

Table S1 The 848 peptides identified by CESI-Q-TOF-MS.

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
1	AAEIYGTKESPQTHY	1693.795	Stomach (0.5 h)	16	AEIYGTKESPQTHYYAV	1955.927	Stomach (0.5 h)
2	AAHVKQV	751.4341	Stomach (0.5 h)	17	AGRDPYKLRP	1171.646	Stomach (0.5 h)
3	AAHVKQVL	864.5181	Stomach (0.5 h)	18	AGRDPYKLRPV	1270.715	Stomach (0.5 h)
4	AAHVKQVLL	977.6022	Stomach (0.5 h)	19	AIAEKKADAVTLDGGMV	1687.882	Stomach (0.5 h)
5	AAHVKQVLLHQ	1242.72	Stomach (0.5 h)	20	AIAEKKADAVTLDGGMV	1850.945	Stomach (0.5 h)
6	AAHVKQVLLHQQ	1370.778	Stomach (0.5 h)	21	AIANLKK	756.4857	Stomach (0.5 h)
7	AAHVKQVLLHQQAL	1554.899	Stomach (0.5 h)	22	AKLGGRPT	798.4712	Stomach (0.5 h)
8	ADRDQYELL	1121.535	Stomach (0.5 h)	23	AKLGGRPTY	961.5345	Stomach (0.5 h)
9	AEDVGDV	703.3025	Stomach (0.5 h)	24	ALNLDGGYIY	1097.539	Stomach (0.5 h)
10	AEIYGTK	780.4017	Stomach (0.5 h)	25	ALYLGSRV	941.497	Stomach (0.5 h)
11	AEIYGTKESPQ	1221.588	Stomach (0.5 h)	26	ANEGLTWNSL	1103.525	Stomach (0.5 h)
12	AEIYGTKESPQT	1322.635	Stomach (0.5 h)	27	ANEGLTWNSLK	1231.62	Stomach (0.5 h)
13	AEIYGTKESPQTH	1459.694	Stomach (0.5 h)	28	ANEGLTWNSLKD	1346.647	Stomach (0.5 h)
14	AEIYGTKESPQTHY	1622.758	Stomach (0.5 h)	29	ANEGLTWNSLKDK	1474.742	Stomach (0.5 h)
15	AEIYGTKESPQTHYY	1785.821	Stomach (0.5 h)	30	AQEKFGK	806.4286	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
31	AQEKFGKNK	1048.567	Stomach (0.5 h)	46	DGAGDVA	603.25	Stomach (0.5 h)
32	AVAVVKKGSNF	1118.645	Stomach (0.5 h)	47	DGAGDVAFVKETTVF	1554.757	Stomach (0.5 h)
33	AVAVVKKGSNFQ	1246.703	Stomach (0.5 h)	48	DGGMVFEAGRDP	1249.54	Stomach (0.5 h)
34	AVAVVKKGSNFQL	1359.787	Stomach (0.5 h)	49	DGGMVFEAGRDPY	1412.603	Stomach (0.5 h)
35	AVAVVKKGSNFQLDQLQG	2057.138	Stomach (0.5 h)	50	DGKEDLI	788.3916	Stomach (0.5 h)
36	AVDRTAGWNIP	1198.61	Stomach (0.5 h)	51	DGKEDLIWK	1102.566	Stomach (0.5 h)
37	AVTLDGGM	778.3531	Stomach (0.5 h)	52	DLLFKDSAL	1020.549	Stomach (0.5 h)
38	AVTLDGGMV	877.4215	Stomach (0.5 h)	53	DLLFKDSALG	1077.571	Stomach (0.5 h)
39	AVTLDGGMVF	1008.495	Stomach (0.5 h)	54	DLLFKDSALGF	1224.639	Stomach (0.5 h)
40	AVVARSVD	815.4501	Stomach (0.5 h)	55	DLLFKDSALGFL	1337.723	Stomach (0.5 h)
41	AVVARSVDGKED	1244.636	Stomach (0.5 h)	56	DRDQYELL	1050.498	Stomach (0.5 h)
42	AVVARSVDGKEDL	1357.72	Stomach (0.5 h)	57	DSALGFL	721.3647	Stomach (0.5 h)
43	AVVARSVDGKEDLIWKL	1898.063	Stomach (0.5 h)	58	EAGRDPY	806.3559	Stomach (0.5 h)
44	AVVKKGSNFQLDQLQG	1730.932	Stomach (0.5 h)	59	EAGRDPYK	934.4508	Stomach (0.5 h)
45	DAVTLDGGMVF	1139.517	Stomach (0.5 h)	60	EAGRDPYKLRP	1300.689	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
61	EAGRDPYKLRPV	1399.757	Stomach (0.5 h)	76	ENTNGESTADWAK	1421.606	Stomach (0.5 h)
62	EAGRDPYKLRPVA	1470.794	Stomach (0.5 h)	77	EPLQGAVAK	911.5076	Stomach (0.5 h)
63	EAGRDPYKLRPVAA	1541.831	Stomach (0.5 h)	78	EPLQGAVAKF	1058.576	Stomach (0.5 h)
