_HUMAN
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		sp Q13068 GAGE4_HUMAN
		sp O76087 GAGE7_HUMAN
10966	GGSDSSKDPIDVNYEK200000000000	sp P09874 PARP1_HUMAN
10967	SPGSTPTTPTSSQAPQK200220000000000000	sp P35658 NU214_HUMAN
10968	SPQSDPADTPTNTK2002000000000000	sp P46013 KI67_HUMAN
10969	TLTDEVNSPDSRRR20000	sp Q8TAQ2 SMRC2_HUMAN
10970	VEMYSGSDDDDFNKLPK132000000000000000	sp O60841 IF2P_HUMAN
10971	YSPTSPTYSPTSPVYTPSPK2.000003E+20	sp P24928 RPB1_HUMAN
10972	DSPAFDPSWR20000020000	sp Q9NQS7 INCE_HUMAN
10973	EVDPSTGELQSLQMPSESGPSSLDPSQEGPTGLK20000000000000000000	sp Q13501 SQSTM_HUMAN
10974	VAEPGAEATSSSTGEESGSEHPPAVPMHNK2.00000002E+21	sp Q14684 RRP1B_HUMAN
10975	KCSLPAEEDSVLEK20000000000000	sp P27816 MAP4_HUMAN
10976	RGSLEMSSDGEPLSR20012200000000	sp Q6ZN18 AEBP2_HUMAN
10977	SRSPSPAPEKK2020000000	sp Q8IYB3 SRRM1_HUMAN
10978	SPQNDHSDTDSEENRDNQFLTTVK2.0202E+19	sp Q86W56 PARG_HUMAN
10979	RVSSRSVSGSPEPAK2200202020000000	sp Q8IYB3 SRRM1_HUMAN
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10981	SLGVLPTLNSGSPEK2000000000020000	sp P42695 CNDD3_HUMAN
10982	SLDGAEFSRPASVSENHDAGPDGDKR2.0000002E+26	sp Q9H4G0 E41L1_HUMAN
10983	KPASPLPATQQEKPSQTPEAGR2E+20	sp Q6ZU35 K1211_HUMAN
10984	SPALHGTPTAGCGSR200000000000200	sp Q8IY92 SLX4_HUMAN
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10987	KRPSLPSSPSPGLPK20020020000000	sp O43294 TGFI1_HUMAN
10988	KRPQSPSPR202000	sp Q99590 SCAFB_HUMAN
10989	KLSSGETVTYQK200000000000	sp Q70EL2 UBP45_HUMAN
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		sp A8CG34 P121C_HUMAN
10991	SQTPERPAK2020000000	sp Q96HA1 P121A_HUMAN
		sp A6NF01 P121B_HUMAN
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10994	FSGWYDADLSPAGHEEAK3000000000000000	sp P18669 PGAM1_HUMAN
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10998	HLDVDLDRQSLSSIDKNPSPR2020000002000	sp Q9Y6D5 BIG2_HUMAN
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11002	QSHSGSISPYPK2020000000	sp Q9UQ35 SRRM2_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
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		sp PODJD0 RGPD1_HUMAN
		sp PODJD1 RGPD2_HUMAN
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11005	SLPSSPELPR20002000000	sp Q15569 TESK1_HUMAN
11006	SRTHSTSSSLGSGESPF SR22000000000000	sp Q9UGV2 NDRG3_HUMAN
11007	LQPERGSPK2000	sp Q9UKV3 ACINU_HUMAN
11008	MSPTPPLFSLPEAR2000000000000000	sp Q96FV9 THOC1_HUMAN
11009	SPAPGAPTR20000000000	sp Q9BQG0 MBB1A_HUMAN
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11012	RPDHSGGSPSPTSEPAR20000000000	sp Q8TAD8 SNIP1_HUMAN
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11014	SASPYPHSLSSPQR20000000200000	sp Q5T200 ZC3HD_HUMAN
11015	PGPTPSGTNVGSSGRSPSK220000000	sp P60468 SC61B_HUMAN
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		sp Q14151 SAFB2_HUMAN
11017	RSPALHGTPTAGCGSR20000000000000200	sp Q8IY92 SLX4_HUMAN
11018	SKSPPKVIQDDSLPAGPPPQIR. 00000000000002E+25	sp Q7Z422 SZRD1_HUMAN
11019	DYSTLTSVSSHDSR200000	sp P16144 ITB4_HUMAN
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11022	GWSQEGPVKSPAECR2000000	sp Q9C0C2 TB182_HUMAN
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11027	SPILEEK20000000	sp Q12830 BPTF_HUMAN
		sp P04198 MYCN_HUMAN
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		sp O60245 PCDH7_HUMAN
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		sp P49792 RBP2_HUMAN
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11050	LSPVPVPR20000000	sp Q9UKM9 RALY_HUMAN
11051	AGAGPGGPPQKPAPSSQR2000	sp Q9Y5J1 UTP18_HUMAN
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11058	GPPSPPAPVMHSPSRK20000010200000	sp Q13573 SNW1_HUMAN
11059	RKSLDSESDSK200000200000	sp Q13185 CBX3_HUMAN
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11094	DLAVPAALTPR2000	sp Q8TEM1 PO210_HUMAN



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11137	ALQTSPQPPEGPGLPR200000000000	sp Q6PFW1 VIP1_HUMAN
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11176	RDSFDDRGPVLDYDHGSR2E+20	sp P43243 MATR3_HUMAN
11177	SPFNSPSPQDSR2020000000	sp P08651 NFIC_HUMAN
11178	DSGSDTASAIIPSTTPSVDSDESUVK2E+22	sp Q6WKZ4 RFIP1_HUMAN
11179	SPPSPPTH200020000000	sp Q96HA1 P121A_HUMAN
11180	GSPSPASVSSGR202000000000	sp P10398 ARAF_HUMAN
11181	NSDSIVSLPQSDRSSSSQESLNRPLSAK2222200000000000	sp Q7Z460 CLAP1_HUMAN
11182	SPQRSPLQSAESSPTAGK2000200000002020000	sp Q8IWB9 TEX2_HUMAN
11183	GSETGSETHESDLAPSDK200200000000	sp P48634 PRC2A_HUMAN
11184	RHSASNLHALAHPAPSPGSCSPK2.00000000002E+19	sp P47974 TISD_HUMAN
11185	SPQNQYPAELMR20000000000000	sp P33993 MCM7_HUMAN
11186	DHAPSIPTSPFGDSLESSTELR200000000000000000	sp Q9Y4B5 MTCL1_HUMAN

11187	RLSPSASPPR2020000	sp	Q8IYB3	SRRM1_HUMAN
11188	KLSSERPSSDGEVVENGITTENGK2. 20002E+23	sp	Q8N556	AFAP1_HUMAN
11189	VPEKPPTPKESPFFYR20002000000	sp	Q9HDC5	JPH1_HUMAN
11190	EEATYGVWAERDSDDERPSFGGK3E+19	sp	Q9UBB9	TFP11_HUMAN
11191	RQSSGSATNVASTPDNR220000000000000	sp	Q7Z460	CLAP1_HUMAN
11192	AAAAGSAPSQAGPGSAPSSAASGASSPAPASGPAPSAVSSSSPTPPVR200002 000000000000	sp	Q96C12	ARMC5_HUMAN
11193	VPGGSPLGVSLVYSDK2000000000000	sp	Q96RK0	CIC_HUMAN
11194	MIDQVFGVLESPELSR200	sp	Q9Y4F5	C170B_HUMAN
11195	KPGSPGSESPK200000000	sp	Q96JM3	CHAP1_HUMAN
11196	LQLQLDLSSR200	sp	P49454	CENPF_HUMAN
11197	RSSLSPSSAYER200000200000	sp	Q9BYW2	SETD2_HUMAN
11198	TSVPSPEQPQPYR2002000000000	sp	Q8TEH3	DEN1A_HUMAN
11199	FAAHISPAQSPKPPPAASPGVR2000020000000200000	sp	P19484	TFEB_HUMAN
11200	SKSPPKSPEEEGAVSS20200000000000200	sp	Q01130	SRSF2_HUMAN
11201	VNQSALAVTPSPSFQQR20000000	sp	Q9UKG1	DP13A_HUMAN
11202	SEAGHASSPDSEVTSLCQK2000000000	sp	Q6NZY4	ZCHC8_HUMAN
11203	EALNIGDITSTVSTVPPVDDTLQSASSHSPTQR2000000	sp	P50570	DYN2_HUMAN
11204	NHKPSRSFSR2020200	sp	P84103	SRSF3_HUMAN
11205	TLSSPSNRPSGETSVPPPPAVGR20000000000	sp	Q8WWM7	ATX2L_HUMAN
11206	AATKSPQPTPGR200000000	sp	094761	RECQ4_HUMAN
11207	VSSPSCDSQGLHAEAPSSEPQSPVAQTSGWPACSTPLPLVSR2. 0002E+25	sp	Q96RE7	NACC1_HUMAN
11208	SRTSPVTR202200000	sp	Q9UQ35	SRRM2_HUMAN
11209	NHSGSRTPPVALNSSR202020000000200	sp	Q9UQ35	SRRM2_HUMAN
11210	SCTPSPDQISHRASLEDAPVDDLTR2. 0202000000002E+25	sp	Q7Z2W4	ZCHV_HUMAN
11211	LVKPSSSSGSR2000000	sp	Q9H6B4	CLMP_HUMAN
11212	TASISSPSEGTPTVGSYGCTPQSLPK20200200020000000	sp	Q6PKG0	LARP1_HUMAN
11213	DGTAPPPQSPGSPGTGQDEEWSDEESPRK2000000200020000	sp	Q9UKS6	PACN3_HUMAN
11214	GWSQEGPVK20000000	sp	Q9COC2	TB182_HUMAN
11215	SRPTSFADELAAR200000000000000	sp	Q641Q2	FA21A_HUMAN
		sp	Q9Y4E1	FA21C_HUMAN
		sp	Q5SNT6	FA21B_HUMAN
11216	RYSGSDSDSISEGKRPK2020202000000000	sp	014646	CHD1_HUMAN
11217	EGQPSPADEKGNDSGEGESDDPEK2E+21	sp	075351	VPS4B_HUMAN
11218	NSSPPSPNKNNNAVDSGIHLTIEMNK2. 2E+26	sp	P20020	AT2B1_HUMAN
		sp	P05534	1A24_HUMAN
		sp	P30447	1A23_HUMAN
11219	GGSYSQAASSDSAQGSVDVSLTACK2E+22	sp	P01891	1A68_HUMAN
		sp	P01892	1A02_HUMAN
		sp	Q09160	1A80_HUMAN
		sp	P10316	1A69_HUMAN
11220	NEIIQSPISQVPSVEK20000	sp	Q7Z3T8	ZFY16_HUMAN
11221	HSDNPSEEGEVKDDGLEKSPMK2. 0002E+21	sp	014647	CHD2_HUMAN
11222	KQQHVISTEEGDMMETNSTDDEK110002200000	sp	Q9H0E3	SP130_HUMAN
11223	TISINSPK2002000	sp	Q9P1Y6	PHRF1_HUMAN
11224	LSQSSQDSSPVR2200020000	sp	095251	KAT7_HUMAN
11225	WDEQTSNTKGDDDEESDEEAVKK200000000	sp	043395	PRPF3_HUMAN
11226	STSYGYSR200	sp	Q13242	SRSF9_HUMAN
11227	VLSQSTPGTPSK20000	sp	Q6MZP7	LIN54_HUMAN
11228	RYSDLTTLVAFSSSVYPTK2000000000000000000	sp	P25205	MCM3_HUMAN
11229	AEPASPDSPKGSSETETEPVALAPGPAPTR2. 00022E+24	sp	P11474	ERR1_HUMAN
11230	LNSPPSSIYK200000000	sp	Q53EP0	FND3B_HUMAN

11231	KFELLTPPLSPSR20000	sp P04198 MYCN_HUMAN
		sp P01106 MYC_HUMAN
11232	MATLTSGGEKPNKDLESCSDDDNQGSK2E+25	sp Q5T3I0 GPTC4_HUMAN
11233	EEGSLSDTEADAVSGQLPDPTTNPAGK2.02E+25	sp Q8N1G4 LRC47_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
11234	EDALDDSVSSSVHASPLASSPVRK2.0200000000002E+19	sp O14715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
		sp A6NKT7 RGPD3_HUMAN
11235	NMAPSQSPVR20000	sp Q9NYB9 ABI2_HUMAN
11236	RGSPTTGFIEQK20000000000	sp Q6ZRV2 FA83H_HUMAN
11237	SHNNFVAILDPEGEHQYK2E+19	sp Q9Y478 AAKB1_HUMAN
11238	QSVTSFPDADAFHHQVHDDLLSSSEEECKDR2E+31	sp P50402 EMD_HUMAN
11239	DTHEDHDTSTENTDESNHDPQFEPIVSLPEQEI2E+22	sp P43487 RANG_HUMAN
11240	GHPAQRPEPSPPEPEPQAPEGSQAGAEGPSSPEASRSPAR200000020000	sp Q6ZW31 SYDE1_HUMAN
11241	MSQPGSPSKTPTPSTPGDTQPNTAPVPPAEDGIK2.0002E+21	sp Q14839 CHD4_HUMAN
11242	ATRPASTSQPDPSPDR200000020000	sp P20749 BCL3_HUMAN
11243	SPHQLLSPSSFSPSATPSQK202000000000	sp Q14181 DPOA2_HUMAN
11244	STSAPQMSPGSSDNQSSSPQPAQQK2.00002002E+23	sp Q14157 UBP2L_HUMAN
11245	ICTLSPSPPLASLAPVADSSTR200200000000000000	sp Q9UER7 DAXX_HUMAN
11246	RVSSDEEHTVDSCISDMK22000000000000100	sp Q70CQ2 UBP34_HUMAN
11247	DLSTSPKPSPIPSPVLGR2200000002000000	sp Q8NDI1 EHBP1_HUMAN
11248	EQDCGEPASPAASISR200000000	sp O60353 FZD6_HUMAN
11249	RSPSPTTPGPSR20200000000	sp P49450 CENPA_HUMAN
11250	SKPNLPSESRSR20200	sp Q08170 SRSF4_HUMAN
11251	ARPQPSGPAPSS20	sp Q96MG7 MAGG1_HUMAN
11252	SPPSVQSLAMR200000000100	sp Q96KQ7 EHMT2_HUMAN
11253	RPESAPAESSPSK220000	sp Q07889 SOS1_HUMAN
11254	SSGTPGCSSGTERPQSPGSSK2000000	sp Q8IYD8 FANCM_HUMAN
11255	TQSEESRPSLQQPATSTTETPASPAHTTPTQSTSGR2000000000000000	sp P15336 ATF2_HUMAN
11256	AVGMPSPVSPK102002000	sp Q9UHB7 AFF4_HUMAN
11257	ICTLSPSPPLASLAPVADSSTR200000000000	sp Q9UER7 DAXX_HUMAN
11258	VGASDGVFR2000000	sp Q9NYQ8 FAT2_HUMAN
11259	NAIASDSEADSDTEVPK200020000000	sp Q8WVC0 LEO1_HUMAN
11260	VASFIPGSSSR200000000000	sp Q14573 ITPR3_HUMAN
11261	ASSPTKTSPTTPEASATNSPCTSKPATPAPSEK2.0220002E+26	sp Q12873 CHD3_HUMAN
11262	TPTCQSSTDCSPYAR200000	sp O95235 KI20A_HUMAN
11263	FGSTGSTPPVSPTPSER2200000000002000	sp Q969V6 MKL1_HUMAN
11264	SPTTPIDPEK220000000	sp Q08AD1 CAMP2_HUMAN
11265	APSASDSKADSDGAKPEPVAMAR2.02020002E+21	sp Q7Z4V5 HDGR2_HUMAN
11266	LLASSPALPSSGAQAR200000000000	sp Q9NYZ3 GTSE1_HUMAN
11267	GLLSDSMTDVPVDTGVAAR200000000000000000	sp O43399 TPD54_HUMAN
11268	KAPPPPISPTQLSDVSSPR20020020000	sp Q9P0K7 RAI14_HUMAN
11269	SHSGTSPDNTAPPPPPRPQPSHSR2.022E+23	sp Q96D71 REPS1_HUMAN
11270	KSPAESPSTPR2000000022000	sp O76021 RL1D1_HUMAN
11271	TSPVVAPTSEPSSPLHTQLLK2200000000000000	sp O75044 SRGP2_HUMAN
11272	LALPPATKATTSEGGATSPSPSYSPDTPANR2300000000000	sp P08651 NFIC_HUMAN
11273	SRSRTPLLPR20202000000	sp Q9UQ35 SRRM2_HUMAN
11274	SRTPSRHSCSGSSPPR20220000	sp Q9UQ35 SRRM2_HUMAN
11275	HKSGSMEEDVDTSPGGDYTSPSSPTSSSR3000020000000	sp P08651 NFIC_HUMAN
11276	SKSPPKSPPEEGAVSS2000200000000000	sp Q01130 SRSF2_HUMAN
11277	SSSTEPPPPVR22000000000	sp Q8N684 CPSF7_HUMAN

11278	RHSSTGDSADAGPPAAGSAR2002000000000000	sp	Q6ZU35	K1211_HUMAN
11279	ALGVISNFQSSPK2000	sp	O94876	TMCC1_HUMAN
11280	DPHSPEDEEQPQGLSDDDILRDSGSDQDLDGAGVR2.0000000002E+32	sp	Q86VM9	ZCH18_HUMAN
11281	IHCGLQDSQK2000	sp	Q07864	DPOE1_HUMAN
11282	QPGQVIGATTPSTGSPTNK200000	sp	Q9ULM3	YETS2_HUMAN
11283	SMSFQGIR2000000	sp	P15924	DESP_HUMAN
11284	WCFSEPGK200000	sp	Q2M3G4	SHRM1_HUMAN
11285	ASEPLLSPPPFGESR2000000000	sp	O14686	KMT2D_HUMAN
11286	LQDSSDPDTGSEEEGSSRLSPPHSPR200200000000	sp	Q92974	ARHG2_HUMAN
11287	EPLENSASESSDELPEKER2020020000000000	sp	P78337	PITX1_HUMAN
11288	GLVSPQSPQK20000	sp	P41182	BCL6_HUMAN
11289	REEQTDTSDGESVTHHIR2002000000000000	sp	P36915	GNL1_HUMAN
11290	AAAYDISEDEED2000000	sp	P49585	PCY1A_HUMAN
11291	GPKSDDGFEIVPIEDPAK200000000000000000	sp	Q8IY81	SPB1_HUMAN
11292	TYSECEGTYSPEISWHHR200000000000000000	sp	P48651	PTSS1_HUMAN
11293	SQVNGEAGSYEMTNQHVK200000000000000000	sp	Q05D32	CTSL2_HUMAN
11294	ARSWSPPEVSR20000000200	sp	Q9NR19	ACSA_HUMAN
11295	METVSNASSSNPSSPGR200200000	sp	Q9NTI5	PDS5B_HUMAN
11296	LSSLRASTSK2200020000	sp	P62753	RS6_HUMAN
11297	LKATVTPSPVK200020000	sp	Q9H1E3	NUCKS_HUMAN
11298	DGDSYDPYDFSDTEEMPQVHTPK202000100002000	sp	P25205	MCM3_HUMAN
11299	MLRPVSR200	sp	Q5VZK9	LR16A_HUMAN
11300	YSPTSPTYSPTSPPK20200000000	sp	P24928	RPB1_HUMAN
11301	LTPVRPAAASPIVSGAR200000000	sp	Q9Y2K7	KDM2A_HUMAN
11302	RTPSDDEEDNLFAPPK2000000000000000	sp	Q641Q2	FA21A_HUMAN
11303	NFYESDDDQKEK200000000	sp	Q9Y4E1	FA21C_HUMAN
11304	YPLCSADAAVSTTPPGSPPPPPPLPEPPVLK2.000002E+21	sp	O00203	AP3B1_HUMAN
11305	EAATQEDPEQLPELEAHGVSESEGEERPVAEEEILLR2020000000000000000000	sp	Q9BVJ6	UT14A_HUMAN
11306	FPEFCSSPSPVEVK20000000	sp	Q68E01	INT3_HUMAN
11307	TAPVASTPSSTPIPALSPPTK200000	sp	Q7Z3K3	POGZ_HUMAN
11308	TIGGGDDSFNTFFSETGAGK20000000000	sp	P68363	TBA1B_HUMAN
11309	LDSEMDSHSENYDTMSSPLPGK1002000030100000000	sp	Q71U36	TBA1A_HUMAN
11310	ASPHATPSTLHFPTSPIIQPGPYFSHPAIR20000000000000000000	sp	Q9BQE3	TBA1C_HUMAN
11311	KPISDNSFSSDEEQSTGPIK2002000000000000	sp	Q6PEY2	TBA3E_HUMAN
11312	TKSSSDLSPPR22020020000	sp	Q13748	TBA3C_HUMAN
11313	AISTPETPLTK200200000	sp	Q9NTI5	PDS5B_HUMAN
11314	LNETELTDLEGQESPPKNYLCIEEEK2E+21	sp	Q12857	NFIA_HUMAN
11315	VSPSTSYTPSR20002000000	sp	O60293	ZC3H1_HUMAN
11316	ASHSSERPPPR20020000000	sp	Q9BRD0	BUD13_HUMAN
11317	KEKTPATTPEAR2000200000	sp	Q14181	DPOA2_HUMAN
11318	LLDRSPSR20000	sp	Q9ULF5	S39AA_HUMAN
11319	RRSPADAIPK2000000000	sp	Q9BTE3	MCMBP_HUMAN
11320	VAAAAGSGSPSPGSPGHDR2000000	sp	O15027	SC16A_HUMAN
11321	QSAGPNSPTGGGGGGSGGTR2000000000000000	sp	Q8NI27	THOC2_HUMAN
		sp	O76039	CDKL5_HUMAN
		sp	P18754	RCC1_HUMAN
		sp	Q9BTU6	P4K2A_HUMAN
		sp	A7E2V4	ZSWM8_HUMAN

		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
		sp Q29960 1C16_HUMAN
11322	GGSCSQAASSNSAQGSDESLIACK2000000000	sp Q29963 1C06_HUMAN
		sp P30504 1C04_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
11323	SVSTSSPAGAAIASTSGASNSSSN200	sp Q86UE8 TLK2_HUMAN
11324	SDGACDSPSSDKENSSQIAQDHQK2000000000	sp Q9UIG0 BAZ1B_HUMAN
11325	LSPERSSDESSENEPHLPHGVSQR2.0002E+24	sp Q9BQ52 RNZ2_HUMAN
11326	ALEAASLSQHPPSLCISDSEEEEEER2.00002E+19	sp O43159 RRP8_HUMAN
11327	DGTAPPPQSPGSPGTGQDEEWSDEESPRK2.000002E+21	sp Q9UKS6 PACN3_HUMAN
11328	SPSGHPHVR20000000	sp Q8WX93 PALLD_HUMAN
11329	NYSVGSRPLKPLSPLR200000000020000	sp O60825 F262_HUMAN
11330	NATQPPNAEEESGSSSASEEEDTKPKPTK2.000000020222E+27	sp Q68E01 INT3_HUMAN
11331	GSGGLFSPSTAHPD GALGQR20000000000000	sp Q9C0C2 TB182_HUMAN
11332	SAYQEYDSDSVPEELKR30002000000000	sp Q96JG6 CC132_HUMAN
11333	RESESEDETTPAAPQLIK202020020000000000	sp O60832 DKC1_HUMAN
11334	GSSPSIRPIQGSQSSSPVEK200000	sp Q92560 BAP1_HUMAN
11335	EVVKPVPITSPAVSK20000000	sp Q96HC4 PDLI5_HUMAN
11336	GPSPEGSSSTESSPEHPPK200000000020000000	sp Q9UQ35 SRRM2_HUMAN
11337	YRPASVPPSPSLSR20000020000	sp P13807 GYS1_HUMAN
11338	LLKPGEPESEYTDDEEDTKDHNKQD203000000000000000	sp O15173 PGRC2_HUMAN
11339	RSSQPSPTAVPASDSPPTK20000000000000000000	sp Q3KQU3 MA7D1_HUMAN
11340	TLSSSAQEDIIR20000000000000	sp Q9H788 SH24A_HUMAN
11341	GPSPEGSSSTESSPEHPPK20002220000000000000	sp Q9UQ35 SRRM2_HUMAN
11342	DHNVGSPKSPHTASPHQHTQTEK20000000002000000000	sp A0JNW5 UH1BL_HUMAN
11343	SPSPEPIYNSEGK200000000000	sp Q15637 SF01_HUMAN
11344	SAFTPATATGSSPSPVLGQGEK2020000000000000	sp Q15811 ITSN1_HUMAN
11345	GQTPLTEGSEDL DGHSDPEESFAR2.0020000002E+19	sp Q86VR2 F134C_HUMAN
11346	SFPLFDNSK2000000000	sp Q5T0W9 FA83B_HUMAN
11347	NKASPQSEFMPSK200000000000	sp Q9NQW6 ANLN_HUMAN
11348	QIGGGIQSITYTHNGDISR200	sp O00411 RPOM_HUMAN
11349	TKSPTDDEVTPSAVVR20000000000000000000	sp Q9Y2H5 PKHA6_HUMAN
11350	EYIPGQPPLSQSSDSSPTR220000	sp P07814 SYEP_HUMAN
11351	GASSPYGAPGTPR20000002000	sp Q5UIP0 RIF1_HUMAN
11352	DKDSGSSSPLPK22000000	sp Q86VQ1 GLCI1_HUMAN
11353	TEMKSPFNPSPPQDSR2000002000	sp P08651 NFIC_HUMAN
11354	STLESEKPGSPEAAETSPPSNIIDHCEK2E+24	sp Q96T23 RSF1_HUMAN
11355	EMEHNTVCAAGTSPVGEIGEEK1.0000000002E+21	sp P18583 SON_HUMAN
11356	SSDQDSTNKEAEAAGVKPAGVRPR2.200002E+24	sp Q92625 ANS1A_HUMAN
11357	GHPAQRPEPSPPEPEPQAPEGSQAGAEGPSSPEASRSPAR200002000000	sp Q6ZW31 SYDE1_HUMAN
11358	SASLSHPGGEGERPAR2000000000000000	sp Q2M3G4 SHRM1_HUMAN
11359	YHGHSMSDPGVSYR1200000000	sp P08559 ODPA_HUMAN
		sp P29803 ODPAT_HUMAN
11360	VLSGNCNHQEGTSSDDELPSAEMIDFQK2.0000000002E+26	sp P16383 GCFC2_HUMAN
11361	SLTELVQQPCPIEASK200000000000000000	sp P39880 CUX1_HUMAN
11362	RRSPSPYYSR200000000	sp P62995 TRA2B_HUMAN
		sp Q13595 TRA2A_HUMAN

11363	IGSPLSPK2002000	sp Q659C4 LAR1B_HUMAN
11364	GQSQTSPDHRSDTSSPEVR2200002000000000	sp Q9UQ35 SRRM2_HUMAN
11365	AADPPAENSSAPEAEQGGAE2000000000000	sp P67809 YBOX1_HUMAN
11366	KAEAAAAPTVPAGPAQPGHVSPTPATTSPGEK20200000	sp Q9Y6R0 NUMBL_HUMAN
11367	SRSWTSFK2020000	sp 075494 SRS10_HUMAN
11368	DDATPVRDEPMDAESITFK20000000000000000	sp Q7Z6E9 RBBP6_HUMAN
11369	DRESRSPFLKPDER202000000000	sp Q9H7N4 SFR19_HUMAN
11370	SQGSPEDESAPQASPGPSK2000000000000	sp Q9P107 GMIP_HUMAN
11371	TPASTPVSQTPQASPMIER20000000002010000	sp Q9UHR4 BI2L1_HUMAN
11372	HGSYEDAVHSGALND2000000000000000	sp P17987 TCPA_HUMAN
11373	SRSRTPLISR20202000000	sp Q9UQ35 SRRM2_HUMAN
11374	GGAPEGPNEAEVTSGKPEQVDPDAEEKSVSGTDVQEECR2000000000000	sp P49321 NASP_HUMAN
11375	SPASCHEHPMADDPSK20000000000000000	sp Q9BXB5 OSB10_HUMAN
11376	SWDSSSPVDRPEPEEAASPTR22000000000000000	sp Q86WB0 NIPA_HUMAN
11377	GEGDAPFSEPGTTSTQRPSSPETATK2200000000000000	sp Q9UQ35 SRRM2_HUMAN
11378	APPPISPTQLSDVSSPR2000020022000	sp Q9P0K7 RAI14_HUMAN
11379	DSGNWDTSGSELSEGELEKR202002000000000	sp 075400 PR40A_HUMAN
11380	SSGHSSSELSPDAVEK20000000020000000	sp Q9UQ35 SRRM2_HUMAN
11381	SRERPSQPEGLR200000000000000	sp Q04637 IF4G1_HUMAN
11382	RSPSPYYSR200000000	sp P62995 TRA2B_HUMAN
		sp Q13595 TRA2A_HUMAN
11383	LPQSSSESSPSPQPTK2022002000000	sp Q9UQ35 SRRM2_HUMAN
11384	AAPAQVRPPSPGNIRPVK2000000000	sp Q14244 MAP7_HUMAN
11385	SWSLIKNTCPPKDDPQVMEDK200000000010000	sp P79522 PRR3_HUMAN
11386	VSPPTPAR20000000	sp Q9H7Z6 KAT8_HUMAN
11387	SESARGDSEDEVLR20002020000000	sp Q8WVC0 LEO1_HUMAN
11388	VAEPGAEATSSTGEESGSEHPPAVPMHNK200020200000000000	sp Q14684 RRP1B_HUMAN
11389	WSAEASGKPSPSDPGSGTATMMNSSSRGSSPTR11000200220000	sp Q5JRA6 MIA3_HUMAN
11390	ASAGPAPGPVVTAEGLHPSLPSPTGNSTPLGSSK2.0000000002E+33	sp Q4KMP7 TB10B_HUMAN
11391	GRLPNSSRPSTPTINVLESK220000000000	sp Q15910 EZH2_HUMAN
11392	SSGHSSSELSPDAVEK20020000000	sp Q9UQ35 SRRM2_HUMAN
11393	MLDITPGCK200000	sp Q96N23 CFA54_HUMAN
11394	WSPPQNYK20000000	sp Q96K21 ANCHR_HUMAN
11395	SSGTNFQGLPSK200000000000	sp Q68EM7 RHG17_HUMAN
11396	SITSPPTSTTK2000000000000	sp Q14938 NFIX_HUMAN
11397	STSATDTHHVMAR20000000001000	sp 014874 BCKD_HUMAN
11398	KGSTFRPHDSFVK220000000000	sp Q9P206 K1522_HUMAN
11399	LSSAGPRSPYCK220000000000	sp Q6PL18 ATAD2_HUMAN
11400	VETVSPSESPK200200000	sp Q99590 SCAFB_HUMAN
11401	STSGRPGCQPSPCVVK20000000000000000	sp 014545 TRAD1_HUMAN
11402	LLSSNPPTSASESARIIGVSHCTQPQVASLSR2.0000002002E+21	sp Q6ZUF6 NC336_HUMAN
11403	FHFPPLDTHSPTNDVQGR2000000000000	sp P41162 ETV3_HUMAN
11404	LTPVSPSSSTEEK200000000000000	sp Q13501 SQSTM_HUMAN
11405	SDGACDSPSSDKENSSQIAQDHQK2000022000000000	sp Q9UIG0 BAZ1B_HUMAN
11406	GLPLGSAVSSPVLFPVGR200200000200000	sp Q8WUM0 NU133_HUMAN
11407	SRHSSISPSTLTLK2002202000000000	sp Q14004 CDK13_HUMAN
11408	ATLLNVPDLSDSIHSANASER2E+20	sp Q07157 ZO1_HUMAN
11409	NSVASPTSPTR20000	sp Q96HB5 CC120_HUMAN
11410	SQDSYPGSPSLSPR2000	sp Q6VN20 RBP10_HUMAN
11411	RPSLPSSPSPGLPK2000200000000	sp 043294 TGF11_HUMAN
11412	IPESETESTASAPNSPR20000002000	sp Q07889 SOS1_HUMAN
11413	SDISDQEEDEESEGCPVSNLSK2.00200000002E+23	sp Q15154 PCM1_HUMAN
11414	RKSHEAEVLK200000000	sp P16949 STMN1_HUMAN

11415	KGNSPNSPPTPK20000000	sp P48634	PRC2A_HUMAN
11416	SILSPGGSCGPIK20020000000000	sp P78347	GTF2I_HUMAN
11417	STQPDVCASPQEKPLR200000000	sp Q3T8J9	GON4L_HUMAN
11418	TNLTSQSSTNLPSPGSPGSPGSPGSPGSPK20000000000	sp Q6IN85	P4R3A_HUMAN
11419	SSGHSSSELSPDAVEK22000020020000000	sp Q9UQ35	SRRM2_HUMAN
11420	VEQATKPSFER20000000	sp Q96E39	RMXL1_HUMAN
11421	EHSYPGSPPLGWPSSETR300200000000000	sp O15320	CTGE5_HUMAN
11422	NKEESSDDEKESSEPPK2200000020000000	sp P35659	DEK_HUMAN
11423	TTSPEPQESPTLPSTEGQVVK2.000002E+20	sp P53814	SMTN_HUMAN
11424	TVGALPR20000000	sp Q9BSV6	SEN34_HUMAN
11425	TPSPGPVESK200000000	sp Q5T0W9	FA83B_HUMAN
11426	NSSTPGLQVPVSPTVPIQNQK2000000020000000000	sp Q03164	KMT2A_HUMAN
		sp P49792	RBP2_HUMAN
		sp Q7Z3J3	RGPD4_HUMAN
11427	EDALDDSVSSSVHASPLASSPVRK2.0200000000002E+19	sp A6NKT7	RGPD3_HUMAN
		sp Q99666	RGPD5_HUMAN
		sp O14715	RGPD8_HUMAN
11428	KKPEDSPSDDDLIVYELTPTAEQK2000000000000000000	sp P49792	RBP2_HUMAN
11429	INPPSSGGTSSSPIK200000	sp P14859	PO2F1_HUMAN
11430	TPSSSPITPPASETK2220002000000000	sp Q9NZJ0	DTL_HUMAN
11431	EQSTPPKSPQ200002000	sp Q92615	LAR4B_HUMAN
11432	QRGSETGSETHESDLAPSDK20202000000000000	sp P48634	PRC2A_HUMAN
11433	SRESLNVDVVK200000000	sp Q07866	KLC1_HUMAN
11434	SSTVATLQGTDPDHG DPR200000000000000	sp Q8IVT2	MISP_HUMAN
11435	VKEEPSPPQSPR20000000	sp Q00613	HSF1_HUMAN
11436	AELGMDSTSQSPPIK1000020000000	sp Q9NR19	ACSA_HUMAN
11437	YSPTSPTYSPK30000000002000	sp P24928	RPB1_HUMAN
11438	TRSPSPTLGESLAPHK2020000000000	sp Q86UU1	PHLB1_HUMAN
11439	AELGMDSTSQSPPIKR10000002000000	sp Q9NR19	ACSA_HUMAN
11440	SLSEEKEDHSDGLAGLK20000000000000000	sp Q6P0Q8	MAST2_HUMAN
11441	AASSSPGSPVASSPSR20000000000000000	sp Q92538	GBF1_HUMAN
11442	EKEEEEEPEGKPSDAESSESDDEK20002202200000	sp Q9BSC4	NOL10_HUMAN
11443	KLSLSEGK2020000	sp Q49A26	GLYR1_HUMAN
11444	RHSSVSDSQPCEPPSVGTEYSQGASPPQHQLK2.2E+31	sp P39880	CUX1_HUMAN
11445	THCVGDSQSSASSPPATSK20000000000	sp Q9H4Z2	ZN335_HUMAN
11446	TLSPATAPTSQGIPTSDEESTPKK2000200000	sp Q9NQS7	INCE_HUMAN
11447	SFNYSNSSTSEVSSTSASK200000000000000	sp Q5VT52	RPRD2_HUMAN
11448	SGTSSPQSPVFR200200000	sp Q8N556	AFAP1_HUMAN
11449	ESEKLENETLNK2002000000000	sp P23588	IF4B_HUMAN
11450	GDTPGHATPGHGATSSAR2000	sp O75533	SF3B1_HUMAN
11451	SPGRPTQGALGEQQDLSNTTSK2E+22	sp P10071	GLI3_HUMAN
11452	LDSQPQETSPELPR2000020000000	sp O14545	TRAD1_HUMAN
11453	LSGEHSESSTRPR22000000	sp Q7Z3K3	POGZ_HUMAN
11454	SHSGTSPDNTAPPPPPRPQPSHSR2E+23	sp Q96D71	REPS1_HUMAN
11455	SPFNPSPPQDSR2000	sp P08651	NFIC_HUMAN
11456	SHTPSPTSPK2002000	sp P40123	CAP2_HUMAN
11457	STSPAGQHHSPISSR200000002000000	sp Q5T200	ZC3HD_HUMAN
11458	GQSQTSPDHRSDTSSPEVR2022000020000000000	sp Q9UQ35	SRRM2_HUMAN
		sp Q9Y4E1	FA21C_HUMAN
11459	SRPTSFADLAAR2000000000	sp Q641Q2	FA21A_HUMAN
		sp Q5SNT6	FA21B_HUMAN
11460	HIVNHDDVFESEELSSDEEMK200002000100	sp Q96RT1	LAP2_HUMAN
11461	EVLTPSTSDNETR202200000000	sp Q70CQ2	UBP34_HUMAN
11462	SHSTEPNLSSFLNDPNMK20000000000000100	sp Q9UPQ0	LIMC1_HUMAN



11463	AMSTTSSISSPQPGK10000002000000	sp Q9UJU6 DBNL_HUMAN
11464	GSVSQPSTSPPKPTGIFQTSANSSFEVK2. 202E+24	sp O15027 SC16A_HUMAN
11465	GPAETAAADSEPTDPEEER200020000000	sp Q5XUX0 FBX31_HUMAN
11466	DHSDSDDQMLVAK202000100000	sp Q9NW75 GPTC2_HUMAN
11467	SGEQITSSPVSPK20000200002000	sp O43314 VIP2_HUMAN
11468	SPHSPSIATPPSQSSSCSDAPMLSTVHSAK2. 002E+31	sp Q9NYF8 BCLF1_HUMAN
11469	DSMSCGLDSEGATPPPVGRLR2000000000	sp Q99956 DUS9_HUMAN
11470	MSSPPSPQK200020000	sp Q9Y2H6 FND3A_HUMAN
11471	YFQSPSRSR2020200	sp Q16629 SRSF7_HUMAN
11472	STSAPQMSPGSSDNQSSSPQPAQK1. 200220002E+19	sp Q14157 UBP2L_HUMAN
11473	TGSNISGASSDISLDEQYKHQLEETKK2. 0020022E+25	sp P22059 OSBP1_HUMAN
11474	AVSLDSPVSVGSSPPVK2000000000200000	sp Q9Y6Q9 NCOA3_HUMAN
11475	IDAGTMAEPSASPSK10002000000	sp Q8IXQ3 CI040_HUMAN
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11477	SSTDLSLPGISR2200000000000	sp Q9HCD5 NCOA5_HUMAN
11478	SPPRPPR20000000	sp Q9UQ88 CD11A_HUMAN
		sp P21127 CD11B_HUMAN
		sp Q9UKB5 AJAP1_HUMAN
11479	SLRSPLPSR202000000	sp Q9NYV4 CDK12_HUMAN
11480	GAAAADLLSSSPESQHGGTQSPGGGQPLLQPTK20000000000000	sp Q9H3S7 PTN23_HUMAN
11481	GRPSKTPSPSQPK20202000000	sp Q9NTI5 PDS5B_HUMAN
11482	LNNSSPNSSGGVK2000000000	sp O00472 ELL2_HUMAN
11483	SRKSPSPAGGGSSPYSR202000002000000	sp Q14004 CDK13_HUMAN
11484	RASSPSLIER200200000	sp A1L170 CA226_HUMAN
11485	FDHESPGTDEDKSG220020000000	sp Q13427 PPIG_HUMAN
11486	KERPISMINASNYNVTSDYAVHPMPVGR1. 00000000022E+24	sp Q9HAU0 PKHA5_HUMAN
11487	SVVSFDK20020000	sp Q15424 SAFB1_HUMAN
11488	SQDSYPGSPSLSPR230020002000	sp Q6VN20 RBP10_HUMAN
11489	RSSQPSPTAVPASDSPPTK202000020000000	sp Q3KQU3 MA7D1_HUMAN
11490	RASPPPK200000	sp Q8IYB3 SRRM1_HUMAN
11491	LSGEHSESSTPRPR20000000200000	sp Q7Z3K3 POGZ_HUMAN
11492	QQSLGGSESPGAPCSPGLSASPASSPGR2E+27	sp Q8N612 F16A2_HUMAN
11493	ALFKPPEDSQDDESDSDAEEEEQTTKR20000202000000000000	sp Q13769 THOC5_HUMAN
11494	QPSPSPPTER2000002000	sp P29590 PML_HUMAN
11495	SAGGSPEGGEDSDREDGNYCPPVK2E+21	sp Q9H6Z4 RANB3_HUMAN
11496	AGSSPAQGAQNEPPR2000000000000	sp Q9UGU0 TCF20_HUMAN
11497	ALSSAVQASPTSPGGSPSSPSSGQR2200000000000000	sp Q03164 KMT2A_HUMAN
11498	RRSPSPYYSR200000200	sp Q13595 TRA2A_HUMAN
		sp P62995 TRA2B_HUMAN
11499	VIEPPAASCSSPR20002000	sp Q3MIN7 RGL3_HUMAN
11500	APSASDSKADSDGAKPEPVAMAR2. 0202020002E+23	sp Q7Z4V5 HDGR2_HUMAN
11501	APSASDSKADSDGAKPEPVAMAR2. 0202020002E+23	sp Q7Z4V5 HDGR2_HUMAN
11502	RGEDAPFSEPGTTSTQRPSPETATK20000000000000	sp Q9UQ35 SRRM2_HUMAN
11503	SKSSDPDLR2020000000	sp O60293 ZC3H1_HUMAN
11504	EGPLGWSSDGLGSLVLLDSPSSPR2000	sp Q9UGP4 LIMD1_HUMAN
11505	NSVLSDPGLDSPRTSPVIMAR200000000200000000	sp O15085 ARHGB_HUMAN
11506	SNSQSDSHDEEVSTPPNPVK2. 02E+20	sp P31321 KAP1_HUMAN
11507	LLSFSPEEFPTLK2000000000	sp Q5JSZ5 PRC2B_HUMAN
11508	VAEPGAEATSSTGEESGSEHPPAVPMHNK2000000000000	sp Q14684 RRP1B_HUMAN

		sp P30504 1C04_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
11509	GGSCSQAASSNSAQGSDESLIACK2000000000000000	sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
		sp P30508 1C12_HUMAN
		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
11510	SDSIRPALNSPVERPSSDQEEGETSAQTER2000000000000000	sp Q9UKV5 AMFR_HUMAN
11511	SSPSKENESSDSEEEVSRPR2000022020000000000	sp Q6KC79 NIPBL_HUMAN
11512	SPSGSAFGSQENLR2000000000000000	sp Q96N67 DOCK7_HUMAN
11513	GGPASVPSSPGTSVK2000000000	sp O43237 DC1L2_HUMAN
11514	SKSPPKSP EEGAVSS200000200000000000	sp Q01130 SRSF2_HUMAN
11515	ELSNSPLRENSFGSPLEFR20000020000000000	sp Q9UQ35 SRRM2_HUMAN
11516	ATSPSSVSGDFDDGHHSVSTPGPSR2. 2202E+22	sp Q86U86 PB1_HUMAN
11517	QLQAPAGSPAPSPSPGGDDKPQVPPR2. 000002E+19	sp Q07912 ACK1_HUMAN
11518	GESAPTLSTSPSPSSPSPTSPSPTLGR2. 2E+20	sp Q9H330 TM245_HUMAN
11519	LGSTGSQPNSEAESVPENVPKPLLPPK2E+23	sp Q8IZ21 PHAR4_HUMAN
11520	LPASPSGSEDLSSVSSSPTSSPK202000000000	sp Q9P2N6 KANL3_HUMAN
11521	EYSPPPSPPLR30000020000	sp Q9HCD6 TANC2_HUMAN
11522	SPQRSPLQSAESSPTAGK2000000000002020000	sp Q8IWB9 TEX2_HUMAN
11523	SGTTPRQGSITSPQANEQSVTPQRR202200000020000000	sp Q9UQ35 SRRM2_HUMAN
11524	SHSESASPSALSSSPNNLSPTGWSQPK2. 002E+19	sp P04049 RAF1_HUMAN
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11526	LPSKELPDSSSPVANNIR20000000000	sp Q9BWT3 PAPOG_HUMAN
11527	YSPSPPK300200000	sp Q8IYB3 SRRM1_HUMAN
11528	SQSIEQESQEK2000000000	sp Q5VTL8 PR38B_HUMAN
11529	DDQVGWSSPPPR200000	sp P17927 CR1_HUMAN
11530	SGSNTGLDYIK2000000000	sp P35367 HRH1_HUMAN
11531	HVTLGPGQSPLSR200000	sp O15061 SYNEM_HUMAN
11532	TKSSSDLSPPRR220000200000	sp Q9BRD0 BUD13_HUMAN
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11534	YIASVQGSTPSPR202000	sp P49792 RBP2_HUMAN
11535	KIPLKPSPSK20000	sp Q99755 PI51A_HUMAN
		sp A2A3N6 PIPSL_HUMAN
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11537	VSPGLPSPNLENGAPAVGPVQPR2. 00002E+22	sp Q9UGP4 LIMD1_HUMAN
11538	HADHSSLTLGSGSSTTR200	sp Q7Z6Z7 HUWE1_HUMAN
11539	QSPGHQSPLASPK2000000002000	sp Q8N1G0 ZN687_HUMAN
11540	TSLPCIPR20000000	sp Q14432 PDE3A_HUMAN
11541	YSPTSPTYSPVYPTSPK2000	sp P24928 RPB1_HUMAN
11542	VTSSMSSPMQPK20000000	sp Q13136 LIPA1_HUMAN
11543	STSAPQMSPGSSDNQSSSPQPAQQK2. 00012E+23	sp Q14157 UBP2L_HUMAN
11544	ALLSSPEGEEK220000000	sp O75592 MYCB2_HUMAN
11545	RTTSPSPEPDR2000000000	sp Q9NQ75 CASS4_HUMAN
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11547	CVSPIVPSPTS200000	sp Q86TC9 MYPN_HUMAN
11548	APQTSSPPVVR2200000000	sp Q8IYB3 SRRM1_HUMAN
11549	TEFLHSQNSLSPR200002000	sp Q96FS4 SIPA1_HUMAN
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11551	TPQSPTLPPAK200000000000	sp P78310 CXAR_HUMAN

11552	QFPPSPSAPK2000000	sp 075381 PEX14_HUMAN
11553	LEAIITPPPAK2000000	sp 075367 H2AY_HUMAN
11554	IFSNQPDSPIASQDK2000000000	sp 075161 NPHP4_HUMAN
11555	DTPGHGSGWAETPR20000200000000	sp 075533 SF3B1_HUMAN
11556	EVNSCTTGSSNSTIIGSQGSETPK2.022E+21	sp Q5T0W9 FA83B_HUMAN
11557	HLSLDDIR2000000	sp Q96NE9 FRMD6_HUMAN
11558	ALQHTESPSETNKPHSK200000000000	sp Q14669 TRIPC_HUMAN
11559	KVVVSPTK200	sp P19338 NUCL_HUMAN
11560	SASPYPSHLSLSPQR2020002000000000	sp Q5T200 ZC3HD_HUMAN
11561	EEASLLSHSPGTSNQSQPCSPKPIR2002000000000000000	sp Q9H4Z3 PCIF1_HUMAN
11562	TSPTTPEASATNSPCTSKPATPAPSEK2E+26	sp Q12873 CHD3_HUMAN
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11564	TASISSPSEGTPTVGSYGCTPQLPK2.022E+25	sp Q6PKG0 LARP1_HUMAN
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11567	VVLGDGVQLPPGDYSTTPGGTLFSTTPGGTR22000000	sp Q13541 4EBP1_HUMAN
11568	RSSSSSAAASTPPGPPAPADPLGYLPLHGGYQYK2.20002E+31	sp Q8NDT2 RB15B_HUMAN
11569	GQSQTSPDHRSDTSSPEVR200002020000000	sp Q9UQ35 SRRM2_HUMAN
11570	GSISTPNSPVK20000	sp Q96AY4 TTC28_HUMAN
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11576	VKDEFSDLSEGDVLESEENDKK200200000200000000	sp Q96E39 RMXL1_HUMAN
11577	SPGAPCQECSR200000000000	sp Q8N554 ZN276_HUMAN
11578	SRYSRSPYSR20200000	sp Q9NWF9 RN216_HUMAN
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11581	GSDHSSDKSPSTPEQGVQR200000000000	sp Q9NVD7 PARVA_HUMAN
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11584	EAAFSPGQQDWSR2000000000	sp Q9UEY8 ADDG_HUMAN
11585	VEEESTGDPFGFDSDESPLVSSK2E+19	sp Q9C0C2 TB182_HUMAN
11586	AFGESSTESDEEEEGCGHHCVR2.202E+19	sp Q7Z5K2 WAPL_HUMAN
11587	FSDSEDEAATVPGNEPTASTPK2.000002E+20	sp O60927 PP1RB_HUMAN
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11594	NACILQASGSPPAKSPTDPK20000002000000	sp Q9P0K7 RAI14_HUMAN
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11596	KSSPSKENESSDSEEEVSRPR2.02000022E+19	sp O43237 DC1L2_HUMAN
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11598	ELEDSPQENVSVTEIIKPCDDPK2E+19	sp Q6NZY4 ZCHC8_HUMAN
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11601	KVEGNFNPFAFPK20000	sp P62995 TRA2B_HUMAN
11602	DLPDGHAASPR2000	sp Q01167 FOXK2_HUMAN
11603	DRTPELLYR20000000	sp Q9H6K1 CF106_HUMAN
		sp Q9P1Y5 CAMP3_HUMAN
		sp Q96T37 RBM15_HUMAN

11604	YETPSDAIEVISPASSAPPQEK2.0000000002E+19	sp 075376 NCOR1_HUMAN
11605	LITSEERSPAK20000	sp Q12888 TP53B_HUMAN
11606	LPDGPTGSSEEEEFLEIPPFNK20020000000000000000	sp Q12888 TP53B_HUMAN
11607	DEDEDEDESSEEDSEDEEPPPK2200020000000000	sp Q5SSJ5 HP1B3_HUMAN
11608	SPSPSPTPEAK2000000000	sp Q8TAQ2 SMRC2_HUMAN
11609	NSNSPPSPSSMNQR20200002000000	sp Q7Z5L9 I2BP2_HUMAN
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11611	EYGSPLK20000	sp P21127 CD11B_HUMAN
		sp Q9UQ88 CD11A_HUMAN
11612	DAQRLSPIPEEVPK2000000000	sp Q96T23 RSF1_HUMAN
11613	LSSPA AFLPACNSPSK200	sp Q86YS7 C2CD5_HUMAN
11614	SRSYTPEYR20200000	sp Q13595 TRA2A_HUMAN
11615	DIEISTEEEEK2200000	sp Q4G0J3 LARP7_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
11616	EDALDDSVSSSVHASPLASSPVR20002000200000	sp A6NKT7 RGPD3_HUMAN
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		sp Q99666 RGPD5_HUMAN
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11619	SRPTSEGS DIESTEPQK20000000000000000000	sp Q4G0J3 LARP7_HUMAN
11620	SSTPLHSPSPIR200002020000	sp O95817 BAG3_HUMAN
11621	EAATQEDPEQLPELEAHGVSESEGEERPVAEEEEILLR2E+34	sp Q9BVJ6 UT14A_HUMAN
11622	SDDSKSSPELVTHLK220000000000	sp Q07960 RHG01_HUMAN
11623	SSSISEEKGSDDEKPR2000002000000000	sp Q9UKV3 ACINU_HUMAN
11624	HLSESSGKPLSTK2000000000	sp Q8IVT2 MISP_HUMAN
11625	SRSHSPMSNR202000000	sp Q13595 TRA2A_HUMAN
11626	NDHLTSTSSPGVIVPESENK2000000000000000	sp O75496 GEMI_HUMAN
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11628	LLSSNEDDANILSSPTDR2000000000002000	sp O75448 MED24_HUMAN
11629	LSPSPSLR20200000	sp Q5T200 ZC3HD_HUMAN
11630	SGSYSGRSPSPYGR300020200000	sp Q9NYV4 CDK12_HUMAN
11631	WGQPPSPTPVPRPPDADPNTPSPKLEGRPER2.0000000000002E+27	sp Q14839 CHD4_HUMAN
11632	KFSAGGSDPPLK2000000	sp O75152 ZC11A_HUMAN
11633	KSPTMEQAVQTASAHLPAPAAVGR2E+23	sp Q8ND56 LS14A_HUMAN
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11635	NEMFESEWSKER10000020000	sp Q5VVM6 CCD30_HUMAN
11636	RESPSPAPKPR20000000	sp Q8IYB3 SRRM1_HUMAN
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11638	LPPPPGSPGLHSPTASPPPTAR20000000020000000	sp O95785 WIZ_HUMAN
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11640	SPSEESAPTTSPESVSGVPSSGSSGR2020000000000000	sp P54725 RD23A_HUMAN
11641	KPSPSESPKPFPAVSPPEPR20000000000000000000	sp Q96JM3 CHAP1_HUMAN
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11643	VAEPGAEATSSSTGEESGSEHPPAVPMHMK2020000000000000	sp Q14684 RRP1B_HUMAN
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11645	SCDELSPVSPQGGYPSEPTR20200000000000000000	sp Q9HCD6 TANC2_HUMAN
11646	ISDSESEDPPRNQASDSENEELPKPR2.0202000000002E+25	sp Q96ST2 IWS1_HUMAN
11647	KADSDSEDKGEESKPK2020000000000000	sp P83916 CBX1_HUMAN
11648	RTSVSPPEQPYPYR2000200000000000	sp Q8TEH3 DEN1A_HUMAN
11649	SQGDEAGGHGEDRPEPLSPK2E+20	sp Q9NZT2 OGFR_HUMAN
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11651	HIEPELAGRDSPIR20000	sp P86790 CCZ1B_HUMAN
		sp P86791 CCZ1_HUMAN
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11653	VRPSSDLNSTGQSPHHK200000000200000	sp P27448 MARK3_HUMAN
11654	WGQPPSPTPVPRPPDADPNTSPKPLEGRPER2.000000000002E+25	sp Q14839 CHD4_HUMAN
11655	IEIPVTPTGQSVSPSPSIPGTPTLK2200000000000	sp Q9H4Z3 PCIF1_HUMAN
11656	HRSPPATPPPK2000000000	sp Q8IYB3 SRRM1_HUMAN
11657	VVPGQFDDADSSDSEN2020000	sp Q9BRS2 RIOK1_HUMAN
11658	YRPASVPPSPSLSR200000000200	sp P13807 GYS1_HUMAN
11659	IPVSPEQAR2000000	sp Q9H981 ARP8_HUMAN
11660	KGTTPPRSPEASPK200000002000	sp Q9HDC5 JPH1_HUMAN
11661	AVTPVSQGSNSSSADPK200000000000000	sp 000512 BCL9_HUMAN
11662	KVSSSSPQSGCPSPTIPAGK2200000000002000000	sp Q76L83 ASXL2_HUMAN
11663	SPIALPVK200000000	sp Q8TF01 PNISR_HUMAN
11664	LQQQHSEQPPLQSPVMTR2001000	sp Q5JTV8 TOIP1_HUMAN
11665	FKSDSGSLGDAK20000000000	sp Q5T5U3 RHG21_HUMAN
11666	SRSPDIQAK2020000000	sp Q9HCS5 E41LA_HUMAN
11667	SRSGSQELDVKPSASPQER2.00022E+20	sp Q9UQ35 SRRM2_HUMAN
11668	RNSVTPLASPEPTKKPR2000002000000000	sp Q16875 F263_HUMAN
11669	APSDSSLGTPSDGRPELR200000000	sp Q15642 CIP4_HUMAN
11670	TPSPSYQR200020000	sp Q5T200 ZC3HD_HUMAN
11671	EPSVPASCAVSATYERSPSPAHLRDDPK3000020000000000	sp Q92615 LAR4B_HUMAN
11672	SELESGKFLPR200020000000	sp Q92628 K0232_HUMAN
11673	FYDLSDSNLSGEDSK30000200002000000	sp Q9H501 ESF1_HUMAN
11674	RASLSEIGFGK20000000	sp Q00537 CDK17_HUMAN
11675	NSDRSEDGYHSDGDYGEHDYR200030200000000000	sp P52756 RBM5_HUMAN
11676	SESFSEDLIPSR200000000000	sp Q9P219 DAPLE_HUMAN
11677	DYPLTPPPSPTVDEPK2000200000000	sp Q5THJ4 VP13D_HUMAN
11678	YGSFEEAEASYPILNR200000000000000	sp 095425 SVIL_HUMAN
11679	AALLAQYADVTDDEDEADEKDDSGATTMNIGSDK2E+24	sp Q96MW1 CCD43_HUMAN
11680	VQRPEDASGGSSPSGTSK20000000	sp Q9UI08 EVL_HUMAN
11681	NMVDLVNTHHLHSSSDEDDRLK2000000000	sp Q9UPN7 PP6R1_HUMAN
11682	IEEVLSPEGSPSKSPSK2000000020000	sp Q9UEY8 ADDG_HUMAN
11683	RSLSGSSPCPK2200000	sp Q9UQ35 SRRM2_HUMAN
11684	VSEEQTPPSPAGAGMSTAMGR10201000	sp Q16666 IF16_HUMAN
11685	SAGGSSPEGGEDSDREDGNYPVVKR2.2E+22	sp Q9H6Z4 RANB3_HUMAN
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11687	GMGPGTPAGYGR20000000	sp P23246 SFPQ_HUMAN
11688	SSTPSSPTGTSSSDSGHHIGWGER2E+20	sp P46019 KPB2_HUMAN
11689	SRSPQRPGWSR202000000000	sp Q9UQ35 SRRM2_HUMAN
11690	SLSELESK20000000	sp P22059 OSBP1_HUMAN
11691	VASPSQGVGSSSPKR200000000200000	sp Q99569 PKP4_HUMAN
11692	SPSTLLPK200000000	sp P27816 MAP4_HUMAN
11693	INHSSSLAR200000	sp 094915 FRYL_HUMAN
11694	GESAPTLSTSPSPSSPSPTSPSPTLGR2E+25	sp Q9H330 TM245_HUMAN
11695	RHSHSHSPMSTR20202002000	sp P62995 TRA2B_HUMAN
11696	NMVDLVNTHHLHSSSDEDDR2220000000	sp Q9UPN7 PP6R1_HUMAN
11697	SRSPSHRTRPR20200000000	sp Q8IYB3 SRRM1_HUMAN
11698	ALEYSNGIFDCQSPSPFMGSLR2.00000000002E+19	sp Q86W92 LIPB1_HUMAN
11699	SRESLNVDVVK200200000000	sp Q07866 KLC1_HUMAN
11700	SRSSSVGSSSYPISPAISR223002000000	sp Q15149 PLEC_HUMAN
		sp Q96HA1 P121A_HUMAN
11701	SQTPERPAK2020000000	sp A8CG34 P121C_HUMAN
		sp A6NF01 P121B_HUMAN