64	EAGRDPYKLRPVAAEIY	1947.021	Stomach (0.5 h)	79	ESPQTHY	860.3664	Stomach (0.5 h)
65	EEYLGTEY	1002.418	Stomach (0.5 h)	80	ETAEEVK	804.3865	Stomach (0.5 h)
66	EIYGTKESPQTHY	1551.721	Stomach (0.5 h)	81	ETAEEVKA	875.4236	Stomach (0.5 h)
67	ENLPEKAD	914.4345	Stomach (0.5 h)	82	ETTVFENLPEK	1305.645	Stomach (0.5 h)
68	ENLPEKADR	1070.536	Stomach (0.5 h)	83	ETTVFENLPEKADRQ	1890.896	Stomach (0.5 h)
69	ENLPEKADRQ	1185.563	Stomach (0.5 h)	84	FENLPEKADRQ	1460.69	Stomach (0.5 h)
70	ENLPEKADRQ	1313.621	Stomach (0.5 h)	85	FENLPEKADRQYEL	1865.88	Stomach (0.5 h)
71	ENLPEKADRQY	1476.684	Stomach (0.5 h)	86	FGSPPGQR	844.4192	Stomach (0.5 h)
72	ENLPEKADRQYE	1605.727	Stomach (0.5 h)	87	FGSPPGQRD	959.4461	Stomach (0.5 h)
73	ENLPEKADRQYEL	1718.811	Stomach (0.5 h)	88	FGSPPGQRDL	1072.53	Stomach (0.5 h)
74	ENLPEKADRQYELL	1831.895	Stomach (0.5 h)	89	FGSPPGQRDLL	1185.614	Stomach (0.5 h)
75	ENTNGESTADWA	1293.511	Stomach (0.5 h)	90	FGSPPGQRDLLFK	1460.778	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
91	FGSPPGQRDLLFKD	1575.805	Stomach (0.5 h)	106	GGRPTYEEYLGTEYV	1732.794	Stomach (0.5 h)
92	FKDSALGF	883.444	Stomach (0.5 h)	107	GGRPTYEEYLGTEYVT	1833.842	Stomach (0.5 h)
93	FKSETKN	852.4341	Stomach (0.5 h)	108	GGRPTYEEYLGTEYVTA	1904.879	Stomach (0.5 h)
94	FKSETKNL	965.5182	Stomach (0.5 h)	109	GILRPYL	830.5014	Stomach (0.5 h)
95	FKSETKNLL	1078.602	Stomach (0.5 h)	110	GILRPYLSWTESLEP	1759.915	Stomach (0.5 h)
96	FKSETKNLLFND	1454.741	Stomach (0.5 h)	111	GILRPYLSWTESLEPL	1872.999	Stomach (0.5 h)
97	GAGDVAFVKETTV	1292.661	Stomach (0.5 h)	112	GILRPYLSWTESLEPLQGA	2129.116	Stomach (0.5 h)
98	GAGDVAFVKETTVF	1439.73	Stomach (0.5 h)	113	GILRPYLSWTESLEPLQGAVA	2299.221	Stomach (0.5 h)
99	GESTADW	764.2977	Stomach (0.5 h)	114	GILRPYLSWTESLEPLQGAVAK	2427.316	Stomach (0.5 h)
100	GESTADWAK	963.4298	Stomach (0.5 h)	115	GILRPYLSWTESLEPLQGAVAK	2574.385	Stomach (0.5 h)
101	GESTADWAKNL	1190.557	Stomach (0.5 h)	116	GMVFEAGRDPY	1240.555	Stomach (0.5 h)
102	GGRPTYEE	907.4035	Stomach (0.5 h)	117	GRDPYKLRP	1100.609	Stomach (0.5 h)
103	GGRPTYEEY	1070.467	Stomach (0.5 h)	118	GRDPYKLRPV	1199.678	Stomach (0.5 h)
104	GGRPTYEEYLGTE	1470.663	Stomach (0.5 h)	119	GRDPYKLRPVA	1270.715	Stomach (0.5 h)
105	GGRPTYEEYLGTEY	1633.726	Stomach (0.5 h)	120	GRPTYEE	850.3821	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
121	GRPTYEEY	1013.445	Stomach (0.5 h)	136	IAEKKAD	773.4283	Stomach (0.5 h)
122	GRPTYEEYL	1126.53	Stomach (0.5 h)	137	IAEKKADA	844.4654	Stomach (0.5 h)
123	GRPTYEEYLGTE	1413.641	Stomach (0.5 h)	138	IAEKKADAVTLDGGMVF	1763.913	Stomach (0.5 h)
124	GRPTYEEYLGTEY	1576.705	Stomach (0.5 h)	139	IIPMGILRPY	1171.679	Stomach (0.5 h)
125	GRPTYEEYLGTEYV	1675.773	Stomach (0.5 h)	140	IIPMGILRPYL	1284.763	Stomach (0.5 h)
126	GRPTYEEYLGTEYVT	1776.821	Stomach (0.5 h)	141	IIPMGILRPYLS	1371.795	Stomach (0.5 h)