11702	QVVSVVQDEEVGLPFEASPESPPPASPDPGVTEIR2E+31	sp Q96TA1 NIBL1_HUMAN
11703	KTTEMQSPMQPVDQASLPGHCR2. 201E+21	sp P01589 IL2RA_HUMAN
11704	MSQPGSPSPKTPPTSTPGDTQPNTPAPVPPAEDGIK2. 02E+31	sp Q14839 CHD4_HUMAN
11705	TDSEKPFRRSQSPK2000000200000	sp Q9Y2W1 TR150_HUMAN
11706	ALPTSKPEGLHSSPVGPSSSK20020000000000	sp Q9UPT8 ZC3H4_HUMAN
11707	TSPTPPESATTVK20000000000000	sp Q5M775 CYTSB_HUMAN
11708	VPPAPVPCPPSPGPSAVPSSPK20000	sp O95817 BAG3_HUMAN
11709	RGTSPRPPEGGLGYSQLGDDDLK2. 000000000002E+21	sp P21127 CD11B_HUMAN
11710	SPSRPILDGK200000000	sp Q9UQ88 CD11A_HUMAN
11711	SVCPGGSSPGSSSGGR200000200000	sp O60890 OPHN1_HUMAN
11712	RESESEDETPPAAPQLIKK2020000000000000	sp Q53LP3 SWAHC_HUMAN
11713	AFQLEEGEETPDCK2000000	sp O60832 DKC1_HUMAN
11714	RRSPSPKPTK202000000	sp P38432 COIL_HUMAN
11715	SPPRPQR200000000	sp Q15287 RNPS1_HUMAN
11716	SLSYSPVER20000000	sp Q6PJG9 LRFN4_HUMAN
11717	SVASNQSEMEFSSLQDMPK20000000000000	sp Q9UQ35 SRRM2_HUMAN
11718	SCTPSPDQISHRASLEDAPVDDLTR2. 020200002E+25	sp Q8N163 CCAR2_HUMAN
11719	APGTPHSHTKPYVR200200000000	sp Q7Z2W4 ZCCHV_HUMAN
11720	RGSPSAAFTFPDTPDDFGK20000000000000	sp Q07020 RL18_HUMAN
11721	SDSDYDLSPK200000000	sp Q9ULT0 TTC7A_HUMAN
11722	SRSPEAGFCVYLK2020000000000000	sp Q08499 PDE4D_HUMAN
11723	NSFSEIRIHPLSQASGGSSNIRQEPAPK2. 0200000002002E+27	sp Q07343 PDE4B_HUMAN
11724	TATPPGYKPGSPSFR200000002000000	sp Q9NTZ6 RBM12_HUMAN
11725	ATGPPSQNSNIGTGRGSEGDCSPEDKNSVTAK20000200000000000	sp O76039 CDKL5_HUMAN
11726	SQGEEEGPLSDK20000000000000	sp P54259 ATN1_HUMAN
11727	MAESPCSPSQPPSPSPDEL PANVK1. 00000002E+27	sp Q9UKJ3 GPTC8_HUMAN
11728	SSTPLHSPSPIR20002020000	sp Q9H063 MAF1_HUMAN
11729	EGSPANWK2000000	sp Q14160 SCRIB_HUMAN
11730	VSSPVFGATSSIK200000000000	sp O95817 BAG3_HUMAN
11731	RPPGPTTSPASTSLSPGQR2000000002200000	sp O96019 ACL6A_HUMAN
11732	GILAADESTGSIK2000000	sp Q99708 COM1_HUMAN
11733	ATSPSSVSGDFDDGHHSVSTPGPSR2. 02E+20	sp Q01970 PLCB3_HUMAN
11734	FHSPSTWSPNKDTPQEK2020000000000000	sp ALDOA_RABIT
11735	SLDGAEFRPASVSENHDAGPDGDKR2. 000000000002E+26	sp P04075 ALDOA_HUMAN
11736	SRSPPVSK20000200	sp Q86U86 PB1_HUMAN
11737	RREEGPPSPDPGASSDAEPEPPSGR200000000000	sp Q99590 SCAFB_HUMAN
11738	TPLSFTNPLHSDSDSDERNSDGAVTQNK2. 002E+19	sp Q9H4G0 E41L1_HUMAN
11739	WGQPPSPTPVPRPPDADPNTSPKPLEGRPER2. 02E+27	sp Q01130 SRSF2_HUMAN
11740	DHNVGSPPKSPTHASQHTQTEK200000000000	sp Q9NTJ3 SMC4_HUMAN
11741	SWDSSPVDRPEPEAASPTTR200000	sp Q05209 PTN12_HUMAN
11742	TSGSDDPGICSNTDSTQAQVLLGK2E+23	sp Q14839 CHD4_HUMAN
11743	TEPAEAEAAAGSPSESPSPAEEELPGSHAEPVPAQGEAPGEQAR2E+46	sp A0JNW5 UH1BL_HUMAN
11744	SQPDPVDTPTSSKPQSK20000000000	sp Q86WB0 NIPA_HUMAN
11745	KASSDSEDSSEEEEEVQGPPAK2. 2020022E+20	sp Q8NDI1 EHBP1_HUMAN
11746	DQPPFGSDSDSVEADKSSPGIHLER2000002200000000	sp P55884 EIF3B_HUMAN
11747	KYDSSSLPPSNSGK20000000000	sp P46013 KI67_HUMAN
11748	RASHTLLPSHR20000000	sp Q14978 NOLC1_HUMAN
11749	AASSSPGSPVASSPSR200000000000	sp Q14676 MDC1_HUMAN
11750	RGTSPRPPEGGLGYSQLGDDDLK2. 00000000002E+20	sp Q68CP9 ARID2_HUMAN
11751	AAAAATAPPSPGPAQPGPR20000000000000	sp Q9UKV3 ACINU_HUMAN
		sp Q92538 GBF1_HUMAN
		sp Q9UQ88 CD11A_HUMAN
		sp P21127 CD11B_HUMAN
		sp Q6SPF0 SAMD1_HUMAN

11752	AQTTPGPSLSGSK200000000200	sp Q9UQ35 SRRM2_HUMAN
11753	DDPGQQTDSSEDEDIIGPMPAK2022000000000000	sp Q8IXQ4 GPAM1_HUMAN
11754	DRQSLDGFYSHGMAEGR2000000000000000	sp Q12802 AKP13_HUMAN
11755	LNEAQPSTIATSMRDSLIDSLT20	sp P56537 IF6_HUMAN
11756	RNSSSPVSPASVPGQR2002000000000	sp Q9H1B7 I2BPL_HUMAN
11757	RSSSPAELDLK2200000000	sp Q2PPJ7 RGPA2_HUMAN
11758	SRSPLHPMSEIPR20200000000000	sp Q9NVW2 RNF12_HUMAN
11759	STAPSAASASASAAAASSPAGGGAELELLEHCGVCR2E+21	sp Q13263 TIF1B_HUMAN
11760	GESAPTLSTSPSPSSPSPSPSPTLGR2. 2E+19	sp Q9H330 TM245_HUMAN
11761	LEDAESSNPNLQSLSTDALPSASK20000000000000	sp Q9Y294 ASF1A_HUMAN
11762	AASSSPGSPVASSPSR20200	sp Q92538 GBF1_HUMAN
11763	TLSPTPASATAPTSQGIPTSDEESTPK200020000	sp Q9NQS7 INCE_HUMAN
11764	STFSLPEEEEEPEPLVFAEQPSVK2. 02E+23	sp Q6Q0C0 TRAF7_HUMAN
11765	TKSSSDLSPPRR20200200200000	sp Q9BRD0 BUD13_HUMAN
11766	SLRQSFR20002000	sp Q6P1M3 L2GL2_HUMAN
11767	TSQLGDSPFYPGK2000000000000	sp Q15334 L2GL1_HUMAN
11768	SLDGAEFRPASVSENHDAGPDGDKR2000000000000000	sp P49790 NU153_HUMAN
11769	IEVLPVDTGAGGYSGNSGSPK2000000000000000	sp Q9H4G0 E41L1_HUMAN
11770	QPPPLAPQSPQGGVMGGSNSNQQQMR2. 000001E+19	sp Q92738 US6NL_HUMAN
11771	APVHFVEPLSPTGVAGHR2000000000	sp P46937 YAP1_HUMAN
11772	RLSSTSLASGHSVR2002000000000	sp Q5T5Y3 CAMP1_HUMAN
11773	DLSTSPKPSPIPSPVLGR2000020002000000	sp Q9BZL6 KPCD2_HUMAN
11774	SWDSSSPVDRPEPEAASPTR2. 0022E+21	sp Q8NDI1 EHBP1_HUMAN
11775	HALTSPSLGGQGR20000000000	sp Q86WB0 NIPA_HUMAN
11776	SSFNVSDVARPEAAGSPPEEGGCTEGTPAK2E+29	sp Q5PRF9 SMAG2_HUMAN
11777	TPAPDPGSPRTPPPTSR20000000200	sp Q92619 HMHA1_HUMAN
11778	SSSVGSSSSYPISPAUSR20002000000	sp Q96LL9 DJC30_HUMAN
11779	VLTQMGSPLNPISSVS2020	sp Q15149 PLEC_HUMAN
11780	FVHGSPSR2000	sp Q99717 SMAD5_HUMAN
11781	SAPTSPTATNISSK2000000000000000	sp Q5T200 ZC3HD_HUMAN
11782	AASADSTTEGTPADGFTVLSTK2000000000000000	sp Q9H792 PEAK1_HUMAN
11783	KAPSASDSKADSDGAKPEPVAMAR2. 0202E+21	sp 075367 H2AY_HUMAN
11784	GSGEKPVAPGDDTESLHSQGEFEFDMPQPPHGHVLR2. 0200000000001E+25	sp Q7Z4V5 HDGR2_HUMAN
11785	NYSSPPPCHLSR2000000000	sp Q12888 TP53B_HUMAN
11786	DKLCSPLSEPGPSK200000000000	sp Q12986 NFX1_HUMAN
11787	EYIPGQPPLSQSSDSSPTR220200000	sp Q5VUA4 ZN318_HUMAN
11788	RLEISPDSSPER200200000	sp P07814 SYEP_HUMAN
11789	TNTPQGVLPSSQLK2000000000000	sp Q99959 PKP2_HUMAN
11790	FLSHSTDLSLNK2000000	sp Q14004 CDK13_HUMAN
11791	GSSPSIRPIQGSQSSSPVEK2200000	sp Q12802 AKP13_HUMAN
11792	SPTPNQDQDGMVEK20000000000000	sp Q92560 BAP1_HUMAN
11793	LIHGEDSDSEGEERSSGSEAGGAGHEEGR2. 020000000022E+27	sp Q3KR37 GRM1B_HUMAN
11794	DKSPGRPLER200000000	sp Q9C0C9 UBE20_HUMAN
11795	HTIIPAKSPEK20000	sp O14578 CTRO_HUMAN
11796	KHSEAEFTPPLKCSPK2000002000000000	sp Q9NZM1 MYOF_HUMAN
11797	KRTSSTCSNESLSVGGTSVTPR2. 22E+19	sp Q3B726 RPA43_HUMAN
11798	ESSEPPAPASSPEAASPTEQGPAGTSK2. 00000022E+25	sp 060343 TBCD4_HUMAN
11799	WQSAFTK200000	sp Q92766 RREB1_HUMAN
11800	CCSPPNELGFR2000000000	sp 075635 SPB7_HUMAN
11801	KHSEAEFTPPLKCSPK2000002000	sp Q9BYW2 SETD2_HUMAN
11802	SGSYGRSPSPYGR20020200000	sp Q3B726 RPA43_HUMAN
11803	TVSFAIEVVPAPPEPPPATPPQAK2. 02E+24	sp Q9NYV4 CDK12_HUMAN
		sp O15047 SET1A_HUMAN

11804	SSTVATLQGTPDHGDPR2000000000000000	sp Q8IVT2	MISP_HUMAN
11805	TPAPEPEPCEASELPAK200000000000000000	sp Q9ULV3	CIZ1_HUMAN
11806	KPVTVSPTTPTSPTEGEAS200000000000000000	sp Q9Y6G9	DC1L1_HUMAN
11807	ISDSESEDPPR20202000000	sp Q96ST2	IWS1_HUMAN
11808	QGSITSPQANEQSVTPQR20000002020000	sp Q9UQ35	SRRM2_HUMAN
11809	EDLSPAFDHSPNK20000020000	sp P17706	PTN2_HUMAN
11810	EGPPAPPPVKPPSPVNIR2000000	sp Q8TF74	WIPF2_HUMAN
11811	HDSSAPDPAPDSADEVMPEAVPEVASEPGLESASHPNVDR2E+29	sp Q9BTC0	DIDO1_HUMAN
11812	EEGSLSDTEADAVSGQLPDPTNPSAGK2.02E+23	sp Q8N1G4	LRC47_HUMAN
11813	TPMYGSQTPLQDGSR2000000200000000	sp 000267	SPT5H_HUMAN
11814	LKSPSQDNTDSYFR2000000000000	sp Q9UJF2	NGAP_HUMAN
11815	SVSKESVASMADSGDDFASDGSSSR20000000000000000000	sp Q92805	GOGA1_HUMAN
11816	GPGQPSSPQR200000	sp 095400	CD2B2_HUMAN
11817	SLSSSPVKK2200000	sp Q9H3Q1	BORG4_HUMAN
11818	EGPSCSPGPF20000000	sp Q4AC94	C2CD3_HUMAN
11819	YSEFTSTSGTGHNQTR200200000000000000	sp P48960	CD97_HUMAN
11820	VLTANSNPSSPSAAK22000000	sp Q9NV56	MRGBP_HUMAN
11821	RSPGPGSPQSPR200000200000	sp Q9P1Y5	CAMP3_HUMAN
11822	GRSPSPIGVGSEMCR20200000001000	sp Q9Y4B5	MTCL1_HUMAN
11823	RLSLDIQSPPNIGLR20000000000000	sp Q9NSI6	BRWD1_HUMAN
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		sp Q9TNN7	1C05_HUMAN
		sp P04222	1C03_HUMAN
		sp Q07000	1C15_HUMAN
		sp Q29865	1C18_HUMAN
		sp Q29960	1C16_HUMAN
11825	GGSCSQAASSNSAQGSDESLIACK2.0002E+20	sp Q29963	1C06_HUMAN
		sp P30504	1C04_HUMAN
		sp P30505	1C08_HUMAN
		sp P30508	1C12_HUMAN
		sp P30510	1C14_HUMAN
		sp Q95604	1C17_HUMAN
		sp P30510	1C14_HUMAN
		sp Q9TNN7	1C05_HUMAN
		sp P04222	1C03_HUMAN
		sp Q07000	1C15_HUMAN
		sp Q29865	1C18_HUMAN
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		sp P30504	1C04_HUMAN
		sp P30505	1C08_HUMAN
		sp P30508	1C12_HUMAN
		sp Q95604	1C17_HUMAN
11827	ASPVGDGNFR2000000000	sp Q86YS7	C2CD5_HUMAN



		sp Q9TNN7 1C05_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29960 1C16_HUMAN
		sp Q29963 1C06_HUMAN
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		sp Q95604 1C17_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
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		sp P30510 1C14_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
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		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P30505 1C08_HUMAN
11831	RNSNSPPSPSSMNQR20200000210000	sp Q7Z5L9 I2BP2_HUMAN
11832	SRSYSPAHR232000000	sp Q9Y2W1 TR150_HUMAN
11833	ISSKSPGHMVILDQTK2200000100000000	sp Q9NWH9 SLTM_HUMAN
11834	VSPAPGSAPETPEDK2000000000200000	sp Q13459 MY09B_HUMAN
11835	SPEQLCAPAEAAPCPAEPER2E+20	sp Q86UD0 SAPC2_HUMAN
11836	TNSGISSASGGSTEPTTPDSERPAQALLR2000000000000000	sp Q86X10 RLGPB_HUMAN
11837	GGEKTPVSGSLKSPVPR20020000000000	sp Q8NI27 THOC2_HUMAN
11838	RESESEDETPPAAPQLIK20200200000000000	sp O60832 DKC1_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
11839	EDALDDSVSSSVHASPLASSPVRK20002000000	sp A6NKT7 RGPD3_HUMAN
		sp O14715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
11840	NHKPSRSFSR2000200	sp P84103 SRSF3_HUMAN
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11843	KMSQPGSPSPK12000200000	sp Q14839 CHD4_HUMAN
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11846	VGSLTPPSSPK22000	sp Q2M2I8 AAK1_HUMAN
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11849	QFIAAQNLGPASGHGTPASSPSSSSLPSPMSPTR200000	sp Q8WX93 PALLD_HUMAN
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11851	HSLSGSSPGMK20000200000	sp Q9UQ35 SRRM2_HUMAN
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11853	VAELALSLSSTSDDEPPSSVSHGAK200000000000000000	sp Q8N1G2 CMTR1_HUMAN
11854	YSPTSPTYSPPTPK20200030002000	sp P24928 RPB1_HUMAN
11855	ETSAATLSPGASSR2002020000000	sp Q9Y6I9 TX264_HUMAN
11856	EKWDCESICSTYSNLYNHPQLIK20002000000000000000	sp Q96GA3 LTV1_HUMAN
11857	TVQSNSPISALAPT GK20000000000000	sp O95292 VAPB_HUMAN

11858	ALGYWGEQDVQEMPAPESPPGR200000	sp Q8NC42 RN149_HUMAN
11859	YSPTSPTYSPVYTPSPK200000000030000000	sp P24928 RPB1_HUMAN
11860	APKPPTDGSTSTPSEDQALGK2000000000000000	sp Q8NHM5 KDM2B_HUMAN
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11862	VDSPSHGLVTSSLCIPSPAR2000000000	sp Q9UER7 DAXX_HUMAN
11863	LMLSTSEYSQSPK202000	sp Q12888 TP53B_HUMAN
11864	GPGGSQGSQSPQGAR20002000000	sp Q96MG7 MAGG1_HUMAN
11865	SEAPNSSEEDSPIKSDK200000200020000000	sp Q8IYH5 ZZZ3_HUMAN
11866	MSCFSRSPSPTPLDR2002100000000	sp Q9UQ35 SRRM2_HUMAN
11867	ESSLPSK2000000	sp 075477 ERLN1_HUMAN
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		sp Q29960 1C16_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp P04222 1C03_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29865 1C18_HUMAN
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		sp P30504 1C04_HUMAN
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		sp P30510 1C14_HUMAN
		sp Q95604 1C17_HUMAN
		sp TAU_HUMAN
11870	SPVVSQDTSR20002000	sp P10636 TAU_HUMAN
11871	NSSLGSPSNLCGSPGSIR200000020000000	sp Q9C0B0 UNK_HUMAN
11872	SDSPVPTAPTSGGPKSTASAVPELATDPELEK2E+33	sp Q86U44 MTA70_HUMAN
11873	GCDSPDPDTSYVLTPTHEEK30020000000	sp Q02078 MEF2A_HUMAN
11874	SSFPDSTEAFDPR2000000000000	sp Q86UL8 MAGI2_HUMAN
11875	HSTASHSSHTSGIEADTKPR2.02022E+19	sp Q96NE9 FRMD6_HUMAN
11876	IFVGGLSPDTPEEK200000000	sp Q14103 HNRPD_HUMAN
11877	SWSASDPFYTNR2000000000000	sp Q9Y2G3 AT11B_HUMAN
11878	SRSPYSPVLR203200000	sp Q14004 CDK13_HUMAN
11879	SPQSDPADTPTNTK200000002000000	sp P46013 KI67_HUMAN
11880	RVSSRSVSGSPEPAK2220202020000000	sp Q8IYB3 SRRM1_HUMAN
11881	SAASPVVSSMPERASESSSEEKDDYEIFVK2.00022100002E+27	sp P18583 SON_HUMAN
11882	KGTTPPRSPEASPK200020000000	sp Q9HDC5 JPH1_HUMAN
11883	TLGTPTQPGSTPR20200002000	sp Q8NFH5 NUP53_HUMAN
11884	NKSPPADPPR200000000	sp 060566 BUB1B_HUMAN
11885	VSELLQGLSFGSDSVEK200200000	sp Q96JS3 PGBD1_HUMAN
11886	SSSELSLSDKSELKK220200000000000	sp Q86X27 RGPS2_HUMAN
11887	EAASSGDDSGRDREPPVQR22000200000000000	sp Q9BW71 HIRP3_HUMAN
11888	RFSLSPSLGPQASR2000200000000	sp Q9NYF3 FA53C_HUMAN
11889	GELSRSPSPFTHTHLAQGYR20202000000000000	sp P38398 BRCA1_HUMAN
11890	SPSRTLSPGR20002020000	sp Q13523 PRP4B_HUMAN
11891	EEEVERPVSSPGDPEQK200000000	sp 043149 ZZEF1_HUMAN
11892	HKSGSMEEDVDTSPGGDYYSPTSSSR1E+25	sp P08651 NF1C_HUMAN
11893	RTGRLGQR20000000	sp Q5HYW3 RGAG4_HUMAN
11894	SFSPKSPLELGEKLSPLPGPGAGDPR2.002000000002E+25	sp Q15742 NAB2_HUMAN
11895	TASPINSARGTSPK20200020000000000	sp Q9BTV5 FSD1_HUMAN
11896	SYSPDGKESPSDK3200000200000	sp P43243 MATR3_HUMAN
11897	AGSSASSPSPASSPHSAVPAADPADSASGSSNK2E+19	sp Q8NDF8 PAPD5_HUMAN
11898	GPSPAPASSPKR20000200000	sp Q9H7N4 SFR19_HUMAN
11899	SRSRSPAHNR2000232000000	sp Q9Y2W1 TR150_HUMAN
11900	GQSQTSPDHRSDTSSPEVR2022200000	sp Q9UQ35 SRRM2_HUMAN

11901	GAPSQDDESGGIEDSPDR20000	sp Q9H4G0 E41L1_HUMAN
11902	SASPYPSHSLSSPQR200000000020000	sp Q5T200 ZC3HD_HUMAN
11903	VEMYSGSDDDDDFNKLPK1300200000000000	sp 060841 IF2P_HUMAN
11904	NVQQDNSEAGTQPQVQTDAQQTSQSPPELTSEENKIPDADK200000000000	sp Q92598 HS105_HUMAN
11905	HLSTPSSVSPEPQDSAK2000000000000	sp Q9NZN5 ARHGC_HUMAN
11906	SSTPGSPTHVSSGSNASR2000000000000000	sp Q14CZ0 CP072_HUMAN
11907	LSTTPSPTSSLHEDGVEDFR202020000000000000	sp 094842 TOX4_HUMAN
11908	SRSISLR20200000	sp Q16629 SRSF7_HUMAN
11909	QLSALSSDGEGQGRPVR2000000000000000	sp 015438 MRP3_HUMAN
11910	RLSSSLNPSK202000000	sp Q13535 ATR_HUMAN
11911	IKEEPVEVDSSPPDSPASSPCSPPLK220000002200200000	sp P17544 ATF7_HUMAN
11912	TPHVQAVQGPLGSPPKR200000	sp Q15149 PLEC_HUMAN
11913	GRNDSGEENVPLDLTR20000000000000	sp Q6R327 RICTR_HUMAN
11914	RGTSPRPPEGGLGYSQLGDDDLK2.00000000003E+21	sp Q9UQ88 CD11A_HUMAN
11915	SQDSYPGSPSLSPR202000	sp P21127 CD11B_HUMAN
11916	LGSLVDEFK20000000	sp Q6VN20 RBP10_HUMAN
11917	IDFTPVSPAPSPTR20000	sp P12956 XRCC6_HUMAN
11918	KPIKSPSKDASSGK20000022000	sp Q7Z309 F122B_HUMAN
11919	HLDVDLDRQLSSIDKNPSEK20200000000000	sp Q13523 PRP4B_HUMAN
11920	NPSVVIKPEACSPQFGK2000000	sp Q9Y6D5 BIG2_HUMAN
11921	SRSRSDYNYR2020200000000	sp Q8WXE1 ATRIP_HUMAN
		sp 075494 SRS10_HUMAN
		sp Q29960 1C16_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
11922	GGSCSQAASSNSAQGSDESLIACK2E+20	sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
		sp P30510 1C14_HUMAN
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11924	EPAPLLSAQYETLSDSD2000	sp 075376 NCOR1_HUMAN
11925	ASQSRPNSSALETLGGEK2000020000000000	sp 094804 STK10_HUMAN
11926	AQGEPVAGHESPKIPYEK200000000	sp 094979 SC31A_HUMAN
11927	SRSQSRNSPLPVPSK2020202000000000	sp Q13247 SRSF6_HUMAN
11928	YTPTSPSYSPSSPEYTPTSPK300020002000000	sp P24928 RPB1_HUMAN
11929	CPEILSDESSDEDEKKNK2002220000000000	sp P35659 DEK_HUMAN
11930	VTDKDCNESTDNDELEDEPEEPFYR220000000000000000	sp Q9Y2K1 ZBTB1_HUMAN
11931	AADISLDNLVEGK2000000000	sp Q12888 TP53B_HUMAN
11932	EMEHNTVCAAGTSPVGEIGEEK1.0002E+21	sp P18583 SON_HUMAN
11933	ETNLDLPLVDTHSK2000000000000000	sp P08670 VIME_HUMAN
11934	DFTNEAPPAPLPDASASPLSPHR20000	sp Q6WCQ1 MPRIP_HUMAN
11935	SLSNTPDYCHDK200000000000	sp Q8NG31 CASC5_HUMAN
11936	SESAPTLHPYSPLSPK200000000000000000	sp Q8WUF5 IASPP_HUMAN
11937	SRSSVGSSSSYPIPAVSR2220002000000	sp Q15149 PLEC_HUMAN
11938	SAPASNAGVLSAHSSGAQTPELSR2.000200000002E+27	sp Q01167 FO XK2_HUMAN
11939	YMNSDTSPELR100000200000	sp Q96ST3 SIN3A_HUMAN
11940	SRDSFLK20000000	sp P46778 RL21_HUMAN
11941	VAFDPEQKPLHGVLTPTSSPASSPLVAK200000022000000	sp Q14684 RRP1B_HUMAN
11942	STPHRTPITPNTK20002000200000	sp Q9NQW6 ANLN_HUMAN

11943	TSPCQSPTISEGPVPSLPIRPSR2. 2E+23	sp 043426 SYNJ1_HUMAN
11944	RESESEDETPPAAPQLIKK2000200000000000000	sp 060832 DKC1_HUMAN
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11946	FFCSDSTDCVSNK20000000000	sp Q9UQ84 EXO1_HUMAN
11947	GSGVAQSPQQPPPPQQQQQPPQQPTPPK2E+22	sp Q6VY07 PACS1_HUMAN
11948	ASPSKPASAPASR20000000000	sp P27816 MAP4_HUMAN
11949	AAIELPSLEVLSDQEEDREQCGVK200000000000000	sp Q9BQI3 E2AK1_HUMAN
11950	NSDSIVSLPQSDRSSSSQESLNRPLSAK2. 00020022E+23	sp Q7Z460 CLAP1_HUMAN
11951	ELSPLISLSPVPPLSPIHSNQQTLPR2. 002000002E+21	sp P51825 AFF1_HUMAN
11952	FNSLPQYSPVTFDR20000000	sp Q8WY36 BBX_HUMAN
11953	GHVTPNLSR2000000	sp Q96CP6 GRM1A_HUMAN
11954	KPPAPSPVQSQSPSTNWSPAVPVK2. 000002022E+19	sp Q8IYB3 SRRM1_HUMAN
11955	GYTSDSEVYTDHGRPGK2000000000000	sp Q92538 GBF1_HUMAN
11956	LSEEAECNPSTPSK200000	sp 094804 STK10_HUMAN
11957	SITSTPLSGK2002000	sp Q9P0K7 RAI14_HUMAN
11958	LSQMQSPIPLTETR102000000000	sp Q2VPK5 CTU2_HUMAN
11959	TRHSPTPQQSNR2020000000	sp Q8IYB3 SRRM1_HUMAN
11960	EGMNPYSYDEYADSDDEDQHDAYLER1. 0000003E+22	sp Q08945 SSRP1_HUMAN
11961	GVAPADSPEAPR2000000	sp Q9P2G1 AKIB1_HUMAN
11962	TGVTSTSDSEEGDDQEGEK2000020000000000000	sp 075475 PSIP1_HUMAN
11963	LPSSPASPSPK2200000000	sp Q9H4G0 E41L1_HUMAN
11964	EIEMSVDDDDINSSK1000000000200	sp Q9NQW6 ANLN_HUMAN
11965	SLRGSDALSETSSVSHIEDLEK2000200000000000000	sp Q6P996 PDXD1_HUMAN
11966	VSPASPAGSPSADFAVHGESLGDR2E+23	sp Q96C92 SDCG3_HUMAN
11967	DGSGTPSRHSLSGSSPGMK200000000002200000	sp Q9UQ35 SRRM2_HUMAN
11968	SEESSDDESHPLR22000000000	sp Q9UQ84 EXO1_HUMAN
11969	DDSHSAEDSEDEKEDHK20200000000000000	sp Q9H1E3 NUCKS_HUMAN
11970	QHEAPSNRPLNELLTPQGSPR200000000000002000	sp P10398 ARAF_HUMAN
11971	SQPSPVTVKPKPPAR2002000000000000	sp Q96AY4 TTC28_HUMAN
11972	TPNDFSEAPSSPSAQDHQPTLGLACSQLHSPQHK2. 02E+25	sp Q6T4R5 NHS_HUMAN
11973	RVSPSPPR2020000	sp Q5VUA4 ZN318_HUMAN
11974	DGSDYDPYDFSDTEEMPQVHTPK3. 00000000001E+20	sp P25205 MCM3_HUMAN
11975	SCTPSPDQISHRASLEDAPVDDLTR2. 000020002E+21	sp Q7Z2W4 ZCCHV_HUMAN
11976	EEATYGVWAERDSDDERPSFGGK2E+20	sp Q9UBB9 TFP11_HUMAN
11977	DTPGHGSGWAETPR20000000002000	sp 075533 SF3B1_HUMAN
11978	AKSPAPGLASGDRPPSVSSVHSEGCNR2. 0000002E+26	sp Q9Y618 NCOR2_HUMAN
11979	DGTAPPPQSPGSPGTGQDEEWSDEESPRK2000000000000000000	sp Q9UKS6 PACN3_HUMAN
11980	ALEQQAESDSSDTEEKDDDDDEEDVGKR2. 02002E+22	sp Q9BXY0 MAK16_HUMAN
11981	SPAAPGQSPTGVCYER2000000000000000000	sp 075534 CSDE1_HUMAN
11982	SLSGTVVSR200000000	sp Q70EL1 UBP54_HUMAN
11983	GTPPLTPGR200000000	sp Q9Y3Q4 HCN4_HUMAN
11984	SPAPGLASGDRPPSVSSVHSEGCNR2. 000000000002E+26	sp Q9Y618 NCOR2_HUMAN
11985	SFSPEER2020000	sp Q9UDY2 ZO2_HUMAN
11986	SAPASPTHPLMSPR202000010000	sp P85037 FOXK1_HUMAN
11987	DGQDAIAQSPEKESK20000000	sp Q9NWH9 SLTM_HUMAN
11988	SAPASPTHPLMSPR2000002000010000	sp P85037 FOXK1_HUMAN
11989	NSSTETDQQPHSPDSSSVHSIR2E+21	sp Q92613 JADE3_HUMAN
11990	AGLESGAEPGDGSDTTK200000000000200	sp 060832 DKC1_HUMAN
11991	LLAASGSNSPTR20000	sp Q9NS37 ZHANG_HUMAN
11992	QVVISPQVSR2000200	sp Q6ZQY3 GADL1_HUMAN
11993	DTSQSDKDLDDALDKLSDSLGQR20000000	sp P20810 ICAL_HUMAN
11994	GHPAQRPEPSPPEPEPQAPEGSQAGAEGSSPEASRSPAR2. 000000000000 2E+19	sp Q6ZW31 SYDE1_HUMAN
11995	QQGHEPVSPR2000	sp Q9BTA9 WAC_HUMAN