127	GSNFQLDQLQGR	1361.669	Stomach (0.5 h)	142	ILRPYLSW	1046.591	Stomach (0.5 h)
128	GSPPGQRD	812.3777	Stomach (0.5 h)	143	ILRPYLSWTE	1276.682	Stomach (0.5 h)
129	GSPPGQRDL	925.4617	Stomach (0.5 h)	144	ILRPYLSWTESLEPL	1815.977	Stomach (0.5 h)
130	GSPPGQRDLL	1038.546	Stomach (0.5 h)	145	IPSKVDS	744.4018	Stomach (0.5 h)
131	GTKESPQTH	983.4672	Stomach (0.5 h)	146	IYGTKESPQTHY	1422.678	Stomach (0.5 h)
132	GTKESPQTHY	1146.531	Stomach (0.5 h)	147	KADAVTLDGGMV	1175.586	Stomach (0.5 h)
133	GTKESPQTHYY	1309.594	Stomach (0.5 h)	148	KADAVTLDGGMVF	1322.654	Stomach (0.5 h)
134	GTKESPQTHYYA	1380.631	Stomach (0.5 h)	149	KADAVTLDGGMVFEAGR	2110.999	Stomach (0.5 h)
135	GWNIPMGL	886.4371	Stomach (0.5 h)	150	KESPQTHY	988.4614	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
151	KGEADAL	702.3548	Stomach (0.5 h)	166	LAVAVVKK	826.564	Stomach (0.5 h)
152	KGEADALNLDG	1101.5302	Stomach (0.5 h)	167	LAVAVVKKKA	897.6011	Stomach (0.5 h)
153	KGEADALNLDGGY	1321.615	Stomach (0.5 h)	168	LAVAVVKKKANEGLT	1411.84	Stomach (0.5 h)
154	KGSNFQL	792.413	Stomach (0.5 h)	169	LAVAVVKKKANEGLTW	1597.919	Stomach (0.5 h)
155	KGSNFQLDQLQGR	1489.7637	Stomach (0.5 h)	170	LAVAVVKKKANEGLTWN	1912.078	Stomach (0.5 h)
156	KLGRPTYEEYLGTEY	1874.905	Stomach (0.5 h)	171	LAVAVVKKKANEGLTWN	2040.173	Stomach (0.5 h)
157	KLRPVAAEIY	1158.6761	Stomach (0.5 h)	172	LEPLQGAVAK	1024.592	Stomach (0.5 h)
158	KLRPVAAEIYGTKESPQ	1886.0261	Stomach (0.5 h)	173	LFKDSALGF	996.528	Stomach (0.5 h)
159	KNLRETAEEVK	1315.7095	Stomach (0.5 h)	174	LGGRPTY	762.4024	Stomach (0.5 h)
160	KSETKNL	818.4498	Stomach (0.5 h)	175	LGGRPTYEE	1020.488	Stomach (0.5 h)
161	LAQVPSH	750.4024	Stomach (0.5 h)	176	LGGRPTYEEY	1183.551	Stomach (0.5 h)
162	LAVAPNH	720.3919	Stomach (0.5 h)	177	LGGRPTYEEYL	1296.635	Stomach (0.5 h)
163	LAVAPNHA	791.429	Stomach (0.5 h)	178	LGGRPTYEEYLGTE	1583.747	Stomach (0.5 h)
164	LAVAPNHAVVS	1076.5978	Stomach (0.5 h)	179	LGGRPTYEEYLGTEY	1746.81	Stomach (0.5 h)
165	LAVAVVK	698.4691	Stomach (0.5 h)	180	LGGRPTYEEYLGTEYV	1845.879	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
181	LGGRPTYEEYLGTEYVT	1946.926	Stomach (0.5 h)	196	LRIPSKVDSALY	1360.771	Stomach (0.5 h)
182	LGGRPTYEEYLGTEYVT	2017.963	Stomach (0.5 h)	197	LRPVAAEI	867.5178	Stomach (0.5 h)
183	LGGRPTYEEYLGTEYVT	2131.047	Stomach (0.5 h)	198	LRPVAAEIY	1030.581	Stomach (0.5 h)
184	LGGRPTYEEYLGTEYVT	2202.084	Stomach (0.5 h)	199	LRPYLSWTESLEPLQGAV	2404.279	Stomach (0.5 h)
185	LHQQALF	855.4603	Stomach (0.5 h)	200	LSKAQEK	802.4548	Stomach (0.5 h)
186	LLFKDSALG	962.5437	Stomach (0.5 h)	201	LSKAQEKF	949.5233	Stomach (0.5 h)
187	LLSKAQEKFGK	1247.724	Stomach (0.5 h)	202	LSKAQEKFG	1006.545	Stomach (0.5 h)
188	LLSKAQEKFGKNK	1489.862	Stomach (0.5 h)	203	LSKAQEKFGK	1134.64	Stomach (0.5 h)
189	LPEKADRDQ	1070.536	Stomach (0.5 h)	204	LSKAQEKFGKN	1248.683	Stomach (0.5 h)