11996	RSSSSGDQSSDSLNSPTLLAL2E+20	sp P15408 FOSL2_HUMAN
11997	GLSLEHQK2000000	sp Q765P7 MTSSL_HUMAN
11998	NSTDSRPVSVTYR200000000000	sp Q8TBZ3 WDR20_HUMAN
11999	AVRPEVTVASSDEVCDGDR20000000000	sp Q75151 PHF2_HUMAN
12000	GSSGSPAHAESYSSGGGGQK2.000000003E+20	sp P15408 FOSL2_HUMAN
12001	MSQPGSPSPK2000002000	sp Q14839 CHD4_HUMAN
12002	SRSRHSSISPSTLTLK20200220200000000	sp Q14004 CDK13_HUMAN
12003	DGSGTPSRHSLSGSSPGMK200000020202200000	sp Q9UQ35 SRRM2_HUMAN
12004	EHHPEEGSSGSEVEEIPETPCESQGEELK2.202E+22	sp Q12888 TP53B_HUMAN
12005	KQGEDNSTAQDTEELEK200000000000	sp Q86UE4 LYRIC_HUMAN
12006	TASRPDDIPDSPSSPK2000000000002000	sp Q5VZK9 LR16A_HUMAN
12007	AGAPGALSPSYDGGLHGLQSK300000000000	sp P15923 TFE2_HUMAN
12008	GGAPDPSPGATATPGAPAQSSPDAR200000000200000	sp Q095365 ZBT7A_HUMAN
12009	SRTSPAPWK2000000	sp Q9UQ35 SRRM2_HUMAN
12010	EQSTPPKSPQ20002000	sp Q92615 LAR4B_HUMAN
12011	YTPTSPSYSPSSPEYTPTSPK2.022E+20	sp P24928 RPB1_HUMAN
12012	DGTAPPQSPGSPGTGQDEEWSDEESPRK200020000	sp Q9UKS6 PACN3_HUMAN
12013	VKIIPTPRSK200	sp Q14703 MBTP1_HUMAN
12014	YADEEIPRSPFK20000	sp Q075369 FLNB_HUMAN
12015	SYSRSPPYPR23200000000	sp Q07Z6E9 RBBP6_HUMAN
12016	LVKPSSSSSGSR200000	sp Q09H6B4 CLMP_HUMAN
12017	NDSGDHSDHMHYQK2000000000000000	sp P78332 RBM6_HUMAN
12018	SPSPGNHTSNSSNASNATVVPQNSSAR2.2E+20	sp Q09BTA9 WAC_HUMAN
12019	APKPPTDGSTSPTSTPSEDQALGK2E+20	sp Q08NHM5 KDM2B_HUMAN
12020	HSTPSNNSNPSGPPSPNSPHR2E+20	sp Q09Y5S2 MRCKB_HUMAN
12021	SPSESSPQHPTTPPARPR2000002000020000000	sp Q014733 MP2K7_HUMAN
12022	TEMKSPFNPSPQDSR200020200000000	sp P08651 NFIC_HUMAN
12023	EEASLLSHSPGTSNQSQPCSPKPIR2E+22	sp Q09H4Z3 PCIF1_HUMAN
12024	IDFSSIAVPGTSSPR2000	sp Q05TDH0 DDI2_HUMAN
12025	WDCESICSTYSNLYNHPQLIK200023200000000000	sp Q096GA3 LTV1_HUMAN
12026	QGRPSPTSPVKPSSPASKPDGPAELPLTDR2.02E+26	sp Q013905 RPGF1_HUMAN
12027	AGEAPTENPAPPTQSSAE20002000	sp P16989 YBOX3_HUMAN
12028	SRSHSDNDRFK202020000000	sp Q075494 SRS10_HUMAN
12029	DTHEDHDTSTENTDESNHDPQFEPIVSLPEQEIK2.2E+26	sp P43487 RANG_HUMAN
12030	IVPQSQVNPESPCK20000	sp Q09ULM3 YETS2_HUMAN
12031	RESCGSSVLTDFEGK200000000000000	sp Q015231 ZN185_HUMAN
12032	NIGRDTPTSAGPNSFNK2000000000000	sp Q08WW12 PCNP_HUMAN
12033	AGGPTTPLSPTR20000	sp P20700 LMNB1_HUMAN
12034	NHKPSRFSR200020200	sp P84103 SRSF3_HUMAN
12035	SPSPLRGNVPSPLPTR202000000002000000	sp Q09Y2V2 CHSP1_HUMAN
12036	DYGNSPLHR200000	sp P26599 PTBP1_HUMAN
12037	TPSVSAPLALSCPR2000000000000	sp Q09UGP4 LIMD1_HUMAN
12038	RLSLGQGDSTEAAATEER2000000000000000	sp Q06ZRV2 FA83H_HUMAN
12039	SLSEYELSSDLQDK200000000000000	sp Q06DN90 IQEC1_HUMAN
12040	ENSPAVSPTTNSTAPFGLKPR2.000000002E+19	sp Q09Y2L9 LRCH1_HUMAN
12041	LDSEMDSSENYDTMSSPLPGK2.2E+21	sp Q09NTI5 PDS5B_HUMAN
12042	TPTSTPVHVK200000000	sp Q09ULM3 YETS2_HUMAN
12043	DDGDEEGLLTHSEEELEHSQDTDADDGALQ2.000000002E+19	sp Q09UKM9 RALY_HUMAN
12044	SRSISRPR202020000	sp Q016629 SRSF7_HUMAN
12045	RHSSVSDSQPCEPPSVGTEYSQGASPPQHQLK2.2E+31	sp P39880 CUX1_HUMAN
12046	LDQPGNLPGSNRR20000	sp P46013 KI67_HUMAN
12047	STFVQSPADACTPPDTSSASEDEGLR2220200000000	sp Q09P265 DIP2B_HUMAN
12048	HQPWQSPERPLSR200000000	sp Q09BQ52 RNZ2_HUMAN
12049	KPGPPLSPEIRSPAGSPELR200000000200000	sp Q096JM3 CHAP1_HUMAN

12050	ASLVALPEQTASEEETPPPLLTK20000000000200	sp Q9UKV3 ACINU_HUMAN
12051	EDSAPSSVAR2000020000	sp Q9BZE4 NOG1_HUMAN
12052	SLSRTPSPPPFR20202000000	sp Q7L4I2 RSRC2_HUMAN
12053	ELESPLTPGK20020000	sp O15014 ZN609_HUMAN
12054	AEGHMGSEPR1020000	sp Q9P2N6 KANL3_HUMAN
12055	AISSDDSSDEDK22000220000000	sp Q6PD62 CTR9_HUMAN
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12057	HSSTPNSSEFSR2000000000	sp Q8N9Q2 SR1IP_HUMAN
12058	EYVNSTSEESHDEDEIRPVQQDLHR200000000000000000	sp Q8NBU5 ATAD1_HUMAN
12059	EGSTQQLQTTSPK2000	sp Q5T8P6 RBM26_HUMAN
12060	DVDVSEDSPPPLPER200000000000	sp Q05209 PTN12_HUMAN
12061	IHLGSSPK2000	sp Q9H410 DSN1_HUMAN
12062	VQIPVSRPDPVSDNEEDSYDEEIHDP200000000000	sp Q07157 ZO1_HUMAN
12063	SSSVGSSSYPIPAVSR22200000000000000000	sp Q15149 PLEC_HUMAN
12064	LSTIQNSPTK20000	sp Q5VYV7 SLX4I_HUMAN
12065	KPPAPSPVQSQSPSTNWSAPVVK220000000000	sp Q8IYB3 SRRM1_HUMAN
12066	HPSGGSSPPR2000200000	sp Q6IQ23 PKHA7_HUMAN
12067	MISTPSPK2000	sp Q53F19 CQ085_HUMAN
12068	SQSLQDPTR200000000	sp Q8N4C8 MINK1_HUMAN
12069	SPGTTPRPPGASK2002000000000	sp Q9Y3X0 CCDC9_HUMAN
12070	RESTGSYDLYPDSDDSEDQVENS3. 2000002002E+20	sp Q70CQ2 UBP34_HUMAN
12071	GSASPHKSSPQLVATPSQHHQQQQQIK2. 0002E+26	sp Q9POU4 CXXC1_HUMAN
12072	VPGSSGHLHK22000000	sp Q9UEW8 STK39_HUMAN
12073	RTPSDDEEDNLFAPPK2020000000000000	sp Q641Q2 FA21A_HUMAN
12074	SSITEPEGPNPNIQK20000000000000	sp Q9Y4E1 FA21C_HUMAN
12075	LGTLAQLHASHCLDDLPLSTPSSPR20020000	sp Q13625 ASPP2_HUMAN
12076	FTSSQTQSTPPKPPSPSFLGLSSFPPLPGAAGNLK2. 0000022E+20	sp Q68DK2 ZFY26_HUMAN
12077	GSPPVPSGPPMEEDGLR200000000000000000	sp Q92615 LAR4B_HUMAN
12078	KSSTVATLQGTDPDHGDP20000000000000	sp Q9H4L4 SENP3_HUMAN
12079	SKPAPSESAGYGIPLKDGDEKTDEEAEGPYSDNEMLTHK2E+32	sp Q8IVT2 MISP_HUMAN
12080	DTHPGSPGSPALRPLPAR20000000000000	sp Q6NUK4 REEP3_HUMAN
12081	SHMSGSPGPGSNTAPSTPVIIGSDKPGMEEK1. 00000000202E+30	sp Q8N0Y2 ZN444_HUMAN
12082	TVTPASSAKTSPAK220000	sp A0FGR8 ESYT2_HUMAN
12083	SGSMEEDVDTSPGGDYTSPSSPTSSSR3000020000000	sp Q16555 DPYL2_HUMAN
12084	GLGTDEDSLIEIICSR2000000000	sp P08651 NFIC_HUMAN
12085	GSPPVPSGPPMEEDGLR20000000010000000	sp P07355 ANXA2_HUMAN
12086	EGAASPAPETPQPTSPETSPK2000000002000	sp A6NMY6 AXA2L_HUMAN
12087	YNLDASEEEDSNKK20000	sp Q9H4L4 SENP3_HUMAN
12088	SGNSDSIER2000000000	sp Q9Y3X0 CCDC9_HUMAN
12089	GSPTDAEGSPGLSR2000000	sp O95218 ZRAB2_HUMAN
12090	SPSRPILDGK20000000000	sp Q7Z2E3 APTX_HUMAN
12091	KVYEDSGIPLPAESPK3002000000000000	sp Q8TER5 ARH40_HUMAN
12092	SHSPSPDPDTPSPVGD2000000000	sp O60890 OPHN1_HUMAN
12093	GSSQPNLSTSHSEQEYK200000000000000000	sp Q8IXM2 BAP18_HUMAN
12094	AKSSDSDSSEDEPPKNQPK2. 20202022E+21	sp Q13586 STIM1_HUMAN
12095	SRSPSPELNNK2000220000000	sp O95208 EPN2_HUMAN
12096	VQMTSPSSTGSPMFK2000000001000	sp Q14978 NOLC1_HUMAN
12097	RESESEDETPPAAPQLIK202000020000000000	sp Q9UQ35 SRRM2_HUMAN
12098	LAEIEEEKPEPDISSEESVSTVEEQENETPPATSSEAEQPK2. 200000002E+22	sp P49790 NU153_HUMAN
12099	MFGGPGTASRPSSSR20000	sp O60832 DKC1_HUMAN
12100	EASAPLAQGEPPGREDLPT200000000000000000	sp Q5SSJ5 HP1B3_HUMAN
		sp P08670 VIME_HUMAN
		sp Q9P1Y6 PHRF1_HUMAN

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12102	SSTPPRQSPSR20000020000	sp Q9UQ35 SRRM2_HUMAN
12103	STSAPQMSPGSSDNQSSSPQPAQQK2200020000000000	sp Q14157 UBP2L_HUMAN
12104	TVSSPPTSPRPGSAATVSASTSNIIPPR220000000	sp 076094 SRP72_HUMAN
12105	DNSPEPNDPEEPQEVSSSTPSDKK2E+21	sp Q9HB58 SP110_HUMAN
12106	STSPAGQHHSPISSR2220000000002000	sp Q5T200 ZC3HD_HUMAN
12107	TFNSPNLK200000	sp Q9Y6N7 ROBO1_HUMAN
12108	KLSLPTDLKPDLDVK2000000000000000	sp Q5T5C0 STXB5_HUMAN
12109	VDENENCSLGLSPSEPPQTLDLVR200000000000	sp Q96P16 RPR1A_HUMAN
12110	DQRPSPIR20000	sp Q5T200 ZC3HD_HUMAN
12111	STSMDSGSSESPASLK2000000000000000	sp Q15811 ITSN1_HUMAN
12112	RSSSPFLSK22000000	sp Q9NYV4 CDK12_HUMAN
12113	SYSPDGKESPSDKK2000000020000	sp P43243 MATR3_HUMAN
12114	RSSQPSPTAVPASDSPPTKQEVK2020000200000000000	sp Q3KQU3 MA7D1_HUMAN
12115	SGGLQTPECLSR2000020000200	sp P85037 FO XK1_HUMAN
12116	SLENETLNKEEDCHSPTS KPPKPDQPLK2E+28	sp P23588 IF4B_HUMAN
12117	SCDEPLTPPHSPTSMQLIHDVSPR221000000002000	sp Q9Y2K7 KDM2A_HUMAN
12118	SLSPGKENVSA LDMEK20000000020000000	sp Q9H307 PININ_HUMAN
12119	REESPMVDVQSPSAQDTQSIASDGT PQGEK1. 00000000000202E+26	sp Q7Z6Z7 HUWE1_HUMAN
12120	TVSPGVSPIHGQGVVENLK20000000000000000	sp Q9NZM3 ITSN2_HUMAN
12121	MVEPENAVTITPLRPEDDYSR1E+22	sp Q14160 SCRIB_HUMAN
12122	SQDSYPGSPSLSPR200200020002000	sp Q6VN20 RBP10_HUMAN
12123	SSSMSSLTGAYTSGIPSSSR2. 001E+20	sp A8CG34 P121C_HUMAN
12124	SLRGSDALSETSSVSHIEDLEK2000202220000000000	sp Q6P996 PDXD1_HUMAN
12125	RSTLMSSPR122000	sp Q8NBR9 CK072_HUMAN
12126	TPESKPTILVK2000000000000	sp Q9Y3B9 RRP15_HUMAN
12127	SEDEDSLEEAGSPAPGPCPR20000020000000000	sp Q8TBB5 KLDC4_HUMAN
12128	KHSPSPPPPTPTESR2000000000200	sp Q92922 SMRC1_HUMAN
12129	AQPFGFIDS DTAEEERIPATPVVIPMK2. 02E+20	sp Q14676 MDC1_HUMAN
12130	TVNSTRET PPK20000000	sp Q5SSJ5 HP1B3_HUMAN
12131	KYSISSDNSDTTDSHATSTSASR2022000000000000	sp Q9NQC1 JADE2_HUMAN
12132	DYGHSSSRDDYPSR20000000000	sp Q96E39 RMXL1_HUMAN
12133	TSSSETEEKKPEKPLILQSLGPK2. 2E+21	sp P38159 RBMX_HUMAN
12134	SPILLPK20000000	sp Q9Y5K6 CD2AP_HUMAN
12135	VNPVTSLENYTCSDSEESSEKDK20000000000000000	sp Q8TD16 BICD2_HUMAN
12136	GPGGSQGSQGPSPQGAR2000000000000000	sp Q14432 PDE3A_HUMAN
12137	EEAPASPLRPLYPQISPLK2000000000000000	sp Q96MG7 MAGG1_HUMAN
12138	GSPSVAASSPPAIPK200000020000000	sp P85037 FO XK1_HUMAN
12139	SVCGHLENTSVGNPNSAENSFR20002000220000000	sp Q92610 ZN592_HUMAN
12140	SAASPVVSSMPERASESSSEEKDDYEIFVK2. 21E+23	sp Q96HH9 GRAM3_HUMAN
12141	EENVDGQEVAPSSPPLAAQSPEGNHGHQEPASTSFHPPR2. 0000002E+27	sp P18583 SON_HUMAN
12142	LHDFLAHSSESEETSSPPR20000	sp Q9NWS9 ZN446_HUMAN
12143	NSSTETDQQPHSPDSSSVHSIR2000000000000000000	sp P46100 ATRX_HUMAN
12144	TPNDFSEAPSSPSAQDHQPTLGLACSQLHSPQHK20000200000	sp Q92613 JADE3_HUMAN
12145	HPSGGSSPPPR2200000	sp Q6T4R5 NHS_HUMAN
12146	SGGLQTPECLSR2000000000200	sp Q6IQ23 PKHA7_HUMAN
12147	AASADSTTEGTPADGFTVLSTK2E+20	sp P85037 FO XK1_HUMAN
12148	TRSWDSSSPVDRPEPEAASPTR2. 002E+21	sp P85037 FO XK1_HUMAN
12149	DMPMDISALSSPSCSPSPQSETPLEK2000000000000	sp 075367 H2AY_HUMAN
12150	SFSISPSRR2020000000	sp Q86WB0 NIPA_HUMAN
12151	ASPVTSPAAAFPTASPANK2000000000200000	sp Q8NDZ2 SIMC1_HUMAN
12152	RGSQSEDESCSLHSQTLSEDER2. 02E+20	sp P18583 SON_HUMAN
		sp Q9UIF9 BAZ2A_HUMAN
		sp Q9UPZ3 HPS5_HUMAN

12153	SKSYSPGRR2020200000	sp Q8IWX8 CHERP_HUMAN
12154	SQLPTLEQDGGTQNPVSSPGMSQELR2001000000	sp P46937 YAP1_HUMAN
12155	EVYQQQYQYSGGR20000	sp Q99729 ROAA_HUMAN
12156	EVEEDESEDEEMSEDEEDDSSGEEVVIPQKK2.00001000000022E+25	sp P19338 NUCL_HUMAN
12157	RPNTSPDR20000	sp Q86VM9 ZCH18_HUMAN
12158	YTPTSPSYSPSSPEYTPSPK3.2E+21	sp P24928 RPB1_HUMAN
12159	RYSPPIQR30000000	sp Q8IYB3 SRRM1_HUMAN
12160	ENQGSHESTLNPETVAINLSDVDLSK2000000000000000000	sp P25445 TNR6_HUMAN
12161	DDSHSAEDSEDEKEDHKNR20000020000000000000	sp Q9H1E3 NUCKS_HUMAN
12162	VDSPLPSDKAPTPPGK200000000200000	sp Q12893 TM115_HUMAN
12163	QKSPEIHR2000000	sp Q9NYF8 BCLF1_HUMAN
12164	ASSPSASSSSFR200000000000	sp Q8IXZ2 ZC3H3_HUMAN
12165	KYLHRPSR200	sp Q8TBK6 ZCH10_HUMAN
12166	SGSSQELDVKPSASPQER22000000020000000	sp Q9UQ35 SRRM2_HUMAN
12167	NSNFSVQHPSSTSPTEK200000000000000	sp Q03468 ERCC6_HUMAN
12168	GCLTTPNSPSMHSR20010000	sp 075128 COBL_HUMAN
12169	LLSSNEDDANILSSPTDR200000	sp 075448 MED24_HUMAN
12170	SRSPGSPVGEETGSPPK20020000000000000	sp P35611 ADDA_HUMAN
12171	HASSPESPKPAPAPGSHR20000000000000000	sp Q9UQ35 SRRM2_HUMAN
12172	ISDSESEDPPRHQASDSENEELPKPR2.0200000000002E+25	sp Q96ST2 IWS1_HUMAN
12173	ARSPDLGPQEQMNP200000000000000	sp Q8IWX8 ZSC29_HUMAN
12174	DRPGSPESPLLDAPFSR20000000000	sp Q8TF72 SHRM3_HUMAN
12175	RSSLSSHSHQSIYR2200202000000000	sp 015027 SC16A_HUMAN
12176	SRTHSTSSSLGSGESPFSR2.02E+19	sp Q9UGV2 NDRG3_HUMAN
12177	MGQAPSQSLPPAQDQPRSPVPSAFSDQSR1E+30	sp Q9UQ35 SRRM2_HUMAN
12178	TSGAGSPQTPPER20000000200000	sp Q96MH2 HEX12_HUMAN
12179	TVSASSTGDLPK220000000	sp P78362 SRPK2_HUMAN
12180	EFSFEAWNAK200000000	sp P78347 GTF2I_HUMAN
12181	RVSSSGKPTSLK22000000000	sp Q9ULD2 MTUS1_HUMAN
12182	GTVPDDAVEALADSLGK20000	sp P20810 ICAL_HUMAN
12183	QKSQTPPRR20200000	sp Q9UQ35 SRRM2_HUMAN
12184	KRESESEDETTPAAPQLIK20020000000000	sp 060832 DKC1_HUMAN
12185	SPPKSPEEEGAVSS2000000000000000	sp Q01130 SRSF2_HUMAN
12186	GPSQEEEDNHSDEEDRASEP2E+19	sp Q86VM9 ZCH18_HUMAN
12187	LSVGEQPDVTHDPYEFLLQSEPAASAK2000000000	sp Q9H8M2 BRD9_HUMAN
12188	QIPAAASLIQTAWR200000000	sp P51787 KCNQ1_HUMAN
12189	HEDLVPPAGSPELSPQPLFRPR200020000000000	sp Q9UPS6 SET1B_HUMAN
12190	RPGAPSLSPAPR200000	sp Q5QP82 DCA10_HUMAN
12191	ISPLSSPCSSPLQGTPASSLVSK2.0022E+22	sp Q14432 PDE3A_HUMAN
12192	WLKSPTPIDPEK20000000000	sp Q08AD1 CAMP2_HUMAN
12193	SQSLNIR200000	sp 000443 P3C2A_HUMAN
12194	MPLLELGGETTPPLSTERSPEAVGSECPSR2.2E+21	sp Q9Y4K1 AIM1_HUMAN
12195	MALEVGMEDGQLSDSDMTVAPSDRPLQLPK1.0000001000002E+33	sp Q9H814 PHAX_HUMAN
12196	VKEEIKSDSSPLPSEK20220002000	sp 014646 CHD1_HUMAN
12197	SRSRSTPPSSAGLGSNSAPPIDSR220000202000000000	sp Q8IWX8 CHERP_HUMAN
12198	APELNTSSPSLK200000	sp Q9UN79 SOX13_HUMAN
12199	HSLASSEYPVR20030000	sp P51570 GALK1_HUMAN
12200	PGPTPSGTNVGSSGRSPSK200000000	sp P60468 SC61B_HUMAN
12201	LEGSDVDSELEDRVDGK2000200000000000	sp Q92614 MY18A_HUMAN
12202	DYTGCTSESLSPVK2000020000	sp 095297 MPZL1_HUMAN
12203	SQRSNSPLPVPPSK2020200000000000	sp Q13247 SRSF6_HUMAN
12204	GSLPQLVSPIVSR200	sp 075970 MPDZ_HUMAN
12205	LRKDLFSSR2200	sp Q9BWF2 TRAI1_HUMAN



12206	SPPGPAGSSPK200000002000	sp	Q9H9D4	ZN408_HUMAN
12207	NDKSEEEQSSSVK220000	sp	P07910	HNRPC_HUMAN
12208	TPTSSPASSPLVAK20000000	sp	Q14684	RRP1B_HUMAN
12209	HISECFPSPGGQLAEVHSVSPEQGA20000000	sp	Q5T1V6	DDX59_HUMAN
12210	TPTSPLK220000	sp	Q06264	SMCA5_HUMAN
12211	SKSYSPGRR2023000000	sp	Q8IWX8	CHERP_HUMAN
12212	GHYTEGAELVDAVLVDVVR3200000000000000	sp	Q9BUF5	TBB6_HUMAN
12213	GEAVLRPGLDAEPELSPEEQ2000000	sp	P04350	TBB4A_HUMAN
12214	SRSPLDKDTYPPSASVVGASVGGHR300002000000000000	sp	Q9BRJ6	CG050_HUMAN
12215	SLDGAEFSPASVSENHDAGPDGDKR2020000000000000	sp	Q96T37	RBM15_HUMAN
12216	EKSATRPSPPER200002020000	sp	Q9H4G0	E41L1_HUMAN
12217	TPNNVSTPAPSPDASQLASSLSSQK2. 2E+20	sp	Q9UQ35	SRRM2_HUMAN
12218	RISSSSVQPCSEEVSTPQDSLAAQCK2E+22	sp	Q8WWI1	LM07_HUMAN
12219	QRLATFFPRK2000000	sp	P42694	HELZ_HUMAN
12220	NGLPGSRPGSPER200020000	sp	Q9Y5P3	RA12_HUMAN
12221	STSAPQMSPGSSDNQSSSPQAQQK1. 000000002E+19	sp	Q92993	KAT5_HUMAN
12222	YIASVQGSTPSPR20000002000	sp	Q14157	UBP2L_HUMAN
12223	SRSLSSPR202020000	sp	P49792	RBP2_HUMAN
12224	HSPATISSSTR20000000000	sp	Q9UQ35	SRRM2_HUMAN
12225	AASPESASTPESLQAR2002002200000000	sp	Q015119	TBX3_HUMAN
12226	AKSPAPGLASGDRPPSVSSVHSEGDCNR2E+26	sp	Q96BT3	CENPT_HUMAN
12227	SGSMEEDVDTSPGGDYITSPSSPTSSSR20000000	sp	Q9Y618	NCOR2_HUMAN
12228	ESSPFLSPLEASR200000000000	sp	P08651	NF1C_HUMAN
12229	SKPNLPSESRSR2000200	sp	Q9UP95	S12A4_HUMAN
12230	SDILKDPPEANSIQSANATK2000200200000000	sp	Q08170	SRSF4_HUMAN
12231	IKEEPEVVDSSPPDSPASSPCSPPLK220002000200000000	sp	Q8N488	RYBP_HUMAN
12232	SLPLISASR2000000000	sp	P17544	ATF7_HUMAN
12233	VQSTADIFGDEEGDLFK2000000000000000	sp	Q5VV41	ARHGG_HUMAN
12234	TGGSSPEALHRPYGCDVEPQALNEAIR2E+23	sp	Q641Q2	FA21A_HUMAN
12235	LHSYSSPSTK200200000	sp	Q5SNT6	FA21B_HUMAN
12236	NALFPEVFSPTPDENSQNSR200000000000	sp	Q9Y4E1	FA21C_HUMAN
12237	LSLQHTQQNADGQEDGESERNSNSSGDPGR2000000000	sp	P42684	ABL2_HUMAN
12238	DPNSATATAPPSPLK20000	sp	Q96T37	RBM15_HUMAN
12239	YSPKTPPR20020000	sp	Q8IVT2	MTSP_HUMAN
12240	ASQPDLDVTPTSSKPQPK200000020000000000	sp	Q075052	CAPON_HUMAN
12241	DSAGVPAPAPDSALDSAPTASAPAPAPALAQAPALSPSLASAPPEAK200000000000	sp	Q92766	RREB1_HUMAN
12242	SRSSSVGSSSYPISPAISR22200220000000000000	sp	P49792	RBP2_HUMAN
12243	LSDCEGSVHCEVPSEYACEGKPPDPDSEGSTK2002000	sp	Q7Z3J3	RGPD4_HUMAN
12244	EKPCVSPTVGEK200000	sp	A6NKT7	RGPD3_HUMAN
12245	GGHSSVSTESESSSFHSS200000000000	sp	PODJD1	RGPD2_HUMAN
12246	GVQKPAGPSTSPDGNSR20000000	sp	PODJD0	RGPD1_HUMAN
12247	YSPSPPKR202000000	sp	P46013	KI67_HUMAN
12248	CLLADLPLPELPGDDLSKSPEEK20000000	sp	Q7Z2T5	TRM1L_HUMAN
12249	TRMTELSR201000000	sp	Q15149	PLEC_HUMAN
12250	RQSSSYDDPWK23000000	sp	Q9NSI6	BRWD1_HUMAN
12251	SSLSSHQSQIYR2202200000000000	sp	Q68DQ2	CRBG3_HUMAN
		sp	P61073	CXCR4_HUMAN
		sp	Q86YV5	SG223_HUMAN
		sp	Q8IYB3	SRRM1_HUMAN
		sp	Q14004	CDK13_HUMAN
		sp	Q8IVH4	MMAA_HUMAN
		sp	Q96D71	REPS1_HUMAN
		sp	Q015027	SC16A_HUMAN