190	LPEKADRDQY	1233.599	Stomach (0.5 h)	205	LSKAQEKFGKNK	1376.778	Stomach (0.5 h)
191	LRIPSKV	811.528	Stomach (0.5 h)	206	LSWTESLEPL	1173.592	Stomach (0.5 h)
192	LRIPSKVD	926.5549	Stomach (0.5 h)	207	LSWTESLEPLQ	1301.65	Stomach (0.5 h)
193	LRIPSKVDS	1013.587	Stomach (0.5 h)	208	LSWTESLEPLQGA	1429.709	Stomach (0.5 h)
194	LRIPSKVDSA	1084.624	Stomach (0.5 h)	209	LSWTESLEPLQGAVA	1599.814	Stomach (0.5 h)
195	LRIPSKVDSAL	1197.708	Stomach (0.5 h)	210	LSWTESLEPLQGAVAK	1727.909	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
211	LSWTESLEPLQGAVAKF	1874.978	Stomach (0.5 h)	226	NEGLTWNSL	1032.488	Stomach (0.5 h)
212	LSWTESLEPLQGAVAKF	2022.046	Stomach (0.5 h)	227	NEGLTWNSLK	1160.583	Stomach (0.5 h)
213	LTTLKNL	801.496	Stomach (0.5 h)	228	NEGLTWNSLKDK	1403.705	Stomach (0.5 h)
214	LTTLKNLR	957.5971	Stomach (0.5 h)	229	NEGLTWNSLKDKK	1531.799	Stomach (0.5 h)
215	LTTLKNLRE	1086.64	Stomach (0.5 h)	230	NLNREDF	906.4195	Stomach (0.5 h)
216	LTTLKNLRET	1187.687	Stomach (0.5 h)	231	NLPEKAD	785.3919	Stomach (0.5 h)
217	LTTLKNLRETA	1258.725	Stomach (0.5 h)	232	NLPEKADR	941.493	Stomach (0.5 h)
218	LTTLKNLRETAE	1387.767	Stomach (0.5 h)	233	NLPEKADR	1056.52	Stomach (0.5 h)
219	LTTLKNLRETAEE	1516.81	Stomach (0.5 h)	234	NLPEKADRQ	1184.579	Stomach (0.5 h)
220	LTTLKNLRETAEEVK	1743.973	Stomach (0.5 h)	235	NLPEKADRQY	1347.642	Stomach (0.5 h)
221	LTTLKNLRETAEEVKA	1815.01	Stomach (0.5 h)	236	NLPEKADRQYEL	1589.769	Stomach (0.5 h)
222	LYLGSRY	870.4599	Stomach (0.5 h)	237	NLRETAEEV	1059.52	Stomach (0.5 h)
223	MGILRPY	848.4578	Stomach (0.5 h)	238	NLRETAEEVKA	1258.652	Stomach (0.5 h)
224	MGILRPYL	961.5419	Stomach (0.5 h)	239	NSLKDKK	831.4814	Stomach (0.5 h)
225	NEGLTWNS	919.4035	Stomach (0.5 h)	240	NTNGESTADWAK	1292.563	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
241	PEKADRDQ	957.4516	Stomach (0.5 h)	256	RPTYEEYLGTEYVT	1719.7992	Stomach (0.5 h)
242	PEKADRDQY	1120.5149	Stomach (0.5 h)	257	RSVDGKEDL	1017.5091	Stomach (0.5 h)
243	QLDQLQG	800.4028	Stomach (0.5 h)	258	RSVDGKEDLI	1130.5931	Stomach (0.5 h)
244	QLFGSPPGQRDL	1313.6728	Stomach (0.5 h)	259	RSVDGKEDLIW	1316.6725	Stomach (0.5 h)
245	QLFGSPPGQRDLL	1426.7569	Stomach (0.5 h)	260	RSVDGKEDLIWK	1444.7674	Stomach (0.5 h)
246	RPYKLRP	1043.5876	Stomach (0.5 h)	261	RSVDGKEDLIWKLL	1670.9355	Stomach (0.5 h)
247	RPYKLRPVA	1213.6931	Stomach (0.5 h)	262	RSVDGKEDLIWKLLS	1757.9676	Stomach (0.5 h)
248	RETAEVKAR	1187.6258	Stomach (0.5 h)	263	RTAGWNIP	913.477	Stomach (0.5 h)
249	RIPSKVDS	900.5029	Stomach (0.5 h)	264	SAGWIIPM	889.4368	Stomach (0.5 h)
250	RIPSKVDSA	971.54	Stomach (0.5 h)	265	SAGWIIPMG	930.4633	Stomach (0.5 h)
251	RIPSKVDSAL	1084.624	Stomach (0.5 h)	266	SAGWIIPMGILRP	1425.7802	Stomach (0.5 h)
252	RPTYEEY	956.4239	Stomach (0.5 h)	267	SAGWIIPMGILRPY	1572.8486	Stomach (0.5 h)
253	RPTYEEYLGTE	1356.6197	Stomach (0.5 h)	268	SAGWIIPMGILRPYL	1685.9327	Stomach (0.5 h)