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12253	ITSPLMEPSSIEK2000100000000	sp P28066 PSA5_HUMAN
12254	SDGAPASDSKPGSSEAAPSSK2000000000000000	sp P80723 BASP1_HUMAN
12255	TDSEKPPFRGSQSPKR20000000020000	sp Q9Y2W1 TR150_HUMAN
12256	SSPASSPELPWRPR2200220000000000	sp Q9NYF3 FA53C_HUMAN
12257	NATQPPNAEEESGSSSASEEEDTKPKPTK20222020000000000000	sp Q68E01 INT3_HUMAN
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12259	NWTEDEMEGGISSPVK20000	sp P08651 NFIC_HUMAN
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12261	SFQDYTGQK2000000000	sp Q9NP64 NO40_HUMAN
12262	TTSRSPVLSR2002000000	sp 095819 M4K4_HUMAN
12263	IKEEPEVEVDSSPPDSPASSPCSPPLK2200020022000000000	sp P17544 ATF7_HUMAN
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12265	LPSSPASPSPK2000	sp Q9H4G0 E41L1_HUMAN
12266	SSPINPAQSGSPNHVDSAYFPGSSTSSSSDNDEGSGGATNHISGNK2E+22	sp Q9NZB2 F120A_HUMAN
12267	TPSPESHRSAPGSEER202000002000000000	sp Q66K74 MAP1S_HUMAN
12268	FHSSDSEEEHK22020000000	sp Q96ST2 IWS1_HUMAN
12269	ERFSPRHELSPQK2000000200000	sp Q9BXP5 SRRT_HUMAN
12270	QRSVSPYSR20200000	sp Q9NYV4 CDK12_HUMAN
12271	SRTASGSSVTSLDGTR2022022000000	sp Q92597 NDRG1_HUMAN
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12273	ASPAPGSGHPEGPGAHLDMNSLDR20000000000000000000	sp 094826 TOM70_HUMAN
12274	SQSPHYFR2000000000	sp Q9UKJ3 GPTC8_HUMAN
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12276	TSLMSAESPTPR200200000	sp P49810 PSN2_HUMAN
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12278	GADNDGSGSESGYTTTPK20300000	sp Q7Z417 NUFP2_HUMAN
12279	QICVVMLETLSQSPPK2000000	sp Q15366 PCBP2_HUMAN
12280	KSDGACDPSDDKENSQIAQDHQK200002000000000000	sp Q9UIG0 BAZ1B_HUMAN
12281	RPTETNPVTSNSDEECNETVK20000200000000000000	sp P46100 ATRX_HUMAN
12282	KPPAPPSPVQSQSPSTNWSPAVPVKK2.00000000002E+20	sp Q8IYB3 SRRM1_HUMAN
12283	ESSPEKEAEEGCEKESEEGCPK2E+22	sp 043719 HTSF1_HUMAN
12284	SSDQDSTNKEAEAAGVKPAGVRPR2.00002E+23	sp Q92625 ANS1A_HUMAN
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12286	RPACDPEPEGSGSSSDEGCTVVRPEK222000020000000	sp 015541 R113A_HUMAN
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12291	LSEAVWQPEEHYSSSPEK20000	sp Q9Y485 DMXL1_HUMAN
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12293	QSSSRFNLSK220000000	sp Q14738 2A5D_HUMAN
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12300	SDGEAKPEPSPSPR200000000002000	sp P26358 DNMT1_HUMAN
12301	RAPSTSPSFEGTQETYTVAHEENV2.2E+21	sp Q9BUT9 F195A_HUMAN
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12309	SPLEHSSPEKEAPSPEK2200000020000	sp Q01082 SPTB2_HUMAN
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12312	SGSDAGEARPPTPASPR200000000000002000	sp Q9UPZ3 HPS5_HUMAN
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12315	NTSVGSTSTGVESILEPCDEIEQK2000200000000000	sp Q96AY4 TTC28_HUMAN
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12317	LDSSCLESVK20000000	sp Q96BD5 PF21A_HUMAN
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12323	DTHEDHDTSTENTDESNHDPQFEPVIVSLPEQEIK2.000002E+33	sp P37275 ZEB1_HUMAN
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12325	LPSSPASPSPK2000200000	sp P39880 CUX1_HUMAN
12326	HDSGGLPLTPR20020002000	sp Q9H4G0 E4IL1_HUMAN
12327	SFVPEEEK200000000	sp Q96MH2 HEX12_HUMAN
12328	APSASDSKADSDGAKPEPVAMAR2.00020002E+21	sp 094880 PHF14_HUMAN
12329	TNPPTQKPPSPMSG20010000	sp Q7Z4V5 HDGR2_HUMAN
12330	SAQGSSSPVPSMVQK20000010000	sp Q8IZP0 ABI1_HUMAN
12331	MSHSSSDTDINEIHTNHK2022220200000000000	sp P35658 NU214_HUMAN
12332	YPRPASVPPSPSLSRHSSPHQSEDEEDPR2.000202020022E+24	sp P46100 ATRX_HUMAN
12333	HTSCSSAGNDSKPVQEAPSVAR200002000000000000	sp P13807 GYS1_HUMAN
12334	TDSQSVRHSPVLAQK200002200000000000	sp Q9P246 STIM2_HUMAN
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12340	SRPQGLTEAEQR20000000000000	sp Q15836 VAMP3_HUMAN
12341	SPTTPIDPEK20020000000	sp P23763 VAMP1_HUMAN
12342	SPTSPSPPR20200020000	sp TAU_HUMAN
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12348	GTFSDQELDAQSLDDEDDNMHHAVYPAVNR2.00000002E+27	sp P35579 MYH9_HUMAN
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12350	SNSPLPSIQLPQSPSASKK2000000000020000000	sp Q569K4 Z385B_HUMAN
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		sp Q08174 PCDH1_HUMAN
		sp Q8IWS0 PHF6_HUMAN
		sp Q9UQ35 SRRM2_HUMAN
		sp Q5VUA4 ZN318_HUMAN

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12357	TGSNISGASSDISLDEQYK220000000000	sp P22059 OSBP1_HUMAN
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12372	DGSGTPSRHSLSGSSPGMK20020202000000	sp Q9UQ35 SRRM2_HUMAN
12373	VSDSESEGPQKGPASDSETEDASR2.0202000000002E+23	sp Q96ST2 IWS1_HUMAN
12374	RGGADTYSVPSVLSR202002000000	sp P14543 NID1_HUMAN
12375	SRSRSPPVSK200020000000	sp Q01130 SRSF2_HUMAN
12376	GVSMSLPSSPLLPR2100002000000	sp Q9C0D5 TANC1_HUMAN
12377	NSVLSDPGLDSPRTSPVIMAR200000200000000000	sp 015085 ARHGB_HUMAN
12378	GGPASVPSSSPGTSVK200000000	sp 043237 DC1L2_HUMAN
12379	TPLSSLSATSNNSTVACIDR220000000000000000	sp P00533 EGFR_HUMAN
12380	SPALHGTPTAGCGSR2000000020000000	sp Q8IY92 SLX4_HUMAN
12381	SSGKTPETLVPTAPK2002000000000000	sp Q14676 MDC1_HUMAN
12382	KHSPSPPTPTESR20000000020000	sp Q92922 SMRC1_HUMAN
12383	DLGTQNHTSELILSSPPGQK2000000	sp Q9ULD2 MTUS1_HUMAN
12384	SELESR2000000	sp Q92974 ARHG2_HUMAN
12385	LDVTILSPSR20000	sp 075369 FLNB_HUMAN
12386	RMSGEP1QTVESIR20000000000000	sp Q5VT52 RPRD2_HUMAN
12387	SRSKSPFR202020000	sp Q14498 RBM39_HUMAN
12388	HCGLSLSTPPGK20200000	sp Q71F23 CENPU_HUMAN
12389	SPNREWEVLQAPHQVITNLPEGVR2E+25	sp 075376 NCOR1_HUMAN
12390	TLSPTPASATAPTSQGIPTSDEESTPKK2E+28	sp Q9NQS7 INCE_HUMAN
12391	GGPASVPSSSPGTSVK20000000	sp 043237 DC1L2_HUMAN
12392	TKPTQAAGPSSPQKPPTPEETK2000200	sp P18887 XRCC1_HUMAN
12393	VASSPVMVSNPATR200010000000	sp P51610 HCFC1_HUMAN
12394	RHSSDINHLVTQGRESPEGSYTDANQEV2.0000002E+27	sp Q5TON5 FBP1L_HUMAN
12395	SRSPSSPELNK2020200000000	sp Q9UQ35 SRRM2_HUMAN
12396	LDAEPKPCSDTNSDSPDEGLDHNHIK2.000202E+19	sp Q13129 RLF_HUMAN
12397	ESEKSLENETLNKEEDCHSPTSKPPKPDQPLK2.00002E+28	sp P23588 IF4B_HUMAN
12398	SSESPNHSLHNEVADDSQLEK2.2E+22	sp Q9ULH0 KDIS_HUMAN
		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
12399	EDALDDSVSSSVHASPLASSPVRK2.00000000000002E+19	sp A6NKT7 RGPD3_HUMAN
		sp Q99666 RGPD5_HUMAN
		sp 014715 RGPD8_HUMAN
12400	NLSPTAVINITSEK2000000000000	sp Q92870 APBB2_HUMAN
12401	GLQVDLQSDGAAAEDIVASEQSLGQK2E+19	sp P33991 MCM4_HUMAN
12402	SRSRSPLAIR20202000000	sp Q9UQ35 SRRM2_HUMAN
12403	QLGDSSGEGPEFVEEEEEVALRSDSEGSYTPGKK20200200000000	sp Q14839 CHD4_HUMAN

12404	GSSDGRGSDSESDLPHR20002000000	sp Q9UPQ0 LIMC1_HUMAN
12405	FNDPSGCAPSPGAYDVK200000000	sp 075330 HMMR_HUMAN
12406	AGSSPCLGSSSAASSPPPPASR200000000	sp Q6ZRS2 SRCAP_HUMAN
12407	AISSDDSSDEDK2200220000000	sp Q6PD62 CTR9_HUMAN
12408	SSWLSTEGGR2002000000	sp Q9BXF6 RFIP5_HUMAN
12409	LEDAESSNPNLQSLSTDALPSASK2.000002E+19	sp Q9Y294 ASF1A_HUMAN
12410	QSPGHQSPLASPK20002000	sp Q8N1G0 ZN687_HUMAN
12411	SAVSPDLHITPIYEGR2000000000000	sp Q99569 PKP4_HUMAN
12412	HSSISPSTLTLK20000002000	sp Q14004 CDK13_HUMAN
12413	TRSPSPDILER2020000000000	sp Q13523 PRP4B_HUMAN
12414	SPQTLAPVGEDAMKTPSPAEDAREPEAK202000000000000	sp Q92797 SYMPK_HUMAN
12415	APAGPSLEETSVSPPK2000	sp Q6AI08 HEAT6_HUMAN
12416	RVSSDEEHTVDSICSDMK2000020000000000	sp Q70CQ2 UBP34_HUMAN
12417	SPSPPTQHTGQPPGQPSAPLSAPR20000000000	sp Q68EM7 RHG17_HUMAN
12418	VSVTPPEESQNSDTPPRPDR200002000000000000	sp Q05209 PTN12_HUMAN
12419	LTPVSLNSPIKGADCQEVQDK2.0000002E+22	sp P46100 ATRX_HUMAN
12420	TNLTSQSSTNLPGSPGSPGSPGSPGSPGSPK2.002002E+19	sp Q6IN85 P4R3A_HUMAN
12421	AVIVSSPK2000	sp Q14966 ZN638_HUMAN
12422	LKSVEDEMDSPGEEPFTYTGQR2.00001E+20	sp Q12857 NF1A_HUMAN
12423	TPSPKEEDEEPESPPEKK20000000002000000	sp Q9H1E3 NUCKS_HUMAN
12424	SGEQITSSPVSPK20000000002000	sp 043314 VIP2_HUMAN
12425	SRSRTSPITR20200200000	sp Q9UQ35 SRRM2_HUMAN
12426	LSSSATLLNSPDR200000000	sp Q14244 MAP7_HUMAN
12427	AKSQYSGQLHEVR200000000000	sp Q9BXB5 OSB10_HUMAN
12428	SSPAELSSSQHLLR2000020000000000	sp Q9UQC2 GAB2_HUMAN
12429	AGDTSQEGHSLEEK200000000000	sp Q99607 ELF4_HUMAN
12430	LSPSPSSR20000000	sp Q03252 LMNB2_HUMAN
12431	RHSSQIEGVR220000000	sp P20700 LMNB1_HUMAN
12432	GPSSPPTTAPMR20000000100	sp Q6RI45 BRWD3_HUMAN
12433	ASGGAQSPLR20000	sp Q9NXL9 MCM9_HUMAN
12434	LSLQHTQQNADGQEDGESERNSNSSGDPGR22000000	sp Q6NXT1 ANR54_HUMAN
12435	TQSLSALPK20000000	sp 075052 CAPON_HUMAN
12436	YSPTSPTYSPK32000000	sp Q96RK0 CIC_HUMAN
12437	ATKMLTATAMPTVATSVQSPYRSTK220020302000	sp P24928 RPB1_HUMAN
12438	NQSHSSPSVSPSR202202020000	sp Q9UPU5 UBP24_HUMAN
12439	ISDSESEDPPRHQASDSENEELPKPR2.020000000002E+23	sp Q7Z6E9 RBBP6_HUMAN
12440	SSSVGSSSYPIPAVSR220000000000000	sp Q96ST2 IWS1_HUMAN
12441	SPVSPGSSSPQLQVK20000000020000000	sp Q15149 PLEC_HUMAN
12442	GTPPLTPSDSPQTR2000000000	sp Q8N3F8 MILK1_HUMAN
12443	GVGSGPHPPDTQQPSPSK200	sp Q6NZY4 ZCHC8_HUMAN
12444	SRSRSGPLR20200200000	sp Q68CZ2 TENS3_HUMAN
12445	SPSSESPQHPTPPARPR200000000000000000	sp Q9UQ35 SRRM2_HUMAN
12446	SKSQSSGSSATHPIVPGAR2200000000000000	sp 014733 MP2K7_HUMAN
12447	SCDEPLTPPHSPTSMLQLIHDPVSPR220000000002000	sp Q9Y4H2 IRS2_HUMAN
12448	SPQTPSQMVPLPSANPPGPK2.0000201E+21	sp Q9Y2K7 KDM2A_HUMAN
12449	ADGSPVNTGVETTTPSK20000020000	sp Q86UU0 BCL9L_HUMAN
12450	NKQDDLNCEPLSPHNITPEPVSK2000020000000	sp P0CF74 LAC6_HUMAN
12451	DRGDAYYDTPRHSYEQR2000000	sp Q6VMQ6 MCAF1_HUMAN
12452	KGSRSDTSAR20000	sp P49760 CLK2_HUMAN
12453	SRSSSVGSSSYPIPAVSR220000000000000000	sp Q5SYE7 NHSL1_HUMAN
		sp Q15149 PLEC_HUMAN

12454	SRSYSR20230000	sp Q8IYB3 SRRM1_HUMAN
		sp Q13595 TRA2A_HUMAN
		sp P62995 TRA2B_HUMAN
		sp Q07955 SRSF1_HUMAN
12455	SATSTSASPTLR220000000	sp 075061 AUXI_HUMAN
12456	HCGLSLSSTPPGK2020200000	sp Q71F23 CENPU_HUMAN
12457	FSPSQVPIQTR20000000000	sp Q08AD1 CAMP2_HUMAN
12458	QDKPMDTSVLSEEGEPFQK20020000000000	sp Q12888 TP53B_HUMAN
12459	SPAGPAATPAQAQAASPR2.0000002E+19	sp Q13428 TCOF_HUMAN
12460	WSDQGAQTSR20000000000	sp Q9COC2 TB182_HUMAN
12461	SLDGAEFRPASVSENHDAGPDGDK2.000000000002E+25	sp Q9H4G0 E41L1_HUMAN
12462	LQDSSDPDTGSEEGSSRLSPPHSPR20000200000000000	sp Q92974 ARHG2_HUMAN
12463	SGSTSPLGFAR20000000	sp 075385 ULK1_HUMAN
12464	TPMYGSQTPLQDGSR20000000000000000	sp 000267 SPT5H_HUMAN
12465	RSSDPWSNSHFPHEN20000000000000	sp Q92545 TM131_HUMAN
12466	TAPTSTIAPGVMASSPALPTQPAEEAAR2000000000000000	sp P16220 CREB1_HUMAN
12467	DEPPPLSPALTPATPSSLDPFPSR2.00000002E+19	sp Q7Z6Z7 HUWE1_HUMAN
12468	GYSPYSVSGSGSTAGSR300000000200200	sp Q15149 PLEC_HUMAN
12469	GPSAAGEQEPDKESGASVDEVAR20020000000	sp P50579 MAP2_HUMAN
12470	DDSHAEDSEDEKEDHK20000020000000000	sp Q9H1E3 NUCKS_HUMAN
12471	KSPSGPVKSPPLSPVGTTPVK2000220000	sp Q9BVC5 ASHWN_HUMAN
12472	EELMSSDLEETAGSTSIPKR22000000000000000	sp 000567 NOP56_HUMAN
12473	GSISLNSSPR2000	sp Q9NYL2 MLTK_HUMAN
12474	HLSTPSSVSPEPQDSAK20000000000000000	sp Q9NZN5 ARHGC_HUMAN
12475	NSVQTPVENSTNSQHVK20000000000000000	sp Q9UHI6 DDX20_HUMAN
12476	SPDSCRPAALPCLPSTQDVPSR2E+19	sp Q9Y283 INVS_HUMAN
12477	MQSLQLDCVAVPSSR1020000000000200	sp Q13546 RIPK1_HUMAN
12478	LHDFLAHSSESEETSSPPR20000000000000	sp P46100 ATRX_HUMAN
12479	WGQPPSPTVPRPPDADPNTSPKPLEGRPER2.00000000000202E+25	sp Q14839 CHD4_HUMAN
12480	TISPSTSNLIPLMR2000200000100	sp 094806 KPCD3_HUMAN
		sp Q15139 KPCD1_HUMAN
12481	SYLEGSSDNQLK20000200000000	sp Q9UHI6 DDX20_HUMAN
12482	RASEELDGLFR20000000000	sp Q14814 MEF2D_HUMAN
12483	LVTSPQSEPPAPFPPPR20000000000000000	sp Q9H792 PEAK1_HUMAN
		sp Q9Y4E1 FA21C_HUMAN
12484	ETVSEAPLLFSDEEEK20000000	sp Q641Q2 FA21A_HUMAN
		sp Q5SNT6 FA21B_HUMAN
12485	VAEPGAEATSSSTGEESGSEHPPAVPMHNK2.0000002E+21	sp Q14684 RRP1B_HUMAN
12486	GNVFSSTAAGTPNK20000020000	sp Q05682 CALD1_HUMAN
12487	SISNPPGSNLR20000000000	sp Q9H4A3 WNK1_HUMAN
12488	ATRPASTSQPDSPDR2000020000	sp P20749 BCL3_HUMAN
12489	SHMSGSPGPGSNTAPSTPVIGGSDKPGMEEK2E+21	sp A0FGR8 ESYT2_HUMAN
12490	YTPTSPSYSPSSPEYTPTSPK2000000000000	sp P24928 RPB1_HUMAN
12491	GSSPSIRPIQSGQSSSPVEK20020000000	sp Q92560 BAP1_HUMAN
12492	QEPAAEYETPESPVPPARSPSPPK200000020200000	sp P54259 ATN1_HUMAN
12493	GPTTGEGALDLSDVHSPPKSPEGK2E+21	sp 095684 FR10P_HUMAN
12494	KRESESEDETTPAAPQLIK20200200000000000	sp 060832 DKC1_HUMAN
12495	IDAGTMAEPSASPSK20000000	sp Q8IXQ3 CI040_HUMAN
12496	KHHESI TPPSRSPK20020000000	sp Q8IY92 SLX4_HUMAN
12497	TKPTQAAGPSSPKPPTPEETK2E+22	sp P18887 XRCC1_HUMAN
12498	TAMPSPGVSQNK102000000000	sp Q86UU0 BCL9L_HUMAN
12499	GEVPGGSAHYGGPSPEK20000	sp 094762 RECQ5_HUMAN
12500	AASPYSQRPASPTAIR200000002000000	sp Q99569 PKP4_HUMAN

12501	NIIGSSPVADFSAIK20000000000	sp P42566 EPS15_HUMAN
12502	FSRSPIR2020000	sp P18583 SON_HUMAN
12503	DGYGSRDSYSSSRSDLYSSGR202200000000000	sp P38159 RBMX_HUMAN
12504	DTHEDHDTSTENTDESNHDPQFEP IVSLPEQEIK2. 002E+25	sp P43487 RANG_HUMAN
12505	QTESQIMGELPFTIASK200000000000000	sp 000370 LORF2_HUMAN
12506	TASPLPLRPDAPVEK20000000000000000	sp Q9HCK8 CHD8_HUMAN
12507	SRPLATGPSSQSHQEK200000	sp Q96RL1 UIMC1_HUMAN
12508	NQKPSQVNGAPGSPTEPAGQK20000000	sp Q9BQG0 MBB1A_HUMAN
12509	KRSSLSPGR2020000	sp Q9BUQ8 DDX23_HUMAN
12510	SRSRSTPPSSAGLGSNSAPPIPDSR2. 0202002E+24	sp Q8IWX8 CHERP_HUMAN
12511	TESPATAAETASEELDNR2000000000	sp Q9NTJ3 SMC4_HUMAN
12512	AASSSPGSPVASSPSR22000000000000000	sp Q92538 GBF1_HUMAN
12513	QFSTASDHEQPGVSG200200000000000	sp Q66K14 TBC9B_HUMAN
12514	GSSDGRGSDSEDLPHR202000000	sp Q9UPQ0 LIMC1_HUMAN
12515	TISINSPK2000	sp Q9P1Y6 PHRF1_HUMAN
12516	LSSLRASTSKSESSQK200022000000000	sp P62753 RS6_HUMAN
12517	IGNEESDLEEACILPHSP INVDK20000000	sp 095714 HERC2_HUMAN
12518	ENSSPSVTSANLDHTKPCWYWDK2E+20	sp Q9BVR0 HRC23_HUMAN
12519	HKSGSMEEDVDTSPGGDYITSPSSPTSSSR1. 00000000000003E+25	sp 075909 CCNK_HUMAN
12520	SSSFSDTLEESSPIAAIFD TENLEK220000000000000	sp P08651 NFIC_HUMAN
12521	SPSESPQHIAHPER20000000000	sp P15924 DESP_HUMAN
12522	APVPEPGLDLSLSPRPDSPQR20000200000	sp Q68DK2 ZFY26_HUMAN
12523	QRSHSGTSPDNTAPPPPPRPQPSHSR2. 0202E+25	sp Q8TAP8 PPR35_HUMAN
12524	ISDSESEDPPRNQASDSENEELPKPR2. 0200000000202E+23	sp Q96D71 REPS1_HUMAN
12525	RSPPADAIPK2000000000	sp Q96ST2 IWS1_HUMAN
12526	LQDSSDPDTGSEEGSSRLSPHSPR2E+23	sp P18754 RCC1_HUMAN
12527	KRSDAGSSTPTTSTR20002000000000	sp Q92974 ARHG2_HUMAN
12528	LYPPGDEEEAELADSSGLYHEGSPSPGSPWK20000	sp P51532 SMCA4_HUMAN
12529	TSPINEDIR200000000	sp Q8NCF5 NF2IP_HUMAN
12530	EYIPGQPPLSQSSDSSPTR20200000	sp Q9P2R6 RERE_HUMAN
12531	SFISPIPSK2000000000	sp P07814 SYEP_HUMAN
12532	SIKPSDSPR2000	sp Q9UGU0 TCF20_HUMAN
12533	GNVFSSPTAAGTPNKETAGLK200000020000000000	sp Q92610 ZN592_HUMAN
12534	SRSRTPPVTR20202000000	sp Q05682 CALD1_HUMAN
12535	SEHSHSTTLPR200000220000	sp Q9UQ35 SRRM2_HUMAN
12536	QEKPAEKPAETPVATSPTATDSTSGDSSR2000000000000	sp P16144 ITB4_HUMAN
12537	GLSLSPQSPPER20000200000	sp P54727 RD23B_HUMAN
12538	LQSGVSLASK200000200	sp Q7Z2Z1 TICRR_HUMAN
12539	SRSPSPAPEKK202000000000	sp Q09666 AHNK_HUMAN
12540	VDKETNTETPAPSPTVVRPK20000000200000000	sp Q8IYB3 SRRM1_HUMAN
12541	APSDTGSLSPPWDQDRR202000000000	sp Q8IX03 KIBRA_HUMAN
12542	STSPAGQHHSP ISSR220000000002200	sp 015320 CTGE5_HUMAN
12543	NESPPQVPVYNNSHK200000000000000	sp Q5T200 ZC3HD_HUMAN
12544	EHSPYGPSPLGWPSSETR20000200000000000	sp Q9UKL3 C8AP2_HUMAN
12545	TKPTQAAGPSSPQKPPTPEETK2E+19	sp 015320 CTGE5_HUMAN
12546	GYSSPEPDIQDSSGSEAQSVKPSTR2. 000000022E+22	sp P18887 XRCC1_HUMAN
12547	GKPPAPSPR2000	sp Q9BZF1 OSBL8_HUMAN
12548	DVYLSR2000	sp Q9NP71 MLXPL_HUMAN
12549	YTPTSPSYSPSSPEYTPTSPK3. 0000003E+21	sp P38159 RBMX_HUMAN
12550	NIVVDDITEQKPEPQDDGKSTESDVK20020000	sp Q96E39 RMXL1_HUMAN
12551	AGGGAQPSQTATGDPSDEDEDSLEEFVR2000200000000000000	sp P24928 RPB1_HUMAN
12552	SPRPQQDPARPQEPTMPPPETPSEGR2E+26	sp Q9Y4W2 LAS1L_HUMAN
		sp Q2YD98 UVSSA_HUMAN
		sp P29590 PML_HUMAN

12553	SLSPNHNTLQTLK200000000000	sp 075592 MYCB2_HUMAN
12554	LTNLSSNSDV2000000	sp Q13641 TPBG_HUMAN
12555	DSGRGDSVSDSGSDALR202000000000	sp Q53EL6 PDCD4_HUMAN
12556	SPSDAGRSSGDESKKPLPSSSR2.0200002E+22	sp Q8IVL1 NAV2_HUMAN
12557	LASTSDIEEK2000000	sp O14974 MYPT1_HUMAN
12558	MDSLHDMLTDPLELGPCGDGHGTR2.0001E+22	sp Q6P4R8 NFRKB_HUMAN
12559	RSSQPSPTAVPASDSPPTK20200000	sp Q3KQU3 MA7D1_HUMAN
12560	FDHESPGTDEDKSG220020000200	sp Q13427 PPIG_HUMAN
12561	KRSELSQDAEPAGSQETK200000000000000000	sp Q9BVJ6 UT14A_HUMAN
12562	STLESEKPGSPEAAETSPPSNIIDHCEK2E+27	sp Q96T23 RSF1_HUMAN
12563	RREEGPPPPSPDGASSDAEPEPPSGR20000000000000000000	sp Q9NTJ3 SMC4_HUMAN
12564	RSSLSHSHQSQIYR2000200000000000	sp O15027 SC16A_HUMAN
12565	NGFPHPPEPCNPSEAASEESNSEIEQEIPVEQK2.000200002E+21	sp Q9NR30 DDX21_HUMAN
12566	RFTPPSTALSPGK200200000000	sp Q01196 RUNX1_HUMAN
12567	VCSSDSDSSLQVVK22000200000000	sp Q9NSI6 BRWD1_HUMAN
12568	LGPQNFIDETSDIENLLKEEEDYKEENNSK2E+21	sp Q14692 BMS1_HUMAN
12569	DSPAFPDSPWR20000	sp Q9NQS7 INCE_HUMAN
12570	TVSEKHQGK200000000	sp Q9NSC7 SIA7A_HUMAN
12571	VPEASSEPFDTSSPQAGR2000000	sp Q9H4I2 ZHX3_HUMAN
12572	ISLEQPPNGSDTPNPEK202000000	sp Q9UHI6 DDX20_HUMAN
12573	SKSPSPPR202020000	sp Q9UKV3 ACINU_HUMAN
12574	YGNLTETRTLLAIQQQALR3.00020202E+19	sp Q9Y2J4 AMOL2_HUMAN
12575	DGSGTPSRHSLSGSSPGMK200202000000000	sp Q9UQ35 SRRM2_HUMAN
12576	SCTPSPDQISHRASLEDAPVDDLTR2.02000020002E+23	sp Q7Z2W4 ZCCHV_HUMAN
12577	SESVVSGLDSPAK200000000	sp O14578 CTRO_HUMAN
12578	QAAVTSTSLTIKSVFTR22220000000200	sp P20273 CD22_HUMAN
12579	LSPERSDSESNENEPHLPHGVSQR2.2E+20	sp Q9BQ52 RNZ2_HUMAN
12580	SDISDQEEDEESEGCVPVSNLSK20000000000000	sp Q15154 PCM1_HUMAN
12581	IEDVTPIPSDSTR2000	sp P62263 RS14_HUMAN
12582	TSPPCSPANLSR200020000200	sp P49585 PCY1A_HUMAN
12583	HAPSLHGSTELLPLSR200000200	sp Q9H5N1 RABE2_HUMAN
12584	LLELTSSYSPDVSDYK200000000	sp P38432 COIL_HUMAN
12585	DIFYYEDDSEGEDIEK200000000	sp Q15361 TTF1_HUMAN
12586	KVATPPNQK200000000	sp Q6NUQ4 TM214_HUMAN
12587	STTTPPRCSPAR2000020000	sp Q6ZRS2 SRCAP_HUMAN
12588	LPQSSSESSPPSPQPTKVSRR2000200000	sp Q9UQ35 SRRM2_HUMAN
12589	SGSTSPLGFAR220000000	sp O75385 ULK1_HUMAN
12590	RPPTPEAQSEEERSDEEPEAK20000200000000000000	sp Q9BTK6 PAGR1_HUMAN
12591	YLVDTGKPNAGSEEISSEDELVEEK2.0000000022E+22	sp Q9NY61 AATF_HUMAN
12592	AEGEWEDQEALDYFSDKESGK2000	sp Q9NYF8 BCLF1_HUMAN
12593	IDGATQSSPAEPK2000000	sp Q32MZ4 LRRF1_HUMAN
12594	GPPASSPAPAPK200000000	sp Q15942 ZYX_HUMAN
12595	EHPHSRSPSPEPR2020200000	sp Q9UDY2 ZO2_HUMAN
12596	SQSPINCIRPSVLGPLK20200000000000000000	sp Q96E09 F122A_HUMAN
12597	TFSDTTHGSVPSDPLGR20000000000000	sp Q86V48 LUZP1_HUMAN
12598	YIAKFQGEDTVVIASKPYAFDR3.000000002E+22	sp P33176 KINH_HUMAN
12599	QPPLSSQRPR2200000	sp Q8NFI3 ENASE_HUMAN
12600	DRTPPHLLYSR200000000000	sp Q8NDT2 RB15B_HUMAN
12601	DGDSYDPYDFSDTEEMPQVHTPK2.0000000000001E+21	sp P25205 MCM3_HUMAN
12602	RSRSASLR20202000	sp Q16629 SRSF7_HUMAN
12603	ESESEDETTPAAPQLIK200200000000000	sp O60832 DKC1_HUMAN
12604	TAHNSEADLEESFNEHELEPSSPK20000000002000	sp Q8IWS0 PHF6_HUMAN
12605	YTPTSPSYSPSPEYTPSPK300000000002000	sp P24928 RPB1_HUMAN
12606	APQSGSPSPGSLAPR22000000000	sp MQZC1 RN225_HUMAN



12607	SETAPAAPAAPAPAEKTPVKK200000	sp	P10412	H14_HUMAN
12608	SRTHSTSSSLGSGESPF202000000000000000	sp	Q9UGV2	NDRG3_HUMAN
12609	SPQNDHSDTDSEENRDNQFLTTVK2. 000000202E+26	sp	Q86W56	PARG_HUMAN
12610	GGSGYVQLSAGYESVDSPTGSENSTHQSNDTSSHDPQEEK3. 00000000000202E+31	sp	Q15007	FL2D_HUMAN
12611	TPTSSPASSPLVAK200000022000000	sp	Q14684	RRP1B_HUMAN
12612	RQSSGSATNVASTPDNR200000	sp	Q7Z460	CLAP1_HUMAN
12613	VSSPLSPLSPGIKSPTIPR200002000000	sp	Q9P2B4	CT2NL_HUMAN
12614	KFELLTPPLSPSR200000200	sp	P04198	MYCN_HUMAN
12615	SGSGSVGNSSRYSPSQNSPIHHIPSR2. 0202000022032E+27	sp	P01106	MYC_HUMAN
12616	VNFSEEGETEEDDQDSSHSSVTTVK2. 0000200000022E+22	sp	Q9NYF8	BCLF1_HUMAN
12617	KSPSPAGGGSPYSR202000000003000	sp	Q5JTV8	TOIP1_HUMAN
12618	RSSPSARPPDVPGQPQAAK2000000000000000000	sp	Q14004	CDK13_HUMAN
12619	TPASGLQPTSTPAPGSATR2E+20	sp	Q96JP5	ZFP91_HUMAN
12620	QQSSSSKGDSPKPR202200000000000	sp	Q96DF8	DGC14_HUMAN
12621	YLSFTPPEK200000	sp	Q5VWQ8	DAB2P_HUMAN
12622	SDGAPASDSKPGSSEAAPSSKETPAATEAPSSTPK200000000000000000	sp	Q13177	PAK2_HUMAN
12623	KQPPVSPGTALVGSQKEPSEVTPK2000000000000	sp	P80723	BASP1_HUMAN
12624	FFTTGSDSESESSLGEELVTKPVGGNYGK2. 020202E+25	sp	P17096	HMGA1_HUMAN
12625	SPGSGSQSSGWHEVEPGMPSPTTLK2E+22	sp	B5ME19	EIFCL_HUMAN
12626	SALSPSKSPAK200000020000	sp	Q99613	EIF3C_HUMAN
12627	SESAPTLHPYSPLSPK200000000002000	sp	Q12857	NFIA_HUMAN
12628	SSTSFANIQENSN200000000000	sp	P49792	RBP2_HUMAN
12629	TLGTPTQPGSTPR20000022000	sp	Q7Z3J3	RGPD4_HUMAN
12630	DLTHSDSESSLHMSDR200000000	sp	Q8WUF5	IASPP_HUMAN
12631	SPASPSQAPYAQR20020000000000	sp	Q86WC4	OSTM1_HUMAN
12632	RLSSSESPQRDPPPPPPPPPLLR2. 02E+19	sp	Q8NFH5	NUP53_HUMAN
12633	STPLASPSPSGRSPQR2000200000000	sp	Q13586	STIM1_HUMAN
12634	EASTTAAAPGEETSPVQVPAR200000000	sp	Q8NE35	CPEB3_HUMAN
12635	LASVPAGAVAVSAAPGSAAPAAGSAPAAAEK2000000000	sp	Q9H2G4	TSYL2_HUMAN
12636	TPPPHSHAAPSTGSPVPATEPPAAK200000020000000	sp	Q8WYP5	ELYS_HUMAN
12637	RPESAPAESSPSK20000020000	sp	Q5SXM2	SNPC4_HUMAN
12638	SSTPLPTISSAENTR22000000000000200	sp	P05387	RLA2_HUMAN
12639	IEIPVTPTGQSVSPSPSIPGTPTLK2000200000	sp	Q9UKP4	ATS7_HUMAN
12640	VKSSTPPRQSPSR22000020000	sp	Q07889	SOS1_HUMAN
12641	SSISLGWKPAPKDGGSPIK2000000000000020000	sp	P42166	LAP2A_HUMAN
12642	TASISSPSEGTPVGSYGCTPQSLPK2E+27	sp	P42167	LAP2B_HUMAN
12643	VLTANSNPSSPSAAK2000000	sp	Q9H4Z3	PCIF1_HUMAN
12644	SRDATPPVSPINMEDQER200020000000000	sp	Q9UQ35	SRRM2_HUMAN
12645	SPVVGDTSPR20002000	sp	Q8WZ42	TITIN_HUMAN
12646	LNSSVLVPYGAESSEDSDEESK22002000000	sp	Q6PKG0	LARP1_HUMAN
12647	VLSGNCNHQEGTSSDDELPSAEMIDFQK2. 00000000002E+26	sp	Q9NV56	MRGBP_HUMAN
12648	IGFPETEEEELEEIASENSDCIFPSAPDVK2000000000000000	sp	P17275	JUNB_HUMAN
12649	ASINLLSDK2002000	sp	P10636	TAU_HUMAN
12650	KEMDPVTQLYTMTSTLEYK100000030100200000	sp	TAU_HUMAN	
12651	QEKPAEKPAETPVATSPTATDSTSGDSSR2E+19	sp	Q9H9J4	UBP42_HUMAN
12652	DLSTSPKPSPIPSPVLGR20002000000	sp	P16383	GCFC2_HUMAN
12653	GMGSDFEDESREGDPEER1020000000000000000	sp	Q9Y3F4	STRAP_HUMAN
		sp	Q04656	ATP7A_HUMAN
		sp	Q13740	CD166_HUMAN
		sp	P54727	RD23B_HUMAN
		sp	Q8NDI1	EHBP1_HUMAN
		sp	Q5TEC3	ZN697_HUMAN

12654	CSSLPPMTPKTMPR20000000	sp	Q9NYZ3	GTSE1_HUMAN
12655	GSSDGRGSDSEDLPHR200000200000000000	sp	Q9UPQ0	LIMC1_HUMAN
12656	TDSREDEISPPPPNPVVK2000000020000000000	sp	P10644	KAPO_HUMAN
12657	SRSGSSQELDVKPSASPQER2022000000000000000	sp	Q9UQ35	SRRM2_HUMAN
12658	SGYSSPGSPGTPGSR200200000000	sp	P10636	TAU_HUMAN
		sp	TAU_HUMAN	
12659	RLSLPADIR20000000	sp	Q00536	CDK16_HUMAN
		sp	Q00537	CDK17_HUMAN
12660	AADGGERPLAASPPGTVK20000000	sp	095785	WIZ_HUMAN
12661	CVPAPGAGASGGTSPSATQPNPAVFIFEHK2000000000000000000	sp	060343	TBCD4_HUMAN
12662	RVSSRSVSGSPEPAAK2202020200000000	sp	Q8IYB3	SRRM1_HUMAN
12663	DRDYSDHPSGGSYRDSYESYGNSR300000000	sp	Q96E39	RMXL1_HUMAN
		sp	P38159	RBMX_HUMAN
12664	ATSPSSVSGDFDDGHHSVSTPGPSR2. 2E+25	sp	Q86U86	PB1_HUMAN
12665	ETSVSKEDTDHEEKASNEVDTK2. 2E+21	sp	075475	PSIP1_HUMAN
12666	RGPDRGSPEYR200000	sp	P49716	CEBPD_HUMAN
12667	DGSGTPSRHSLSGSSPGMK20200202000000000	sp	Q9UQ35	SRRM2_HUMAN
12668	LPSKELPDSSSPVANNIR2000000020000000000	sp	Q9BWT3	PAPOG_HUMAN
12669	TPPITTK20000000	sp	Q5T200	ZC3HD_HUMAN
12670	SQSSSQDNPLAPGYLPPHYK2. 022E+21	sp	Q6ZRV2	FA83H_HUMAN
12671	TQDPSSPGTTPQAR22000000	sp	Q13112	CAF1B_HUMAN
12672	SPPVQPHTPVTISLGTAPSLQNNQPVEFNHAINVVK2. 000000000002E+37	sp	Q96ST3	SIN3A_HUMAN
12673	SSGHSSSELSPDAVEK220022200000000000	sp	Q9UQ35	SRRM2_HUMAN
12674	VHSTSSLDSQK2000000000	sp	Q96JI7	SPTCS_HUMAN
12675	STPLASPSPSGRSPQR200000002000000000	sp	Q8WYP5	ELYS_HUMAN
12676	TFSDENVL20020000	sp	Q8NB49	AT11C_HUMAN
12677	LSPVGVVSSSQGK200000000000000	sp	Q00587	BORG5_HUMAN
12678	VSPASPAGSPSADFAVHGESLDR2. 000000002E+23	sp	Q96C92	SDCG3_HUMAN
12679	SSEGGVGVGPGGGDEPPTS2E+20	sp	Q9Y4H2	IRS2_HUMAN
12680	HSPVQMN20000000	sp	Q0JRZ9	FCHO2_HUMAN
12681	SPPKSPEKLPQSSSESSPSPQPTK2. 0002E+26	sp	Q9UQ35	SRRM2_HUMAN
12682	NTPSQHSHSIQHSPER2020000000020000	sp	Q9NYF8	BCLF1_HUMAN
12683	GRSPDELPSAGDGGK20000000000000000	sp	060610	DIAP1_HUMAN
12684	TKSSSDLSPRR20020200200000	sp	Q9BRD0	BUD13_HUMAN
12685	RGSVSSVSPKPPSSFK2020000000000000	sp	Q9NZJ0	DTL_HUMAN
12686	HRSNSFS200000	sp	Q09666	AHNK_HUMAN
12687	RVSHSPPPKQR2020000000	sp	Q8IYB3	SRRM1_HUMAN
12688	YIASVQGSTPSR2002000	sp	P49792	RBP2_HUMAN
12689	RLSSTSLASGHSVR22002000000	sp	Q9BZL6	KPCD2_HUMAN
12690	TNSDSALHQSTMTPTQPESFSSGSQDVHQK2000000000000000000	sp	Q6UUU9	CRTC1_HUMAN
12691	GNULLHFPSSQGEEKEK20000000000	sp	Q12888	TP53B_HUMAN
12692	IEIPVPTGQSVSPSPSIPGTPLK2000000000000	sp	Q9H4Z3	PCIF1_HUMAN
12693	EPLENSASESSDTELPEKER2000220000000000	sp	P78337	PITX1_HUMAN
12694	TESVSDKEDKPLAPSGGTEGPEQPPPCPSQTGSPPVGLIK20200000000	sp	Q8WWM7	ATX2L_HUMAN
12695	VAEPGAEATSSSTGEESGSEHPPAVPMHNK2. 20002E+19	sp	Q14684	RRP1B_HUMAN
12696	GGAPEGPNEAEVTSKGPEQVPDAEEKSVSGTDVQECCR2E+27	sp	P49321	NASP_HUMAN
12697	EPALPLEPDSGGNTSPGVTANGEAR2000000000000000000	sp	Q92974	ARHG2_HUMAN
12698	SLAASSPPPCSQPSQCPEEKQESSPMK2000000000000000000	sp	Q7Z6J9	SEN54_HUMAN
12699	YPLCSADAAVSTTPPGSPPPPPLPEPPVLK2. 0000002E+27	sp	Q9BTC0	DIDO1_HUMAN
12700	RSLSPKPR20200000	sp	Q13523	PRP4B_HUMAN
12701	AGSSPCLGSSSAASSPPPPASR2002000000000	sp	Q6ZRS2	SRCAP_HUMAN
12702	LLGLTPPEGGADRSEEEASSTEKPTK200000000000000	sp	Q9UGY1	NOL12_HUMAN

12703	EVHDELEDLSPPPPPLSPPPTTSPHK2220000	sp Q9NQX3 GEPH_HUMAN
12704	EELTPGAPQHQLPPVPGSPEPYPGQQAPGPEPSQPLLELTPR3E+21	sp Q9H6F5 CCD86_HUMAN
12705	RFTPPSTALSPGK200000020000	sp Q01196 RUNX1_HUMAN
12706	SESGSPSAGGHSKPPHSPLVLK2000000000002000000	sp P01106 MYC_HUMAN
12707	VLHSPPAVR2000000	sp Q9Y4B5 MTCL1_HUMAN
12708	SMGTGDTPGLEVPSSPLRK200000	sp Q86WB0 NIPA_HUMAN
12709	RPGSVSSTDQER2200000	sp Q9H1B7 I2BPL_HUMAN
12710	SSSKESLVR20020000	sp Q13439 GOGA4_HUMAN
12711	SPQPVCSPAGSEGTK2000000	sp Q6KC79 NIPBL_HUMAN
12712	DSTSQHDDDNISTTSGFSSRASDKDITVSK2. 2220022002E+19	sp P49790 NU153_HUMAN
12713	KHHESITPPSRSPK2020000	sp Q8IY92 SLX4_HUMAN
12714	HSHKPHPVETSQPSDK20000000	sp Q03188 CENPC_HUMAN
12715	GQSQTSPDHRSDTSSPEVR2200002020000000	sp Q9UQ35 SRRM2_HUMAN
12716	GSPVSSPQNSPGTQR2000000	sp Q9P2P5 HECW2_HUMAN
12717	SVCPCGGSSPGSSSGGR200000	sp Q53LP3 SWAHC_HUMAN
12718	GHPAQRPEPSPPEPEQAPEGSQAGAEGPSSPEASR20000000	sp Q6ZW31 SYDE1_HUMAN
12719	QKPESDDSDRENKGEDTEMQNSDFHSDSHMDR2E+25	sp Q96ST2 IWS1_HUMAN
12720	SDGSGESAQPPEDSSPPASSESSSTR20000000000000	sp Q7Z6Z7 HUWE1_HUMAN
12721	DMESPTKLDVTLAK10002000000000	sp P27816 MAP4_HUMAN
12722	SPEETATQVPSLESITLK2000000000000000000	sp Q9HCK8 CHD8_HUMAN
12723	SSSVGSSSYPISPAISR23000000000	sp Q15149 PLEC_HUMAN
12724	SRSQSRYSADSYSSTR23000000000000	sp A7MD48 SRRM4_HUMAN
12725	DTLSNPQSPQSPYNSPKPKHK2. 000002000203E+21	sp Q96PU5 NED4L_HUMAN
12726	SALNLNQPVSVSSVPVK20000	sp Q86XJ1 GA2L3_HUMAN
12727	NATQPPNAEEESGSSSASEEEDTKPKPKR2. 0000000020222E+28	sp Q68E01 INT3_HUMAN
12728	DSAAPSPIPGADNLSADPVVSPPTSIPFK2E+24	sp Q5T1M5 FKB15_HUMAN
12729	GHSEIITNR20000000	sp P0C091 FREM3_HUMAN
12730	SPVSGDTSR2000	sp TAU_HUMAN
		sp P10636 TAU_HUMAN
12731	DTDMNKKDK201000000	sp Q6S545 POTEH_HUMAN
		sp A6NI47 POTEM_HUMAN
		sp Q6S5H5 POTEG_HUMAN
12732	KRESESEDETTPAAPQLIK2000020000000000	sp O60832 DKC1_HUMAN
12733	RFSPRR200000	sp Q15287 RNPS1_HUMAN
12734	LKTPSTPVACSTPAQLK2002000000000000	sp Q6PL18 ATAD2_HUMAN
12735	VVLGDGVQLPPGDYSTTPGGTLFSTTPGGTR200200000000	sp Q13541 4EBP1_HUMAN
12736	TPTSPLK20020000	sp O60264 SMCA5_HUMAN
12737	DMSLKDSGDLSHRPK200000000	sp O95251 KAT7_HUMAN
12738	KPGPPLSPEIRSPAGSPELR2000020000000000	sp Q96JM3 CHAP1_HUMAN
12739	EAFEQPQTSSTPPR20000	sp P52564 MP2K6_HUMAN
12740	IKPVTENLVLSHTGFCQSSGDEALSK2200000000	sp P46100 ATRX_HUMAN
		sp Q09160 1A80_HUMAN
		sp P05534 1A24_HUMAN
12741	GGSYSQAASSDSAQGSVSLTACK20002002000000	sp P30447 1A23_HUMAN
		sp P01891 1A68_HUMAN
		sp P01892 1A02_HUMAN
		sp P10316 1A69_HUMAN
12742	ARSRTPPSAPSQR20000000200	sp Q9UQ35 SRRM2_HUMAN
12743	SPAEGSERLSLPLR20002020000	sp Q66K74 MAP1S_HUMAN
12744	QGSITSPQANEQSVTPQRR200000000000200000	sp Q9UQ35 SRRM2_HUMAN
12745	SCTSPDQISHRASLEDAPVDDLTR2. 0200000020002E+25	sp Q7Z2W4 ZCCHV_HUMAN
12746	SPTAPSVFSPTGNR2000000000000000	sp Q9H3P2 NELFA_HUMAN
12747	VAHEPVAPPEDKESESEAKVDGETASDESER2000000000000200	sp O95674 CDS2_HUMAN
12748	FHALSSPQSPFPSTPTR20000000000	sp O43166 SI1L1_HUMAN

12749	HSSETFSSTTTVTPVSPSFAHNP2. 02E+22	sp	Q9Y4H2	IRS2_HUMAN
12750	YRPASVPPSPSLSR20002020000	sp	P13807	GYS1_HUMAN
12751	LSGEHSESSTPRPR20000002000000	sp	Q7Z3K3	POGZ_HUMAN
12752	SPAEGSERLSLSPLR2020000	sp	Q66K74	MAP1S_HUMAN
12753	LDSQPQETSPELPR20000000	sp	Q14545	TRAD1_HUMAN
12754	TESVSDKEDKPPLAPSGGTEGPEQPPPCPSQTGSPPVGLIK2. 0000000000 02E+24	sp	Q8WWM7	ATX2L_HUMAN
12755	LDSSEMDHSENYDTMSSPLPGK1000000032000000000	sp	Q9NTI5	PDS5B_HUMAN
12756	MMLTLISK200000	sp	Q07617	SPAG1_HUMAN
12757	SETSSNPSSPEICPNK22000000000000	sp	Q684P5	RPGP2_HUMAN
12758	SKSPLPPEEEAK2000000000000	sp	Q1KMD3	HNRL2_HUMAN
12759	TQPSSGVDSAVGTLPATSPQSTSVQAK200000000000	sp	Q13428	TCOF_HUMAN
12760	SSTPLPTISSAENTR22000000	sp	P42167	LAP2B_HUMAN
12761	SFSADGFEMLQNPVK20000000000000	sp	P42166	LAP2A_HUMAN
12762	VGPGNHGTEGSGGERHSDTSDRR200000000000000	sp	Q9HCM1	K1551_HUMAN
12763	SSSTEPPPPVR20200000000	sp	Q99442	SEC62_HUMAN
12764	TPTSSPASSPLVAK220000000000	sp	Q8N684	CPSF7_HUMAN
12765	VQIPVSRPDPPEVSDNEEDSYDEEIHDP2. 00000000000002E+24	sp	Q14684	RRP1B_HUMAN
12766	AAPEASSPPASPLQHLLPGK20000000000	sp	Q07157	ZO1_HUMAN
12767	MISTPSPK2002000	sp	Q96TA1	NIBL1_HUMAN
12768	TEMKSPFNPSQDSR2020002000	sp	Q53F19	CQ085_HUMAN
12769	DLDRPESQSPK202000	sp	P08651	NF1C_HUMAN
12770	TLSVAAAFNEDEDSEPEEMPPEAK200000000000	sp	Q8WUA4	TF3C2_HUMAN
12771	NCPSPVLIDCPHPNCNK200000000000000	sp	Q8WW12	PCNP_HUMAN
12772	DSFSHSPGAVSSLK200000000000	sp	Q15014	ZN609_HUMAN
12773	DSDYVYPSLESDENPIFK2000000000000	sp	Q9H6R7	CB044_HUMAN
12774	WATDQEDCSQDLAGTDLGPQK200000000000000	sp	Q07515	PHF2_HUMAN
12775	HIAFTPESQRR20020000	sp	Q07512	COBL_HUMAN
12776	KAPSASDSKADSDGAKPEPVAMAR2. 00020002E+21	sp	Q5H9R7	PP6R3_HUMAN
12777	NQSPTEAEKPASSSLPSSPPQLTR20000000000000	sp	Q7Z4V5	HDGR2_HUMAN
12778	IQSSLSVNSDISKK202000002000	sp	Q99567	NUP88_HUMAN
12779	SDSSGGYNLSDIIQSPSSTGLLK200002000000000	sp	Q9P1T7	MDF1C_HUMAN
12780	GNEPGSDRSPSPSK2002000000	sp	Q9P2D0	IBTK_HUMAN
12781	QGGSPSSPSPGQSQVTPQDQEK20000000000000000	sp	Q5VT52	RPRD2_HUMAN
12782	SGSYGRSPSPYGR200000000200000	sp	Q9H0L4	CSTFT_HUMAN
12783	DYSTLTSVSSHDSR2000000200200	sp	Q9NYV4	CDK12_HUMAN
12784	SEHSHSTLPR2220000	sp	P16144	ITB4_HUMAN
12785	GRSECDSSPEPK20002200000	sp	P16144	ITB4_HUMAN
12786	SPYGGGRPGR20000000000	sp	Q9UQ35	SRRM2_HUMAN
12787	TSPPCSPANLSR2000000000000	sp	P46781	RS9_HUMAN
12788	HSSISPVR20020000	sp	P49585	PCY1A_HUMAN
12789	EKSEGGSEYGTGPR202002000000000	sp	Q9NYV4	CDK12_HUMAN
12790	SEPLPTFNQFSPISSPEPEDPIKDNGFGIKPK2. 000020002E+27	sp	Q12873	CHD3_HUMAN
12791	APSASDSKADSDGAKPEPVAMAR2. 02000002E+21	sp	Q92610	ZN592_HUMAN
12792	QSISFSGLPGR200000000000	sp	Q7Z4V5	HDGR2_HUMAN
12793	EYVNSTSEESHDEDEIRPVQQDLHR2E+21	sp	P26651	TTP_HUMAN
12794	STSTPTSPGPR2000000	sp	Q8NBU5	ATAD1_HUMAN
12795	LKLSPSSSR20000200	sp	Q8WWM7	ATX2L_HUMAN
12796	LDSQEPGRQTPDR20000	sp	P20700	LMNB1_HUMAN
12797	SKLSPSPSLRK202000000	sp	Q03252	LMNB2_HUMAN
12798	GRSPPERVPVGP2000000002000	sp	Q8TB03	CX038_HUMAN
12799	KAPPPPISPTQLSDVSSPR2000020020000	sp	Q5T200	ZC3HD_HUMAN
		sp	Q9Y4B5	MTCL1_HUMAN
		sp	Q9P0K7	RAI14_HUMAN

12800	RWVGSSSPISKEK2200000000	sp Q96B77 TM186_HUMAN
12801	GNLHFPSSQGEEKEK2000000000	sp Q12888 TP53B_HUMAN
12802	LSLTSDEEGDPLALGPESGEPQPPQLK2E+25	sp Q5T1M5 FKB15_HUMAN
12803	SSPGAVAGLSNAPGTPR220000000000000000	sp Q9BWG4 SSBP4_HUMAN
12804	AALDGGVASAASPESKPSGTK200000000000	sp Q9Y4K1 AIM1_HUMAN
12805	VETVSQPSESPKDTIDK2000000000000000	sp Q99590 SCAFB_HUMAN
12806	AGSSGNSCITYQPSVSGEHK200000000000000000	sp Q99755 PI51A_HUMAN
12807	TPPHSHPAAPSTGSPVPATEPPAAK2000000020000000	sp Q9UKP4 ATS7_HUMAN
12808	TKCEILSDESSDEDEKK2220000000	sp P35659 DEK_HUMAN
12809	SPLPGPSEEK200000000000	sp Q13112 CAF1B_HUMAN
12810	KSRSPVDLR202000000	sp Q13523 PRP4B_HUMAN
12811	SYDVPPPPMEPDHPFYSNISK2E+21	sp Q8N0Y7 PGAM4_HUMAN
12812	QGQSQAASSSSVTSPIK2000000000	sp P18669 PGAM1_HUMAN
12813	NVSPEFVPCGEGGFLHK200000000000000000	sp O60583 CCNT2_HUMAN
12814	SSPINPAQSGSPNHVDSAYFPGSSTSSSSDNDEGSGGATNHISGNK2E+36	sp O94986 CE152_HUMAN
12815	SKSQSSGSSATHPIVPGAR2E+20	sp Q9N9ZB2 F120A_HUMAN
12816	SDGPASVVEGPKDPSPCK20000200000000000000	sp Q9Y4H2 IRS2_HUMAN
12817	NAVASVQNGPGGGPSSSSIGSASKSDESSTEETDK200220000000	sp Q15911 ZFH3_HUMAN
12818	VGSLTPPSSPK20002000	sp Q8NI27 THOC2_HUMAN
12819	CVSPIPVSPTR20000200000	sp Q2M2I8 AAK1_HUMAN
12820	KPVLSARNMFMFGLSPVRNPR2000200000020000000	sp Q86TC9 MYPN_HUMAN
12821	APSSPVAKPGPVK200000000000	sp A6NL46 YF016_HUMAN
12822	THSVPATPTSTPVNPEAESK2.0002E+21	sp Q9GZR2 REX04_HUMAN
12823	LHSYSSPSTK230000000	sp Q96FF9 CDCA5_HUMAN
12824	WAHDKFSGEEGEIEDDESSTENREEK2.000000000002E+20	sp Q96T37 RBM15_HUMAN
12825	APSASDSKADSDGAKPEPVAMAR2.0000020002E+23	sp Q9Y2W1 TR150_HUMAN
12826	VGPGNHGTEGSGGERHSDTSDRR202020000	sp Q7Z4V5 HDGR2_HUMAN
12827	QEDSVTELTVEDSGESLEDLMAK2E+20	sp Q99442 SEC62_HUMAN
12828	VRPSSDLNSTGQSPHHK2000000000000000	sp Q9GZZ9 UBA5_HUMAN
12829	YQEDSDPERSDYEEQLQK200000000000000000	sp P27448 MARK3_HUMAN
12830	ISDSESEDPPRHQASDSENEELPKPR2.000000000002E+25	sp Q6P6C2 ALKB5_HUMAN
12831	AQPSSSEDELNVFFK2220000000000000	sp Q96ST2 IWS1_HUMAN
12832	VGDSTPVSEKPVSAVDANASESP20200	sp Q5T5U3 RHG21_HUMAN
12833	ASSHSSQTQGGGVTKK2020000000000000	sp Q9H8Y8 GORS2_HUMAN
12834	ADSGTPPPLSLSPAPETK2020000000	sp P02545 LMNA_HUMAN
12835	VAEPGAEATSTGEEGSEHPPAVPMHNK2.000002E+20	sp O15027 SC16A_HUMAN
12836	NVIPDTPPSTPLVPSR200000000000	sp Q14684 RRP1B_HUMAN
12837	SGTTPKPVINSTPGR2000000020000	sp Q9UKA4 AKA11_HUMAN
12838	YRPENTPEPVSTSVSHYGAEPSTVSPCPSSSAK2000200000	sp Q99459 CDC5L_HUMAN
12839	DMSPLSETEMALGK2000020000000	sp P07947 YES_HUMAN
12840	HSCSGSPPR2000020000	sp P27816 MAP4_HUMAN
12841	VLLSPSIPSVGDDQTLSPGSK200000000000000000	sp Q9UQ35 SRRM2_HUMAN
12842	IEVLDSASK200000	sp Q13625 ASPP2_HUMAN
12843	KPIETGSPK200000	sp Q8N5I9 CLO45_HUMAN
12844	SKSPSPLKNK20202000000	sp Q05519 SRS11_HUMAN
12845	VEAVEEPEAAPILGPGEPGPAPSRTPSR20202000	sp O94913 PCF11_HUMAN
12846	GYSSPEPDIQDSSGSEAQSVKPSTR3.000000000002E+24	sp Q8IX07 FOG1_HUMAN
12847	DSELSDTDSGCCLGQSESDK2000000000000000	sp Q9BZF1 OSBL8_HUMAN
12848	SMSVDETDKSPCEAGR2000000000000000	sp Q96T88 UHRF1_HUMAN
12849	DNPAQDFSTLYGSSPLER200000	sp Q53SF7 COBL1_HUMAN
12850	SAGGRPGSGPQLGTGR2000000000	sp Q15648 MED1_HUMAN
12851	NINMDNDLEVLTSPPTR20000	sp O14908 GIPC1_HUMAN
12852	QATPGVPAQQSPSM20000	sp Q9Y5T5 UBP16_HUMAN
		sp O60341 KDM1A_HUMAN