254	RPTYEEYLGTEY	1519.6831	Stomach (0.5 h)	269	SAGWIIPMGILRPYLSWT	2076.0866	Stomach (0.5 h)
255	RPTYEEYLGTEYV	1618.7515	Stomach (0.5 h)	270	SAGWIIPMGILRPYLSWTESLEPLQGAVAK	3298.7424	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
271	SALYLGSRY	1028.529	Stomach (0.5 h)	286	SVDGKEDLIWKL	1401.75	Stomach (0.5 h)
272	SDRAAHVKQVL	1222.678	Stomach (0.5 h)	287	SVDGKEDLIWKLL	1514.834	Stomach (0.5 h)
273	SKAQEKFGK	1021.556	Stomach (0.5 h)	288	SVDGKEDLIWKLLS	1601.867	Stomach (0.5 h)
274	SKAQEKFGKKNK	1263.694	Stomach (0.5 h)	289	SVDGKEDLIWKLLSK	1729.961	Stomach (0.5 h)
275	SLEPLQGA	813.4232	Stomach (0.5 h)	290	SWTESLEPL	1060.508	Stomach (0.5 h)
276	SLEPLQGAVA	983.5287	Stomach (0.5 h)	291	TAGWNIP	757.3759	Stomach (0.5 h)
277	SLEPLQGAVAK	1111.624	Stomach (0.5 h)	292	TAGWNIPM	904.4113	Stomach (0.5 h)
278	SLEPLQGAVAKF	1258.692	Stomach (0.5 h)	293	TAGWNIPMGL	1058.522	Stomach (0.5 h)
279	SLEPLQGAVAKFF	1405.761	Stomach (0.5 h)	294	TAVDRTAGWNIP	1299.657	Stomach (0.5 h)
280	SPPGQRD	755.3562	Stomach (0.5 h)	295	TESLEPL	787.3963	Stomach (0.5 h)
281	SPPGQRDL	868.4403	Stomach (0.5 h)	296	TESLEPLQG	972.4764	Stomach (0.5 h)
282	SPPGQRDLL	981.5243	Stomach (0.5 h)	297	TESLEPLQGA	1043.514	Stomach (0.5 h)
283	SVDGKEDL	861.408	Stomach (0.5 h)	298	TESLEPLQGAVA	1213.619	Stomach (0.5 h)
284	SVDGKEDLIW	1160.571	Stomach (0.5 h)	299	TESLEPLQGAVAK	1341.714	Stomach (0.5 h)
285	SVDGKEDLIWK	1288.666	Stomach (0.5 h)	300	TESLEPLQGAVAKF	1488.782	Stomach (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
301	TLDGGMV	707.316	Stomach (0.5 h)	316	VDRTAGWNIP	1127.572	Stomach (0.5 h)
302	TLDGGMVFEAGRDPY	1626.735	Stomach (0.5 h)	317	VKETTVF	822.4487	Stomach (0.5 h)
303	TTLKNLR	844.513	Stomach (0.5 h)	318	VKETTVFE	951.4913	Stomach (0.5 h)
304	TTLKNLRE	973.5556	Stomach (0.5 h)	319	VKETTVFENLPEKADRD	2118.059	Stomach (0.5 h)
305	TTVFENL	822.4123	Stomach (0.5 h)	320	VKQVLLHQQAL	1275.766	Stomach (0.5 h)
306	VAVVKKKA	713.48	Stomach (0.5 h)	321	VLKGEAD	730.3861	Stomach (0.5 h)
307	VAVVKKANEGLT	1227.719	Stomach (0.5 h)	322	VLKGEADALNL	1141.634	Stomach (0.5 h)
308	VAVVKKANEGLTW	1413.798	Stomach (0.5 h)	323	VLRPTEG	770.4286	Stomach (0.5 h)
309	VAVVKKANEGLTWN	1527.841	Stomach (0.5 h)	324	VLRPTEGY	933.492	Stomach (0.5 h)
310	VAVVKKANEGLTWNSL	1727.957	Stomach (0.5 h)	325	VLRPTEGYLA	1117.613	Stomach (0.5 h)
311	VAVVKKG	699.4643	Stomach (0.5 h)	326	VTAIANLK	828.5069	Stomach (0.5 h)
312	VAVVKKGSNF	1047.608	Stomach (0.5 h)	327	VTAIANLKK	956.6018	Stomach (0.5 h)
313	VAVVKKGSNFQ	1175.666	Stomach (0.5 h)	328	VVARSVDGKEDL	1286.683	Stomach (0.5 h)
314	VAVVKKGSNFQL	1288.75	Stomach (0.5 h)	329	VVKKANEGLTWNSLKD	1929.068	Stomach (0.5 h)
315	VDGKEDLIWK	1201.634	Stomach (0.5 h)	330	VVKKGSNF	877.5022	Stomach (0.5 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
331	VVKKGSNFQ	1005.561	Stomach (0.5 h)	346	YAVAVVKKGSNFQL	1522.851	Stomach (0.5 h)
332	VVKKGSNFQLDQLQG	1659.894	Stomach (0.5 h)	347	YAVAVVKKGSNFQLDQL	1879.02	Stomach (0.5 h)
333	VVSRSDRAAHVKQ	1451.796	Stomach (0.5 h)	348	YAVAVVKKGSNFQLDQLQG	2064.1	Stomach (0.5 h)
334	WAKNLNR	900.4929	Stomach (0.5 h)	349	YAVAVVKKGSNFQLDQLQG	2220.202	Stomach (0.5 h)
335	WAKNLNRE	1029.536	Stomach (0.5 h)	350	YEEYLG	873.3756	Stomach (0.5 h)
336	WAKNLNRED	1144.563	Stomach (0.5 h)	351	AAHVKQVL	730.3861	Stomach (1 h)
337	WIIPMGILRPY	1357.758	Stomach (0.5 h)	352	AEIYGTK	1141.634	Stomach (1 h)
338	WIIPMGILRPYL	1470.842	Stomach (0.5 h)	353	AEIYGTKESPQT	770.4286	Stomach (1 h)
339	WTESLEPLQGAVAK	1527.793	Stomach (0.5 h)	354	AEIYGTKESPQTH	933.492	Stomach (1 h)
340	YAVAVVK	748.4483	Stomach (0.5 h)	355	AEIYGTKESPQTHY	1117.613	Stomach (1 h)
341	YAVAVVKK	876.5433	Stomach (0.5 h)	356	AEIYGTKESPQTHYY	828.5069	Stomach (1 h)
342	YAVAVVKKG	933.5647	Stomach (0.5 h)	357	AIAEKKADA	956.6018	Stomach (1 h)
343	YAVAVVKKGSN	1134.64	Stomach (0.5 h)	358	AKLGGRPT	1286.683	Stomach (1 h)
344	YAVAVVKKGSNF	1281.708	Stomach (0.5 h)	359	AKLGGRPTY	1929.068	Stomach (1 h)
345	YAVAVVKKGSNFQ	1409.767	Stomach (0.5 h)	360	AVAVVKKGSNFQ	877.5022	Stomach (1 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
361	AVAVVKKGSNFQL	1359.787	Stomach (1 h)	376	EAGRDPYKLRP	1300.689	Stomach (1 h)
362	AVTLDGGM	778.3531	Stomach (1 h)	377	EAGRDPYKLRPV	1399.757	Stomach (1 h)
363	AVTLDGGMV	877.4215	Stomach (1 h)	378	EAGRDPYKLRPVA	1470.794	Stomach (1 h)
364	AVTLDGGMVF	1024.49	Stomach (1 h)	379	EAGRDPYKLRPVAA	1541.831	Stomach (1 h)
365	AVVARSVDG	872.4716	Stomach (1 h)	380	EIYGTKESPQTHY	1551.721	Stomach (1 h)
366	AVVARSVDGK	1000.567	Stomach (1 h)	381	ENLPEKA	799.4075	Stomach (1 h)
367	AVVARSVDGKE	1129.609	Stomach (1 h)	382	ENLPEKADR	1070.536	Stomach (1 h)
368	AVVARSVDGKED	1244.636	Stomach (1 h)	383	ENLPEKADRDRD	1185.563	Stomach (1 h)
369	AVVARSVDGKEDL	1357.72	Stomach (1 h)	384	ENLPEKADRDRDQ	1313.621	Stomach (1 h)
370	AVVKKGSNFQL	1189.682	Stomach (1 h)	385	ENLPEKADRDRDQY	1476.684	Stomach (1 h)
371	AVVSRSDRAAHVKQ	1522.833	Stomach (1 h)	386	ENLPEKADRDRDQYE	1605.727	Stomach (1 h)
372	DGAGDVA	603.25	Stomach (1 h)	387	ENTNGESTADWAKNL	1648.733	Stomach (1 h)
373	DPYKLRPVA	1057.592	Stomach (1 h)	388	EPLQGAVAK	911.5076	Stomach (1 h)
374	EAGRDPY	806.3559	Stomach (1 h)	389	EPLQGAVAKF	1058.576	Stomach (1 h)
375	EAGRDPYKLR	1203.636	Stomach (1 h)	390	ESPQTHY	860.3664	Stomach (1 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
391	FGKNGKN	763.3977	Stomach (1 h)	406	GSPPGQR	697.3507	Stomach (1 h)
392	FGKNKSR	835.4664	Stomach (1 h)	407	GSPPGQRDL	925.4617	Stomach (1 h)
393	FGSPPGQR	844.4192	Stomach (1 h)	408	GTKESPQTHY	1146.531	Stomach (1 h)
394	FGSPPGQRD	959.4461	Stomach (1 h)	409	GTKESPQTHYY	1309.594	Stomach (1 h)
395	FGSPPGQRDL	1072.53	Stomach (1 h)	410	GWIIPMGILR	1154.663	Stomach (1 h)
396	FGSPPGQRDLL	1185.614	Stomach (1 h)	411	GWIIPMGILRP	1251.716	Stomach (1 h)