12853	SPSQESIGAR200000000	sp Q8WXE0 CSKI2_HUMAN
12854	SKSPASVDR2020020000	sp Q13243 SRSF5_HUMAN
12855	SSSPIPLTPSK20000000000	sp Q8N5Y2 MS3L1_HUMAN
12856	SPASDSETEDALKPQISDSESEEP2. 020200000002E+23	sp Q96ST2 IWS1_HUMAN
12857	MPGMSPANPSLHSPVPDASHSPR200200000002000	sp Q06244 MED14_HUMAN
12858	SKSEDSALFTHHSLGSEVVK200000000000	sp Q99708 COM1_HUMAN
12859	SLPPQLLEGPSSTPSSHSDTR2000000000000000	sp Q6ZRS2 SRCAP_HUMAN
12860	VETVSPSESPKDTIDK20000000000000	sp Q99590 SCAFB_HUMAN
12861	SRSPSHTRPR20000020000	sp Q8IYB3 SRRM1_HUMAN
12862	IEEVLSPEGSPSKSPSKK20002000200000	sp Q9UEY8 ADDG_HUMAN
12863	RLSGSEDEEDSGK2022000000000	sp Q9BW71 HIRP3_HUMAN
12864	SPVGKSPSTGTYGSSQK200000000000	sp Q9Y2W1 TR150_HUMAN
		sp Q29960 1C16_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
12865	GGSCSQAASSNSAQGSDESLIACK20002000000000	sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
		sp P30510 1C14_HUMAN
12866	SRSIDKDR202000000	sp Q8TF01 PNISR_HUMAN
12867	SPASDSETEDALKPQISDSESEEP2. 00202E+26	sp Q96ST2 IWS1_HUMAN
12868	QRSDDESPTSSGSSDADQRDPAAPEPEEQEER2E+31	sp Q969T4 UB2E3_HUMAN
12869	TSSPPSPQHSPALR2200000000200000	sp Q13029 PRDM2_HUMAN
12870	ALSPGRESFK200002000	sp Q13207 TBX2_HUMAN
12871	SHTSEGAHLDITPNSGAAGNSAGPK2. 00000000002E+25	sp Q92597 NDRG1_HUMAN
12872	TVHPNSPGIPYR20000000	sp Q8WUI4 HDAC7_HUMAN
12873	VQNHLPASGPTQPPVVSSTNEGSPSPPEPTGK20200000000	sp Q043294 TGF11_HUMAN
12874	SRSHDRLPYQR202000000000	sp P49761 CLK3_HUMAN
12875	YETPSDAIEVISPASSAPPQEK2200000000	sp Q075376 NCOR1_HUMAN
12876	EGMNPYSDEYADSDEDQHDAYLER30000000000000000000	sp Q08945 SSRP1_HUMAN
12877	KRESESEDETTPAAPQLIK20000000000000	sp Q060832 DKC1_HUMAN
12878	IFDEVMGCFSDSPQSPTFPEAGHTSLYDEDK2. 000002E+23	sp Q13085 ACACA_HUMAN
12879	RPMSSPEMK2000000	sp P51610 HCFC1_HUMAN
12880	KAPPPPISPTQLSDVSSPR2000020022000	sp Q9P0K7 RAI14_HUMAN
12881	DISGSPSKK2000	sp Q96GX9 MTNB_HUMAN
12882	TPASTPVSGTPQASPMIER20000000000010000	sp Q9UHR4 BI2L1_HUMAN
12883	LIPLDEECMSDETYPVETLEPGGSGGSPDGAGGSK2000000000	sp Q96QC0 PP1RA_HUMAN
12884	KADTEGKSPSHSSNRPFPTSTGGSK2. 02E+19	sp Q15648 MED1_HUMAN
12885	ASSPTKTSPTTPEASATNSPCTSKPATPAPSEK2. 2022E+27	sp Q12873 CHD3_HUMAN
12886	YPFKSPSIGFMNK30002000001000	sp P62256 UBE2H_HUMAN
12887	ILSDVTHSAVFGVPASK2000000000000	sp Q2M2I8 AAK1_HUMAN
12888	ALSSGGITSPPLSPALPK20000000000000000000	sp Q6Y7W6 PERQ2_HUMAN
12889	MKTPVQYSQQNSPQK2000300000000000	sp P46013 KI67_HUMAN
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12891	LDSPTMSR20000	sp P19634 SL9A1_HUMAN
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12893	TASPLPLRPDAPVEK2000000000000000	sp Q9HCK8 CHD8_HUMAN
12894	TKSPKPAESPQSATK20000020000000	sp Q5VUA4 ZN318_HUMAN
12895	VGPGNHGTEGSGGERHSDTSDR20200000	sp Q99442 SEC62_HUMAN
12896	SPQGQLDTSESKPDSFFLEPLMPAVLKPAK2E+30	sp Q5T5Y3 CAMP1_HUMAN

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12898	DQRSPSPIR200000	sp Q5T200 ZC3HD_HUMAN
12899	RPTETNPVTSNSDEECNETVK2000	sp P46100 ATRX_HUMAN
12900	TSPTTPEASATNSPCTSKPATPAPSEK2.000202E+23	sp Q12873 CHD3_HUMAN
12901	LKSSTSFANIQENSN220000000000	sp Q86WC4 OSTM1_HUMAN
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12903	SLTWPPSGSPSHAGTTPPENGLSEHPCETEQINAK2.02E+29	sp P23497 SP100_HUMAN
12904	RPFLSGDEDPK20000000	sp Q8WYR1 PI3R5_HUMAN
12905	STLESEKPGSPEAAETSPPSNIIDHCEK2000000000000	sp Q96T23 RSF1_HUMAN
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12907	LNNTPDSDWLQLKPLAGTDDSVVSEDR20020000	sp Q9ULF5 S39AA_HUMAN
12908	LSPEPGR2000000	sp Q9NWK9 BCD1_HUMAN
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12910	DTLSNPQSPQSPYNSPKPKQHK2000003000000000	sp Q96PU5 NED4L_HUMAN
12911	SPGSTPTTPTSSQAPQK200200000002000000	sp P35658 NU214_HUMAN
12912	LALSPEDKPIRLSPSK20000000000200	sp O15417 TNC18_HUMAN
12913	SYSPDGKESPSDKK30000002000000	sp P43243 MATR3_HUMAN
12914	GHSPTLRPK200000000	sp Q53LP3 SWAHC_HUMAN
12915	HGSFVNKPTR200000000	sp P29353 SHC1_HUMAN
12916	RSSLSPSSAYER2002000000000	sp Q9BYW2 SETD2_HUMAN
12917	TDGSISGDRQPVTVADYISR200000000000000000	sp P51116 FXR2_HUMAN
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		sp P01892 1A02_HUMAN
12919	GGSYSQAASSDSAQGSVDVSLTACK2.3E+22	sp P05534 1A24_HUMAN
		sp Q09160 1A80_HUMAN
		sp P10316 1A69_HUMAN
		sp P30447 1A23_HUMAN
12920	SPQEGPTWSR20000000000	sp Q9Y4F5 C170B_HUMAN
12921	SKEFVSSDESSGENK22200000	sp Q08945 SSRP1_HUMAN
12922	EWNGVSESDSPVKRPGR2000200000000	sp P18858 DNL11_HUMAN
12923	LTPSPDIIVLSDNEASSPR2000	sp Q8WXI9 P66B_HUMAN
12924	EVTPILAQTQSSGPQTSTTPQVLSPTITSPNALPR2E+34	sp P30260 CDC27_HUMAN
12925	RSSLPPNSPKPVR200002000000	sp Q7Z401 MYCPP_HUMAN
12926	AGPDLPSLPSHALEDEGWAAAAPSPGSAR2E+20	sp Q9H3Q1 BORG4_HUMAN
12927	GFYPQYGALETR30000000200	sp Q63ZY3 KANK2_HUMAN
12928	SYSPDGKESPSDKK2000000	sp P43243 MATR3_HUMAN
12929	VENSPQVDGSPGLEGLLGGIGEK2E+21	sp Q8WVB6 CTF18_HUMAN
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12934	GWGSPGRPK2000000	sp A7E2V4 ZSWM8_HUMAN
12935	SAPSSPTLDCEK2002000000000	sp Q9GZM8 NDEL1_HUMAN
12936	ASAGPAPGPVVTAEGLHPSLPSPTGNSTPLGSSK2000000000000	sp Q4KMP7 TB10B_HUMAN
12937	GPTTGEALDLSDVHSPPKSPEGK20002000000000	sp O95684 FR10P_HUMAN
12938	TGSNAAQYK20000000	sp Q8N556 AFAP1_HUMAN
12939	TSCKDDEAVVQAPR20000000000000	sp P35658 NU214_HUMAN
12940	VAEPGAEATSSTGEESGSEHPPAVPMHNK2.2020002E+21	sp Q14684 RRP1B_HUMAN
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12942	KPLTSSSAAPQRPISTQR2000000000000000	sp Q15691 MARE1_HUMAN
12943	NMVDLVNTHHLHSSSDEDDRLK2000000000000000	sp Q9UPN7 PP6R1_HUMAN
12944	SDSDSSTLPR20000	sp Q9ULE0 WWC3_HUMAN
12945	KSTYWMKMR23000100	sp Q96ME7 ZN512_HUMAN

12946	SYESSEDCSEAAGSPAR20000	sp Q6P6C2 ALKB5_HUMAN
12947	RQSSGSATNVASTPDNR2000020000000000	sp Q7Z460 CLAP1_HUMAN
12948	EMEHNTVCAAGTSPVGEIGEEK200000000000000000	sp P18583 SON_HUMAN
12949	SPSEESAPTTSPESVSGSVPSSGSSGR2E+22	sp P54725 RD23A_HUMAN
12950	KTSIKSHMSGSPGPGSNTAPSTPVIIGSDKPGMEEK2.02000002E+29	sp A0FGR8 ESYT2_HUMAN
12951	LSYLPATVEPASPTPAHSLPR20000000000	sp Q9NRA0 SPHK2_HUMAN
12952	NHKPSRSFSR2020000	sp P84103 SRSF3_HUMAN
12953	RYPSSISSSPQK2000020000	sp Q14157 UBP2L_HUMAN
12954	SSTPPRQSPSR22000020000	sp Q9UQ35 SRRM2_HUMAN
12955	SGPSPLSSPNGK2200000	sp O75179 ANR17_HUMAN
12956	YPSSISSSPQK200020000	sp Q14157 UBP2L_HUMAN
12957	TVTPASSAKTSPAK200220000	sp Q16555 DPYL2_HUMAN
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12959	VGDPPQPLPEEPMEVQGAERASPEPQR2000000	sp P46379 BAG6_HUMAN
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12963	DVYLSR2000	sp Q96E39 RMXL1_HUMAN
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12970	GSKPSPLSVK2002000	sp O15027 SC16A_HUMAN
12971	SSLGSHQLPR22000000000	sp Q96S94 CCNL2_HUMAN
12972	SQSLPTLLSPVR20000000020000	sp O95817 BAG3_HUMAN
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12977	REESPMDVDQPSAQTQSIASDGTQGEK1.00000000000002E+26	sp Q15691 MARE1_HUMAN
12978	DSQSASGEERPPEADGK2000000000000	sp Q7Z6Z7 HUWE1_HUMAN
12979	HLGGSGSVVPGSPCLDR2000000	sp P48634 PRC2A_HUMAN
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12981	LQPSSPENSLDPFPPR200000000	sp P10636 TAU_HUMAN
12982	AQPFIDSDTDAEEER20000000	sp TAU_HUMAN
12983	LDSSPVSSPR220000000	sp Q8WWM7 ATX2L_HUMAN
12984	RDSSESQLASTESDKPTTGR200000000	sp Q14676 MDC1_HUMAN
12985	RNSVERPAEPVAGAATPSLVEQK2.000000000002E+22	sp O60293 ZC3H1_HUMAN
12986	RRSPSLSSK20002000	sp Q96B23 CRO25_HUMAN
12987	SSTATHPGPAVQLNK200000000000	sp Q9BTC0 DIDO1_HUMAN
12988	YMSGQTSR2000	sp Q8IYB3 SRRM1_HUMAN
12989	ATLGPTPTTPPQPPDPSQPPPGPMQH200000000000000000	sp Q14684 RRP1B_HUMAN
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12991	YQESPGIQMK20000000	sp O60927 PP1RB_HUMAN
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12993	DLHQPSLSPASPHSQGFER2002000000000	sp Q9UHI6 DDX20_HUMAN
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12995	TLLSESSSQSSKSPSLSSK20000000	sp Q9BZF1 OSBL8_HUMAN
12996	LEPVSPSPPHTDPELELVPPR2000200000000000	sp Q9BTA9 WAC_HUMAN
12997	VLSSAASLPGSELSSRPEGSQGGELSR2E+26	sp Q9UPU5 UBP24_HUMAN
		sp Q9Y618 NCOR2_HUMAN
		sp O15027 SC16A_HUMAN



12998	SSQQPSTPQQAPPGQPQQGTFVAHK2E+19	sp	Q86VM9	ZCH18_HUMAN
12999	RHSSVSDSQCEPPSVGTEYSQGASQPQHQK2E+31	sp	P39880	CUX1_HUMAN
13000	AEPPKPEVVDSTESIPVSEDESDAMVDDPNDEDFVFRPR2. 00000002E+27	sp	Q14202	ZMYM3_HUMAN
13001	EPEPAAGPQAEESATVSAPAPMSPTR20000	sp	Q9BWG6	SCNM1_HUMAN
13002	IGNEESDLEEACILPHSPINVDK2000000000020000000	sp	O95714	HERC2_HUMAN
		sp	Q9BVR0	HRC23_HUMAN
13003	SGSPSDNSGAEEMEVSLAKPK2. 0000002E+21	sp	P31749	AKT1_HUMAN
13004	LQSEPEsir20000000	sp	P09496	CLCA_HUMAN
13005	TASETRSEGSEYEEIPKR2030000000	sp	P48634	PRC2A_HUMAN
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13007	KVSSSPQSGCPSPTIPAGK2000000000002000000	sp	Q76L83	ASXL2_HUMAN
13008	VQRPHSSPPR20000	sp	Q9NUJ3	T11L1_HUMAN
13009	SDGACDSPSSDKENSSQIAQDHQK22000000000	sp	Q9UIG0	BAZ1B_HUMAN
13010	GTSPSSSRPQR20000000000	sp	Q8N1F7	NUP93_HUMAN
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13013	GPGPGQSGPGGPGPR2000000000	sp	Q8IYL2	TRM44_HUMAN
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13015	TPLSSLSATSNNSTVACIDR20000000	sp	P00533	EGFR_HUMAN
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13017	SEVQQPVHPKPLSPDRASSLSESSPPK22020200000	sp	Q9UHB6	LIMA1_HUMAN
13018	STPESGSDKESVGSSTSNEGGR20000000000000	sp	Q6GQQ9	OTU7B_HUMAN
13019	IPGSPPEMGR200000000	sp	P50395	GDIB_HUMAN
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13026	GNLELRPGGAHPGTCSRPGS20000000	sp	Q96SN8	CK5P2_HUMAN
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13040	KPGSLFAALMATATSSLR1000022000	sp	Q96J66	ABCCB_HUMAN
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13042	TSGGDHAPDSPSGENSPAPQGR20000020000000	sp	Q96NY9	MUS81_HUMAN
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		sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30505 1C08_HUMAN
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		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
		sp P30510 1C14_HUMAN
13047	AISSSDSSDEDK20200220000000	sp Q6PD62 CTR9_HUMAN
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13049	SPSPASTSTVPMNTDAPTAPK2.00000000001E+21	sp Q659A1 ICE2_HUMAN
13050	AESSSSISSASLSAANTPTVGVSR2.2022E+21	sp Q0JRZ9 FCHO2_HUMAN
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13054	EYIPGQPPLSQSSDSSPTR20020000000	sp P07814 SYEP_HUMAN
13055	ITSYFLNEGSQARPR23000000000000	sp O75792 RNH2A_HUMAN
13056	LCSSEVGLLK20000000	sp Q9NWC5 TM45A_HUMAN
13057	YTPTSPSYSPPSPEYTPSPK2000000000030000000	sp P24928 RPB1_HUMAN
13058	GSGVAQSPQQPPPPQQQQQQPPQPTPPK20000	sp Q6VY07 PACS1_HUMAN
13059	SPVVSQDTSR200020002000	sp TAU_HUMAN
		sp P10636 TAU_HUMAN
13060	RGTGSESSRASGLVGR2002002002000000	sp Q9UH99 SUN2_HUMAN
13061	VVFSSDESSEEDLSAEMTEDETPKK2.2002200002001E+22	sp Q15021 CND1_HUMAN
13062	EGQSPADEKGNDSGEGESDDPEK2000000000000	sp O75351 VPS4B_HUMAN
13063	ISDSESEDPPRHQASDSENEELPKPR2.00000000202E+21	sp Q96ST2 IWS1_HUMAN
13064	RSSPSARPPDVPGQQPQAAK20000000000000000	sp Q96JP5 ZFP91_HUMAN
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13067	ISDSESEDPPRHQASDSENEELPKPR2.0200000000202E+23	sp Q96ST2 IWS1_HUMAN
13068	SRSRTPPAIR202000000	sp Q9UQ35 SRRM2_HUMAN
13069	RFSSGGEEDDFDR220000000000	sp O94929 ABLM3_HUMAN
13070	LVKPSSSSSGSRSSR2200	sp Q9H6B4 CLMP_HUMAN
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13072	SSPELPPSGGTTSGSR20000020000000000	sp Q96D31 CRCM1_HUMAN
13073	NAASFPLRSPQVPCSPAGSETPK2.00000000002E+21	sp Q6KC79 NIPBL_HUMAN
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13075	SSPVPVSDTQK20000000000	sp Q76FK4 NOL8_HUMAN
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		sp Q13595 TRA2A_HUMAN
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13078	HSTPSNSSNPSGPPSPNSPHR20000000000000000	sp Q9Y5S2 MRCKB_HUMAN
13079	TCPSLSPTSPLNNK202002000000	sp O75925 PIAS1_HUMAN
13080	TGSNISGASSDISLDEQYK200020000000000	sp P22059 OSBP1_HUMAN
13081	RLEISPDSSPER200020000	sp Q99959 PKP2_HUMAN
13082	EVHDELEDLSPPPPPLSPPTTSPHK20000000002020000	sp Q9NQX3 GEPH_HUMAN
13083	QISYNYSDLQSNVTEETPEGEEHHPVADTENK3E+30	sp Q8NC51 PAIRB_HUMAN
13084	ISDSESEDPPR202000000	sp Q96ST2 IWS1_HUMAN
13085	FTDTRKDEQER20200000000	sp P13639 EF2_HUMAN
13086	ASSESSYLSILPK200000000000	sp Q86SQ0 PHLB2_HUMAN
13087	VASPSQGVGSSSPKR200000	sp Q99569 PKP4_HUMAN

13088	VAHSDKPGSTSTASFR200000	sp Q9P0L0 VAPA_HUMAN
13089	RSPSPLGTSVR20200000000	sp Q86SQ0 PHLB2_HUMAN
13090	KPEKPLFSSASPQDSSPR220000000000	sp 000712 NFIB_HUMAN
13091	RDSL TGSSDLYK200000	sp Q14671 PUM1_HUMAN
13092	IEEVLSPEGSPSKSPSK2000200000000	sp Q9UEY8 ADDG_HUMAN
13093	ADAPDAGAQSDELPSYHQNDVSLDR2000000000020000	sp Q92538 GBF1_HUMAN
13094	SSSKGSVEEIMSQPK2200020000100000	sp Q8IWB9 TEX2_HUMAN
13095	RLPGESGSCEDGQSAPAPPR202000000000000000	sp Q6UXQ8 Y0002_HUMAN
13096	LFVTPPEGSSR2000000000	sp Q14244 MAP7_HUMAN
13097	SPSHSSNRPTPTSTGGSK200000000000000000	sp Q15648 MED1_HUMAN
13098	KPSVSEEVQATPNK200000000000	sp Q9UKJ3 GPTC8_HUMAN
13099	HKSDSPESDAEREK200200000000	sp 075400 PR40A_HUMAN
13100	YESQLSTNEEK200000	sp Q13433 S39A6_HUMAN
13101	SPGMEPPFHR2000000000000000	sp Q99973 TEP1_HUMAN
13102	KLEGNPQGSNGVK200020000000	sp P61006 RAB8A_HUMAN
13103	SSSPCRTPEPDNDAHLR200020000000000000	sp Q9Y2K6 UBP20_HUMAN
13104	GKDSLDDGVDLK202000000000	sp P07948 LYN_HUMAN
13105	DGSDYDPYDFSDTEEEMPQVHTPK2020001000000000	sp P25205 MCM3_HUMAN
13106	TSSDPNLNHCQEV200000000000000000	sp Q9NYA4 MTMR4_HUMAN
13107	TPASTPVSGTPQASPMIER2000000002010000	sp Q9UHR4 BI2L1_HUMAN
13108	TDSEKPFRRSQSPKR2020000	sp Q9Y2W1 TR150_HUMAN
13109	ESAAPASPAPSPAPSPTAPPQK2.00002E+22	sp Q3KQU3 MA7D1_HUMAN
13110	YTPTSPSYSPSSPEYTPSPK2020000000000000	sp P24928 RPB1_HUMAN
13111	NMVDLVNTHLHSSDDEDDLK1.000000000022E+22	sp Q9UPN7 PP6R1_HUMAN
13112	SVPVNNLPERSPDSDPR2000000000	sp P49454 CENPF_HUMAN
13113	LPEPSVLSEVTKPPPCLNSSPPSR200000	sp Q9BZL4 PP12C_HUMAN
13114	EVQTPQYLNPFDPEAFVTIKDSPPQSTK200	sp Q8NDI1 EHBP1_HUMAN
13115	GHPAQRPEPSPPEPEPQAPEGSQAGAEGSSPEASRSPAR2E+19	sp Q6ZW31 SYDE1_HUMAN
13116	AKSPTSPSPPRNSDQEGGK2.02020000002E+19	sp Q8IYB3 SRRM1_HUMAN
13117	FSTYTSKDKENKLEASGGR20020000	sp Q8WU90 ZC3HF_HUMAN
13118	GPSQEEEDNHSDEEDRASEPKSQDQDSEVNLSR2.00000020002E+24	sp Q86VM9 ZCH18_HUMAN
13119	LQDSSDPDTGSEEEGSSRLSPPHSPR2.2E+23	sp Q92974 ARHG2_HUMAN
13120	YTPTSPSYSPSSPEYTPSPK20000000000000000000	sp P24928 RPB1_HUMAN
13121	LQDSSDPDTGSEEEGSSRLSPPHSPR2.2000202E+23	sp Q92974 ARHG2_HUMAN
13122	GSDALSETSSVSHIEDLEK20002020000000000000	sp Q6P996 PDXD1_HUMAN
13123	IEHAPSPSSGGTLK20000000	sp Q14004 CDK13_HUMAN
13124	ELVEPLTPSGEAPNQALLR20000000000000	sp P00533 EGFR_HUMAN
13125	EAAVSASDILQESAIHSPGTVEK2E+19	sp Q96C57 CL043_HUMAN
13126	SWEDSPER200020000	sp Q9UDY2 ZO2_HUMAN
13127	QLGDSSGEGPEFVEEEVALRSDSEGSDYTPGK2E+29	sp Q14839 CHD4_HUMAN
13128	VWSPLVTEEGK20000000000	sp 000151 PDL1_HUMAN
13129	TSSAVWNSPPLQGAR2200002000000000	sp Q9NQS7 INCE_HUMAN
13130	SGGGFGPDGSGTTVPAR20000000000000000000	sp Q9H4L4 SENP3_HUMAN
13131	ESYTEEIVSEASHVSGISR2.00000000002E+19	sp Q5QJE6 TDIF2_HUMAN
13132	FSSPTEELDYR20000000000	sp 095359 TACC2_HUMAN
13133	LSSAGPRSPYCK200000200000	sp Q6PL18 ATAD2_HUMAN
13134	AQGPAASAEKPKVEAPAANSQDQTVTVKE2000000	sp P80723 BASP1_HUMAN
13135	HLDGEEDGSSDQSQASGTTGGR200000000000	sp Q9UNF1 MAGD2_HUMAN
13136	LDSSEMDSSENYDTMSSPLPGK2.0010000000021E+21	sp Q9NTI5 PDS5B_HUMAN
13137	SPSPPDGSPAATPEIR202000020000000000	sp 000499 BIN1_HUMAN
13138	QQPPEPEWIGDGESTSPSDK20000000	sp Q8NE71 ABCF1_HUMAN
13139	SAWSPELFS20	sp Q8IXT1 DDIAS_HUMAN
13140	KRSPSPSPTPEAK2020200000000	sp Q8TAQ2 SMRC2_HUMAN
13141	GKSSEPVVIMK2000000000	sp P46013 KI67_HUMAN

13142	YPSSISSSPQKDLTQAK200020000000000	sp Q14157 UBP2L_HUMAN
13143	SRTSPVTR200000000	sp Q9UQ35 SRRM2_HUMAN
13144	SRSLVDYENANK20000000000	sp Q9UNH7 SNX6_HUMAN
13145	EGEETEGSEEDDKENDKTEEMPNDVLENK2.000000002E+23	sp P49321 NASP_HUMAN
13146	STSAPQMSPGSSDNQSSSPQPAQQK1.000020002E+19	sp Q14157 UBP2L_HUMAN
13147	EYIPGQPPLSQSSDSSPTR20000020000	sp P07814 SYEP_HUMAN
13148	SRSRSPTPSSAGLGSNSAPPIDSR2.020022E+22	sp Q8IWX8 CHERP_HUMAN
13149	SLENETLNKEEDCHSPTSKPPKPDQPLK2.000000002E+23	sp P23588 IF4B_HUMAN
13150	TPSSEEISPTKFPGLYR200000200000000	sp Q15311 RBP1_HUMAN
13151	MPDEPEEPVVAVSSPAVPPPTK20000000000	sp O60885 BRD4_HUMAN
13152	VAFDPEQKPLHGVLKPTTSSPASSPLVAK220020000000	sp Q14684 RRP1B_HUMAN
13153	SRSGSSQELDVKPSASPQR2.02E+20	sp Q9UQ35 SRRM2_HUMAN
13154	GEQVSQNGLPAEQGSPPR2000	sp Q01082 SPTB2_HUMAN
13155	SIPLECLSSPK2000	sp Q92667 AKAP1_HUMAN
13156	VVSSEGGPSVDLSFQPSQPLSK20000000000	sp P49815 TSC2_HUMAN
13157	SRVSVSPGR2002020000	sp Q8IYB3 SRRM1_HUMAN
13158	RVPGSSGHLHK22000000	sp Q9UEW8 STK39_HUMAN
13159	SLENETLNKEEDCHSPTSKPPKPDQPLK2.00002E+28	sp P23588 IF4B_HUMAN
13160	LSPYPSSPMR30020000	sp O94967 WDR47_HUMAN
13161	RHSSVSDSQCEPPSVGTEYSQGASPPQHQHLK2.02000000000003E+28	sp P39880 CUX1_HUMAN
13162	KPPGTPPPSALPLGAPPPDALPPGPTPEDSPSLETGSR2E+30	sp Q9BQQ3 GORS1_HUMAN
13163	ISDSESEDPPRHQASDSENEELPKPR2.0202E+25	sp Q96ST2 IWS1_HUMAN
13164	AQTTPGPSLSGSKSPCPQEK2000000000000000000	sp Q9UQ35 SRRM2_HUMAN
13165	DDGDEEGLLTHSEEELEHSQDADADDGALQ2E+19	sp Q9UKM9 RALY_HUMAN
13166	RESESEDETPPAAPQLIKK2000200200000000000	sp O60832 DKC1_HUMAN
13167	QRSHSGTSPDNTAPPPPPRPPQPSHSR2.20002E+21	sp Q96D71 REPS1_HUMAN
13168	HIVNHDDVFEESEELSSDEEMK22000100	sp Q96RT1 LAP2_HUMAN
13169	LNETELTDLEGQESPPKNYLCIEEEK2E+24	sp Q9ULF5 S39AA_HUMAN
13170	NATQPPNAEEESGSSSASEEEDTKPKPTK2.000000002022E+27	sp Q68E01 INT3_HUMAN
13171	DILKPSPGKSPGTRPEDEAEGKPPQR2.0002E+21	sp P31641 SC6A6_HUMAN
13172	TRSWDSSSPVDRPEPEAASPTR2E+21	sp Q86WB0 NIPA_HUMAN
13173	DEPQTVPDVPSFGESPLSPIDMDTQER2000000020000000000	sp P17535 JUND_HUMAN
13174	RQPTVHFSDVSSDDDED200220000000	sp P15822 ZEP1_HUMAN
13175	RSFISPV202000000	sp P18583 SON_HUMAN
13176	SISSDGRPLER2000000000	sp O43166 SIIL1_HUMAN
13177	HASPDKWNDKPK20000000000	sp Q96JM3 CHAP1_HUMAN
13178	ATSVKGYTGANQSR2000300000000	sp Q5VTH9 WDR78_HUMAN
13179	SNPEGAEDR2000000000	sp Q9NWX8 BABA1_HUMAN
13180	ISETLAGIYDDNSLSQDFPGDESPPDAQPTQPTPAPGTPPQVDSTR2.0000002002E+25	sp Q9H7N4 SFR19_HUMAN
13181	ESDKEDGRESYDTPSQR20000000000	sp Q9Y6M7 S4A7_HUMAN
13182	GVPAGNSDTEGGQGR200000000	sp Q9UKV3 ACINU_HUMAN
13183	RSSLSSHSHQSIYR20020000000000	sp O15027 SC16A_HUMAN
13184	NKSNEDQSMGNWQIK2000001000000	sp P02545 LMNA_HUMAN
13185	LEDAFIVISDSGEEPKEENGLQK20200000000000000	sp Q96RL1 UIMC1_HUMAN
13186	TSIDSIDSGVELTTPK20020000000000	sp O15403 MOT7_HUMAN
13187	SPPADPPR200000000	sp O60566 BUB1B_HUMAN
13188	LSPLPGPGAGDPR20000000000000	sp Q15742 NAB2_HUMAN
13189	GLSLSPQSPERR200002000000	sp Q7Z2Z1 TICRR_HUMAN
13190	TVIIHDRPDITHPR200000000020000	sp Q9NWH9 SLTM_HUMAN
13191	ATLSNPKPK20000000	sp Q96JM3 CHAP1_HUMAN
13192	VQMTSPSSTGSPMFK2200000001000	sp P49790 NU153_HUMAN
13193	TANSPGSGARPDPVR2000000000000	sp Q13206 DDX10_HUMAN