397	FKDSALGF	883.444	Stomach (1 h)	412	GWIIPMGILRPYL	1527.864	Stomach (1 h)
398	FKSETKN	852.4341	Stomach (1 h)	413	HLAVAPNHAVVS	1213.657	Stomach (1 h)
399	FKSETKNL	965.5182	Stomach (1 h)	414	IAEKKAD	773.4283	Stomach (1 h)
400	FKSETKNLL	1078.602	Stomach (1 h)	415	IAEKKADA	844.4654	Stomach (1 h)
401	GESTADWAKNL	1190.557	Stomach (1 h)	416	IAEKKADAVTL	1157.666	Stomach (1 h)
402	GILRPYL	830.5014	Stomach (1 h)	417	IAEKKADAVTLDGGMV	1632.839	Stomach (1 h)
403	GRDPYKLRP	1100.609	Stomach (1 h)	418	IIPMGILR	911.5626	Stomach (1 h)
404	GRDPYKLRPV	1199.678	Stomach (1 h)	419	IIPMGILRP	1008.615	Stomach (1 h)
405	GRDPYKLRPVA	1270.715	Stomach (1 h)	420	IIPMGILRPY	1171.679	Stomach (1 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
421	IIPMGILRPYL	1284.763	Stomach (1 h)	436	LAVAPNHA	791.429	Stomach (1 h)
422	IPMGILRPYL	1171.679	Stomach (1 h)	437	LAVAPNHAVVSR	1232.699	Stomach (1 h)
423	IYGTKESPQTHY	1422.678	Stomach (1 h)	438	LDGTRKPVTE	1114.598	Stomach (1 h)
424	KAQEKFGKNKSR	1419.795	Stomach (1 h)	439	LFKDSALGF	996.528	Stomach (1 h)
425	KESPQTHY	988.4614	Stomach (1 h)	440	LGGRPTY	762.4024	Stomach (1 h)
426	KESPQTHYY	1151.525	Stomach (1 h)	441	LPEKADR	827.4501	Stomach (1 h)
427	KGEADAL	702.3548	Stomach (1 h)	442	LPEKADRQY	1233.599	Stomach (1 h)
428	KGEADALN	816.3977	Stomach (1 h)	443	LRIPSKVD	926.5549	Stomach (1 h)
429	KGEADALNL	929.4818	Stomach (1 h)	444	LRIPSKVDS	1013.587	Stomach (1 h)
430	KGEADALNLDG	1101.53	Stomach (1 h)	445	LRIPSKVDSA	1084.624	Stomach (1 h)
431	KGEADALNLDGGY	1321.615	Stomach (1 h)	446	LRIPSKVDSAL	1197.708	Stomach (1 h)
432	LAQVPSH	750.4024	Stomach (1 h)	447	LSKAQEKFGKNK	1376.778	Stomach (1 h)
433	LAQVPSHA	821.4395	Stomach (1 h)	448	LSKAQEKFGKNKSR	1619.911	Stomach (1 h)
434	LAQVPSHAVVA	1090.614	Stomach (1 h)	449	LSKAQEKFGKNKSRS	1706.943	Stomach (1 h)
435	LAVAPNH	720.3919	Stomach (1 h)	450	LSKAQEKFGKNKSRSF	1854.011	Stomach (1 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
451	LTTLKNL	801.496	Stomach (1 h)	466	RETAEVVKAR	1187.626	Stomach (1 h)
452	LTTLKNLR	957.5971	Stomach (1 h)	467	RIPSKVDS	900.5029	Stomach (1 h)
453	LTTLKNLRE	1086.64	Stomach (1 h)	468	RIPSKVDSA	971.54	Stomach (1 h)
454	LTTLKNLRETA	1258.725	Stomach (1 h)	469	RSAGWIIP	898.5025	Stomach (1 h)
455	LTTLKNLRETAE	1387.767	Stomach (1 h)	470	RSVDGKED	904.425	Stomach (1 h)
456	LTTLKNLRETAEE	1516.81	Stomach (1 h)	471	RSVDGKEDL	1017.509	Stomach (1 h)
457	LTTLKNLRETAEEVK	1743.973	Stomach (1 h)	472	RTAGWNIP	913.477	Stomach (1 h)
458	LYLGSRY	870.4599	Stomach (1 h)	473	SAGWIIPMGILR	1312.733	Stomach (1 h)
459	MGILRPYL	961.5419	Stomach (1 h)	474	SAGWIIPMGILRP	1409.785	Stomach (1 h)
460	NLPEKADR	941.493	Stomach (1 h)	475	SAGWIIPMGILRPY	1572.849	Stomach (1 h)
461	PEKADR	829.393	Stomach (1 h)	476	SAGWIIPMGILRPYL	1685.933	Stomach (1 h)