13194	TASISSSPSEGTPTVGSYGCTPQSLPK2E+27	sp	Q6PKG0	LARP1_HUMAN
13195	SGYSSPGSPGTPGSR200000000000	sp	P10636	TAU_HUMAN
13196	SMVSPVPSPTGTISVPNSCPASPR1.000000000202E+23	sp	TAU_HUMAN	
13196	SMVSPVPSPTGTISVPNSCPASPR1.000000000202E+23	sp	P85037	FO XK1_HUMAN
13197	QHSNPGHAGFPVSVHNTTINPTSPTTATMANANR200020000000000000	sp	Q9UPN9	TRI33_HUMAN
13198	HRSLSPVPR20200000	sp	A2AJT9	CX023_HUMAN
13199	SPESQNLIDGTK20000000000000	sp	Q9HCE3	ZN532_HUMAN
13200	QRGSETDSEIHEASDKDSLK2.020202E+21	sp	Q9Y520	PRC2C_HUMAN
13201	LNLAPAAATTSNEPSGNNPPTHLSLDPTNAENTASQSPR200000	sp	Q9UER7	DAXX_HUMAN
13202	QSHSGSISPYPK2000200000	sp	Q9UQ35	SRRM2_HUMAN
13203	GPEENYSRPEAPNEFYDGDHDNDKESDVEI300000000000000000	sp	015379	HDAC3_HUMAN
13204	MVSQSQPGSR2000000	sp	075122	CLAP2_HUMAN
13205	VLSTSSTLTR200000000	sp	P16144	ITB4_HUMAN
13206	NFSPNAPAHFSFVPAR2000000000000000	sp	Q6P0Q8	MAST2_HUMAN
13207	LLQTTNNSPMNSKPQIK200000000000	sp	000178	GTPB1_HUMAN
13208	KDSSSEVFSDAAK200000	sp	Q5T5U3	RHG21_HUMAN
13209	SQGITHHTR200	sp	014512	SOC S7_HUMAN
13210	TPTPSTPGDTQPNT PAPVPPAEDGIK2.02E+26	sp	Q14839	CHD4_HUMAN
13211	LQDSSDPDTGSEEEGSSRLSPPHSPR2.000202E+22	sp	Q92974	ARHG2_HUMAN
13212	LSTIQNSPTK200000000	sp	Q5VYV7	SLX4I_HUMAN
13213	IFDEVMGCFSDSPPQSPTFPEAGHTSLYDEDKVP R1.000202E+30	sp	Q13085	ACACA_HUMAN
13214	GHTASESDEQQWPEEK200000000000	sp	Q9NTI5	PDS5B_HUMAN
13215	HGSYEDAVHSGALND20000002000000	sp	P17987	TCPA_HUMAN
13216	TFSDGLATR20000000	sp	Q86UU1	PHLB1_HUMAN
13217	HSSVSDSQCEPPSVGTEYSQGASQPQHQ LK2E+30	sp	P39880	CUX1_HUMAN
13218	SHISDQSPLSSK2000000000	sp	P42166	LAP2A_HUMAN
13219	TRDDGDEEGLLTHSEEELEHSQD TDADDGALQ2E+21	sp	Q9UKM9	RALY_HUMAN
13220	GTPPLTPSDSPQTR200000	sp	Q6NZY4	ZCHC8_HUMAN
13221	QVPVMQGTGMQGASIQGGSQPGGFSPGQNVTPQDHEK1.00001E+34	sp	P33240	CSTF2_HUMAN
13222	SRSHSPMSNR20002000000	sp	Q13595	TRA2A_HUMAN
13223	SLRSPLPSR20200000000	sp	Q9NYV4	CDK12_HUMAN
13224	LLGDSDEEEQKEAASSGDDSGRDR2.020000000022E+21	sp	Q9BW71	HIRP3_HUMAN
13225	SVSASSGPCSPAAPHPPPPR2.022E+19	sp	Q86UY5	FA83A_HUMAN
13226	SHMSGSPGPGSNTAPSTPVI GSDKPGMEEK1.202E+30	sp	A0FGR8	ESYT2_HUMAN
13227	QGSITSPQANEQSVTPQR202000000000000000	sp	Q9UQ35	SRRM2_HUMAN
13228	SKSPASVDR2020000000	sp	Q13243	SRSF5_HUMAN
13229	RSSDTSGNEASEIESVK202200000000000000	sp	Q71F23	CENPU_HUMAN
13230	VAAAAGSGSPPGSPGHDR2000000000000000	sp	Q9BTU6	P4K2A_HUMAN
13231	STSGPRPGCQPSSPCVPK20000000002000000	sp	014545	TRAD1_HUMAN
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13233	ESSPIPSPTSDR220000000000	sp	Q01082	SPTB2_HUMAN
13234	SSPHSTYYMGQSSSGSIHSNLEQEK3E+19	sp	Q9ULH0	KDIS_HUMAN
13235	QGSITSPQANEQSVTPQRR200000	sp	Q9UQ35	SRRM2_HUMAN
13236	NEENTEPGAESSENADDPNKDTS ENADGQSDENKDDY TIPDEYR2E+40	sp	P43243	MATR3_HUMAN
13237	AGMSSNQSISSPVLDVPR102000000000000000	sp	Q9UQ35	SRRM2_HUMAN
13238	VHGEPTSDLS DID200020000	sp	095696	BRD1_HUMAN
13239	RPTETNPVTSNSDEECNETVK200022020000000000	sp	P46100	ATRX_HUMAN
13240	NTPSQHSHSIQHSPER20000200020000	sp	Q9NYF8	BCLF1_HUMAN
13241	SRSFTLDESLK200000000	sp	Q86WR7	PRSR2_HUMAN
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13243	TVSASSTGDLPK20000000000	sp	P78362	SRPK2_HUMAN
13244	GPTSSPCEEEGDEGEEDRTSDLRDEASSVTR2.00000000000002E+27	sp	Q86VM9	ZCH18_HUMAN
13245	QESCSPHHPQVLAQQGSGSSPK2E+20	sp	Q8N1G0	ZN687_HUMAN

13246	RQSPSPSTRPIR20002000000	sp Q8IYB3 SRRM1_HUMAN
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13249	SSSPFLSKR220000000	sp Q9NYV4 CDK12_HUMAN
13250	HSHSPASSPKPLKK200020000000	sp Q9HDC5 JPH1_HUMAN
13251	RHSSNTDNIPPER200200000000	sp Q96EX2 RNFT2_HUMAN
13252	FSSVSSPQPR2000000	sp Q9P206 K1522_HUMAN
13253	GSGDISSDSIDHSPAK20000	sp Q9H5I1 SUV92_HUMAN
13254	IPMTPTSSFVSPPPPTASPHSNR20000200000000	sp Q7Z5L9 I2BP2_HUMAN
13255	NAVASVQNGPGGGPSSSIGASKSDESSTEETDK200200000000	sp Q8NI27 THOC2_HUMAN
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13257	KSDGACDPSDDKENSQIAQDHQK2000000000000000000	sp Q9UIG0 BAZ1B_HUMAN
13258	ASIGQSPGLPSTTFK2000000000000000	sp Q5VT52 RPRD2_HUMAN
13259	STPLASPSPPGR2000002020000	sp Q8WYP5 ELYS_HUMAN
13260	SPQPVPSPRPQSQPPHSSPSPR2.00000200002E+22	sp Q09472 EP300_HUMAN
13261	LDPFADGGKTPDPK200000	sp 075533 SF3B1_HUMAN
13262	TVLPTVPESPEEVK20000000	sp 075152 ZC11A_HUMAN
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		sp P49792 RBP2_HUMAN
		sp Q7Z3J3 RGPD4_HUMAN
13264	EDALDDSVSSSVHASPLASSPVRK2.02222000000002E+19	sp A6NKT7 RGPD3_HUMAN
		sp O14715 RGPD8_HUMAN
		sp Q99666 RGPD5_HUMAN
13265	LLEDGEDFNLGDALDSSNSMQTIQK20000000	sp P05783 K1C18_HUMAN
13266	SLTNSHLEKK202000000	sp Q9H2H9 S38A1_HUMAN
13267	ALRPPTSPGVFGALQNFK2000000000000	sp 075151 PHF2_HUMAN
13268	ERPISLGFPLPAGDGLLTPDAQK2.00000000000002E+20	sp 060271 JIP4_HUMAN
13269	LHSYSSPSTK2000000	sp Q96T37 RBM15_HUMAN
13270	KNSSTDQGSDEEGLQK20000000020000	sp Q12789 TF3C1_HUMAN
13271	SLPSSPLR200200000	sp O14830 PPE2_HUMAN
13272	NMTVEQLLTGSPTSPTVEPEKPTR2E+22	sp Q9C0C9 UBE20_HUMAN
13273	STPLASPSPPGRSPQR20000020000	sp Q8WYP5 ELYS_HUMAN
13274	LFSLSNPSLSTTNLSGPSR2000000000000000	sp Q6UUV7 CRT3_HUMAN
13275	TSSLDITGPYLTGQWPR300000000	sp Q86VQ1 GLC11_HUMAN
13276	SRASPSR200000	sp Q9UQ35 SRRM2_HUMAN
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13278	SLSSSIGSNSTYLTSK2200000000000000	sp Q86SQ4 GP126_HUMAN
13279	SHSGTSPDNTAPPPPPRPQPSHSR2.20002E+21	sp Q96D71 REPS1_HUMAN
13280	SRGSSPGLR20002200000	sp Q9UQ35 SRRM2_HUMAN
13281	GEIKDSSPSSPSPK22000002000	sp Q6WKZ4 RFIP1_HUMAN
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13284	GTGGSESRASGLVGR2002022000000000	sp Q9UH99 SUN2_HUMAN
13285	ESAAPASPAPSPAPTAPPQK2.0000200000002E+22	sp Q3KQU3 MA7D1_HUMAN
13286	LCSPLEPGDPSK200000000000	sp Q5VUA4 ZN318_HUMAN
		sp Q9UQ88 CD11A_HUMAN
13287	RGTSPRPPEGLGYSQLGDDDLK2.000000000002E+21	sp P21127 CD11B_HUMAN
		sp O95685 PPR3D_HUMAN
13288	SLPSSPER200020000	sp P02545 LMNA_HUMAN
13289	ASSHSSQTQGGGSVTK2000000000000	sp O60245 PCDH7_HUMAN
13290	SSSPLPTVQLHPQSPTAGK2.000000000002E+19	sp Q9NTI5 PDS5B_HUMAN
13291	LDSEMDSSENYDTMSSPLPGK32000000000	sp Q8WWQ0 PHIP_HUMAN
13292	LSRGSISSTSEVHSPNVGLR2220000000000000	sp Q8IYB3 SRRM1_HUMAN
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		sp P30505 1C08_HUMAN
		sp Q9TNN7 1C05_HUMAN
		sp Q29960 1C16_HUMAN
		sp P30508 1C12_HUMAN
		sp Q95604 1C17_HUMAN
13295	GGSCSQAASSNSAQGSDESLIACK2000000000000000	sp Q07000 1C15_HUMAN
		sp Q29963 1C06_HUMAN
		sp P30510 1C14_HUMAN
		sp P04222 1C03_HUMAN
		sp Q29865 1C18_HUMAN
		sp P30504 1C04_HUMAN
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13298	VYSLPDGTFSSDEDEEEEEEEEEEEEEET2E+20	sp O15371 EIF3D_HUMAN
13299	RTSLPCIPR20000000	sp Q14432 PDE3A_HUMAN
13300	SVSSFPVPQDNVDTHPGSGK2E+20	sp Q676U5 A16L1_HUMAN
13301	EKEPGEQASVPLSPK2000	sp P42858 HD_HUMAN
13302	RGGADTYSVPSVLSR232000000000	sp P14543 NID1_HUMAN
13303	ETSVSKEDTDHEEKASNEVDTK2000200000020000000	sp O75475 PSIP1_HUMAN
13304	DLEPGEVPSDSDEDEGEHK20200000000	sp Q96T58 MINT_HUMAN
13305	FVCVGGSPSR20000	sp Q16831 UPP1_HUMAN
13306	AGEAPTENPAPPTQSSAE20000000	sp P16989 YBOX3_HUMAN
13307	DSFIENSSNCTSGSSKPNSPSISPSILSNTEHK2.022E+22	sp Q99700 ATX2_HUMAN
13308	SVCPGGSSPGSSSGGR20000200000	sp Q53LP3 SWAHC_HUMAN
13309	RNSNSPPSPSMNR20000200010000	sp Q7Z5L9 I2BP2_HUMAN
13310	LEPTNVQTVTCSR2000	sp Q99741 CDC6_HUMAN
13311	LPNLSSPAEGPPGPPSGPAPR20000000000000000000	sp O60784 TOM1_HUMAN
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13313	LSPIPEEVPK2000000000	sp Q96T23 RSF1_HUMAN
13314	TVSLTPSPTTQVETPDLVDHNTSPLFR200000	sp Q05209 PTN12_HUMAN
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13316	ELPAAAAIGATSLVAAPHSSSSSPSK2200000000	sp Q14C86 GAPD1_HUMAN
13317	TGVTSTSDSEEEGDDQEGEKK220000000000000000	sp O75475 PSIP1_HUMAN
13318	APPPSRPPRPSV200	sp Q9H6X2 ANTR1_HUMAN
13319	SKHEEEWTDDDLVESL2000000000	sp P51946 CCNH_HUMAN
13320	LKSSTSFANIQENSN220000000200	sp Q86WC4 OSTM1_HUMAN
13321	NSSGSESSLPNLAR20000000000000	sp Q2M2I8 AAK1_HUMAN
13322	KPVSEKTESDTIPYLK202000000	sp Q96PC5 MIA2_HUMAN
13323	LKSSTSFANIQENSN20020000000000	sp Q86WC4 OSTM1_HUMAN
13324	RPLSGPDVGTQPAGLASGAK20000000000000000000	sp O96013 PAK4_HUMAN
13325	RHSSVSDSQPEPPSVGTEYSQGASPPQHQLK2.02E+28	sp P39880 CUX1_HUMAN
13326	ADTEGKSPSHSSSNRPFTPPTSTGGSK2.02E+19	sp Q15648 MED1_HUMAN
13327	EKSATRPSPPER2002020000	sp Q9UQ35 SRRM2_HUMAN
13328	ALPTSKPEGLHSSPVGPSSSK2000000000	sp Q9UPT8 ZC3H4_HUMAN
13329	HSLSGSSPGMK20000200100	sp Q9UQ35 SRRM2_HUMAN
13330	DRPGSPESPLLDAPFSR20000000000000	sp Q8TF72 SHRM3_HUMAN
13331	CSPTVAFVEFPSSPQLK200000000200000	sp Q9UPQ0 LIMC1_HUMAN
13332	SRLTPVSPESSTEEK200200000000000000	sp Q13501 SQSTM_HUMAN
13333	SRSPRPK2000000000	sp Q8WUQ7 CATIN_HUMAN
13334	RSSLSSHSHQSQIYR20220200000000	sp O15027 SC16A_HUMAN
13335	AESSGGGTVPSAGILEQGSPGSGSPPKPK2E+29	sp Q68EM7 RHG17_HUMAN
13336	VRPSSDLNNTGQSPHHK200000	sp P27448 MARK3_HUMAN
13337	DRSPHRSSPSDTRPK2000020000	sp Q8IWS0 PHF6_HUMAN
13338	SPDSCRQALPCLPSTQDVPSR2E+22	sp Q9Y283 INVS_HUMAN

13339	RSPASPHPPFFTHR20020000000000	sp Q17RY0 CPEB4_HUMAN
13340	TPLSSLSATSNNSTVACIDR2020000000000000	sp P00533 EGFR_HUMAN
13341	YSEFTSTSTSGTGHNQTR2020000000	sp P48960 CD97_HUMAN
13342	VAEPGAEATSSTGEESGSEHPPAVPMHNC20000020000000000000	sp Q14684 RRP1B_HUMAN
13343	SGSDAGEARPPTPASPR2000000002000000	sp Q96FS4 SIPA1_HUMAN
13344	ITSPLMEPSSIEK200100000000	sp P28066 PSA5_HUMAN
13345	ATNESEDEIPQLVPIGKK200000000000000000	sp 076021 RL1D1_HUMAN
13346	KLPSDSGDLEALEGK20000000000	sp 060524 NEMF_HUMAN
13347	QPSSPYEDKDKDK20000000000	sp Q9UQ35 SRRM2_HUMAN
13348	IKPVTENLVLSSTGFCQSSGDEALSK202000000000000000	sp P46100 ATRX_HUMAN
13349	APSASDSKADSDGAKPEPVAMAR2.0000020002E+23	sp Q7Z4V5 HDGR2_HUMAN
13350	NACILQASGSPPAKSPTDPK200002000000	sp P41182 BCL6_HUMAN
13351	RRSPSPYYSR200003000	sp P62995 TRA2B_HUMAN
13352	NLSPTPASPNOGPPPQVPVSPGPPK2000000	sp Q13595 TRA2A_HUMAN
13353	SYLLGNSSPR20000	sp Q9COE8 LNP_HUMAN
13354	ATAPQTQHVSPMR2000000000100	sp P02545 LMNA_HUMAN
13355	KWEDSETPK202000	sp P29692 EF1D_HUMAN
13356	MEQQYESSSDGTEKLPK200020000000	sp P31629 ZEP2_HUMAN
13357	SRTVSASSTGDLPK202020000000	sp P46100 ATRX_HUMAN
13358	RLSSTSLASGHSVR2222000002000	sp P78362 SRPK2_HUMAN
13359	YFQSPSR30000000	sp Q9BZL6 KPCD2_HUMAN
13360	AGASIPSSGAASPR2000	sp Q16629 SRSF7_HUMAN
13361	SEDESETEDEEEKSQEDQEQK2E+21	sp Q9P270 SLAI2_HUMAN
13362	VPAQCTVPSHPQSTP VHSPDR200020000	sp P25205 MCM3_HUMAN
13363	LSPLLPGGGGR20000000000	sp Q9NXL9 MCM9_HUMAN
13364	GLGHPPSSPLLK200000	sp 015320 CTGE5_HUMAN
13365	IPRPSVSGCSR200	sp 095785 WIZ_HUMAN
13366	LGDDNLCNTPHLPPCSPPK20000	sp 075122 CLAP2_HUMAN
13367	QQLEEEENFSRPQIDENPLDDNSEEMEDAPK20000000000	sp Q99741 CDC6_HUMAN
13368	DSSKGEDSAEETEAKPAVVAPVVEAVSTPSAAFPDATAEQGPILTK2.2E+48	sp Q5F1R6 DJC21_HUMAN
13369	KSASSRELMIVGR2022000100000	sp Q12906 ILF3_HUMAN
13370	EDALDDSVSSSVHASPLASSPVR202000000000020000	sp P53794 SC5A3_HUMAN
13371	SQSNPILGSPFFSHFDGQDSYAAAVR2E+24	sp P49792 RBP2_HUMAN
13372	IISPGSSTPSSTR200000000200	sp Q7Z3J3 RGPD4_HUMAN
13373	DYDDMSPR12000	sp A6NKT7 RGPD3_HUMAN
13374	GTSPAGPGTFKPGSPTVGPGLPPAGPSGLPSLPPPAAPASGPPLSATQIK2.0000002E+51	sp 014715 RGPD8_HUMAN
13375	SRSYSR20202000	sp Q99666 RGPD5_HUMAN
13376	WSPAYSFSSDSPLDSSPK200020000	sp Q9NYL2 MLTK_HUMAN
13377	SKKEDSSSENAEPDLDNEDEEPAVEIEPEPEPQVTPAPPPAK2.202E+41	sp Q5VT52 RPRD2_HUMAN
13378	ILDATDQESLELKPTSR20002000000000	sp P61978 HNRPK_HUMAN
13379	EPSVPASCAVSATYERSPSPAHLPPDDPK2300000000000000	sp P54259 ATN1_HUMAN
13380	HSSPHISR2000000	sp Q8IYB3 SRRM1_HUMAN
13381	EGTPPPPPSR2000000000	sp Q13595 TRA2A_HUMAN
		sp P62995 TRA2B_HUMAN
		sp Q07955 SRSF1_HUMAN
		sp Q8NCG7 DGLB_HUMAN
		sp P49711 CTCF_HUMAN
		sp Q8IZA0 K319L_HUMAN
		sp Q92615 LAR4B_HUMAN
		sp 060583 CCNT2_HUMAN
		sp Q9Y618 NCOR2_HUMAN



13382	RLSDCESTDVKR20000000000	sp Q7Z333 SETX_HUMAN
13383	SKFDSDEEEEDTENVEAASSGK200000000000020000	sp Q8TF01 PNISR_HUMAN
13384	TPEKLDNTPASPPRSPAEPNDIPIAK20002000000000000	sp 095359 TACC2_HUMAN
13385	LAWPINSNPPNTFV200000000	sp Q15760 GPR19_HUMAN
13386	EASEVKPPR20000000	sp 060341 KDM1A_HUMAN
13387	SRSPQRPGWSR200000000200	sp Q9UQ35 SRRM2_HUMAN
13388	VGSLTPSSPK2020002000	sp Q2M2I8 AAK1_HUMAN
13389	SHSLSPSPVVK2020000000	sp Q96GE4 CEP95_HUMAN
13390	SHSGTSPDNTAPPPPPRPQPSHSR2. 02E+25	sp Q96D71 REPS1_HUMAN
13391	VAEPGAEATSSTGEESGSEHPPAVPMHNK2. 2020002E+21	sp Q14684 RRP1B_HUMAN
13392	RSTSLPSR22000000	sp P18054 LOX12_HUMAN
13393	TPSSEEISPTK200000000	sp Q15311 RBP1_HUMAN
13394	NTPSQHSHSIQHSPER20020200020000	sp Q9NYF8 BCLF1_HUMAN
13395	MTPNSTGEVPSPIPK12000200000000000	sp Q8TDU6 GPBAR_HUMAN
13396	EDFDSTSSSSSTPPLQPR20000000	sp Q13620 CUL4B_HUMAN
13397	SFSPLQPGQAPTGR2000000000000	sp Q7Z3C6 ATG9A_HUMAN
13398	TLETQLAPDCCPSDQDPAPAHPSPHASPMNK2000201000	sp Q9BUR4 WAP53_HUMAN
13399	ILATDVSSKNTPDSKK2002000	sp Q03188 CENPC_HUMAN
13400	VEMYSGSDDDDFNKLPK2000000000000000	sp 060841 IF2P_HUMAN
13401	HAASYSDSENQGSYSGVIPPPGR200000000000000000	sp Q96D71 REPS1_HUMAN
13402	ADGYEPPVQESV200	sp P61247 RS3A_HUMAN
13403	LVKPSSSSSGSRSSR200000002200	sp Q9H6B4 CLMP_HUMAN
13404	MPGMSPANPSLHSPVPDASHSPR2E+19	sp 060244 MED14_HUMAN
13405	LQLSDEESVFEEALMSPDTR20000001200000	sp Q86SQ0 PHLB2_HUMAN
13406	MHNLHGTKPPPSEGSDEEEEEDEEDEEER2E+24	sp Q9BQ67 GRWD1_HUMAN
13407	QPGQVIGATTPSTGSPTNK20000000	sp Q9ULM3 YETS2_HUMAN
13408	SPSSESSQHTPPARPR200000000000000000	sp 014733 MP2K7_HUMAN
13409	DSSDSADGRATPSENLPSSAR2E+20	sp Q8N684 CPSF7_HUMAN
13410	HSTASHSSHTSGIEADTKPR202220000000000000	sp Q96NE9 FRMD6_HUMAN
13411	YSPSQNSPIHHIPSR32000020000000000	sp Q9NYF8 BCLF1_HUMAN
13412	KYSTSWTAPR3002000000	sp Q8NEM8 CBPC3_HUMAN
13413	ERSPSPSFR20000000	sp P49585 PCY1A_HUMAN
13414	TFVLTPELSPGK20000	sp Q96DY7 MTBP_HUMAN
13415	DQHLNNSSSPQR20000	sp Q15154 PCM1_HUMAN
13416	LDIPTGTPQR200000	sp P52732 KIF11_HUMAN
13417	KVTDQISDIVYK20000000300	sp P20929 NEBU_HUMAN
13418	SFSSSPSPSPTSPHRPSIR2. 02220202E+20	sp Q86VM9 ZCH18_HUMAN
13419	SPESDRPR200200000	sp Q96T37 RBM15_HUMAN
13420	SKSHESQLGMR2000000	sp Q8IVT5 KSR1_HUMAN
13421	RGGHSSVSTESSESSFHSS2000000000000000	sp P61073 CXCR4_HUMAN
13422	GPTSSPCEEEGDEGEEDRTSDLRDEASSVTR2. 2E+29	sp Q86VM9 ZCH18_HUMAN
13423	SPDTLAPVASER20000000000000	sp 043310 CTIF_HUMAN
13424	LKWTNTGTPHQTPYQLDPANITLPEMEIR30000000200010000	sp 000754 MA2B1_HUMAN
13425	GSSPSIRPIQGSQGSSSPVEK2E+20	sp Q92560 BAP1_HUMAN
13426	NQSPTEAEKPASSSLPSSPPPQLLTR22000000000	sp Q99567 NUP88_HUMAN
13427	DTGSEVPSGSGHGCPPTPPAPANFEDVAPTGSGEFGATRESPK20000020000000	sp 015047 SET1A_HUMAN
13428	SPAGSQQFGYSPGQQQTHPQGSRR2. 00000000002E+25	sp Q8TAP9 MPLKI_HUMAN
13429	HSSSPLSPGGPHTLTKPTTSSSER2. 2E+23	sp Q9Y618 NCOR2_HUMAN
13430	TFSDENVL20000000	sp Q8NB49 AT11C_HUMAN
13431	SVCPGGSSPGSSSGGR200002000000	sp Q53LP3 SWAHC_HUMAN
13432	DQPDGSSLSPAQSPSQSPPAASSLR202000000000000	sp Q6WCQ1 MPRIP_HUMAN
13433	VAPSPDSSDTDSSEDPRLR202000000000000	sp Q9NP71 MLXPL_HUMAN
13434	SRSRSPHEAGFCVYLK202020000000000000	sp Q9NTZ6 RBM12_HUMAN

13435	LNLGTDSOSSPQK20000	sp Q16512 PKN1_HUMAN
13436	SQDADSPGSSGAPENLTFK200000000000	sp P55196 AFAD_HUMAN
13437	SPTPPSSAGLGSNSAPPIPDSR220000202000000000	sp Q8IWX8 CHERP_HUMAN
13438	LTTEATQTPVSNK2002000	sp Q9BXS6 NUSAP_HUMAN
13439	LPLVPESPRR20000	sp Q86WB0 NIPA_HUMAN
13440	YSPSQNSPIHHIPSR3200000000000000	sp Q9NYF8 BCLF1_HUMAN
13441	VVPGQFDDADSSDSENRLK220200000000	sp Q9BRS2 RIOK1_HUMAN
13442	YSPTSPTYSPPTPK320000030002000	sp P24928 RPB1_HUMAN
13443	VAGGAAPSKPASAKK20000	sp Q14978 NOLC1_HUMAN
13444	AAPSTAPAEATPPKPGAEAEAPPK2E+19	sp O95817 BAG3_HUMAN
13445	ITEVSCKSPQDPVK2000000000000000	sp P46013 KI67_HUMAN
13446	EVHDELEDLSPPPPPLSPPTTSPHK20000200000	sp Q9NQX3 GEPH_HUMAN
13447	YSRSPYSR20200000	sp Q9BRL6 SRSF8_HUMAN
13448	KAEGEPQEE SPLKSK2000200	sp Q9NYF8 BCLF1_HUMAN
13449	EELMSSDLEETAGSTSIPK20000000000	sp O00567 NOP56_HUMAN
13450	RHSSDINHLVTQGRESPEGSYTDDANQEV2.0000000000002E+28	sp Q5TON5 FBP1L_HUMAN