462	PEKADRQY	1120.515	Stomach (1 h)	477	SDRAAHVKQVL	1222.678	Stomach (1 h)
463	PSKVDSAL	815.4389	Stomach (1 h)	478	SKAQEKFGKNKSR	1506.827	Stomach (1 h)
464	QLFGSPPGQRD	1200.589	Stomach (1 h)	479	SKAQEKFGKNKSRS	1593.859	Stomach (1 h)
465	RPYKLRPVA	1213.693	Stomach (1 h)	480	SLEPLQGA	813.4232	Stomach (1 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
481	SLEPLQGAVA	801.496	Stomach (1 h)	496	VARSVDGKEDL	1187.615	Stomach (1 h)
482	SLEPLQGAVAK	957.5971	Stomach (1 h)	497	VAVVKKANEGL	1126.671	Stomach (1 h)
483	SLEPLQGAVAKF	1086.64	Stomach (1 h)	498	VAVVKKANEGLTWNSL	1727.957	Stomach (1 h)
484	SPPGQRDL	1258.725	Stomach (1 h)	499	VAVVKKANEGLTWNSL	1856.052	Stomach (1 h)
485	SVDGKEDL	1387.767	Stomach (1 h)	500	VAVVKKGSNFQ	1175.666	Stomach (1 h)
486	TAEVKAR	1516.81	Stomach (1 h)	501	VAVVKKGSNFQL	1288.75	Stomach (1 h)
487	TAEVKARY	1743.973	Stomach (1 h)	502	VDRTAGWNIP	1127.572	Stomach (1 h)
488	TAGWNIPMGL	870.4599	Stomach (1 h)	503	VKETTVE	951.4913	Stomach (1 h)
489	TAVDRTAGWNIP	961.5419	Stomach (1 h)	504	VKKGSNFQ	906.4923	Stomach (1 h)
490	TESLEPLQG	941.493	Stomach (1 h)	505	VKKGSNFQL	1019.576	Stomach (1 h)
491	TESLEPLQGAVA	829.393	Stomach (1 h)	506	VLKGEADAL	914.5073	Stomach (1 h)
492	TESLEPLQGAVAK	1120.515	Stomach (1 h)	507	VLKGEADALN	1028.55	Stomach (1 h)
493	TESLEPLQGAVAKF	815.4389	Stomach (1 h)	508	VLKGEADALNL	1141.634	Stomach (1 h)
494	TTLKNLR	1200.589	Stomach (1 h)	509	VLLHQQAL	920.5443	Stomach (1 h)
495	TTLKNLRE	1213.693	Stomach (1 h)	510	VLRPTEG	770.4286	Stomach (1 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
511	VVARSVDGKED	1173.599	Stomach (1 h)	526	AEDVGDVA	774.3396	Stomach (2 h)
512	VVARSVDGKEDL	1286.683	Stomach (1 h)	527	AGDDQGLD	789.3141	Stomach (2 h)
513	VVKKGSNFQ	1005.561	Stomach (1 h)	528	AKLGGRPTY	961.5345	Stomach (2 h)
514	VVSRSDRAAHVKQ	1451.796	Stomach (1 h)	529	AKLGGRPTYEE	1219.62	Stomach (2 h)
515	WAKNLNR	900.4929	Stomach (1 h)	530	ALNLDGGY	821.3919	Stomach (2 h)
516	WAKNLNRED	1144.563	Stomach (1 h)	531	AVTLDGGMVF	1008.495	Stomach (2 h)
517	WIIPMGILR	1097.642	Stomach (1 h)	532	AVVARSVDGKE	1129.609	Stomach (2 h)
518	WIIPMGILRP	1194.695	Stomach (1 h)	533	AVVARSVDGKED	1244.636	Stomach (2 h)
519	WIIPMGILRPY	1357.758	Stomach (1 h)	534	AVVARSVDGKEDL	1357.72	Stomach (2 h)
520	WIIPMGILRPYL	1470.842	Stomach (1 h)	535	AVVARSVDGKEDLIW	1656.884	Stomach (2 h)
521	WTESLEPLQGAVAK	1527.793	Stomach (1 h)	536	AVVKKGSNFQL	1189.682	Stomach (2 h)
522	YAVAVVKKGSN	1134.64	Stomach (1 h)	537	EAGRDPY	806.3559	Stomach (2 h)
523	YAVAVVKKGSNFQ	1409.767	Stomach (1 h)	538	EAGRDPYKLRPVA	1470.794	Stomach (2 h)
524	YAVAVVKKGSNFQL	1522.851	Stomach (1 h)	539	EEVKARY	893.4606	Stomach (2 h)
525	YGYTGAFR	933.4345	Stomach (1 h)	540	ENLPEKA	799.4075	Stomach (2 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
541	ENLPEKADR	1070.536	Stomach (2 h)	556	GWNIPMGL	902.432	Stomach (2 h)
542	ENLPEKADRQ	1313.621	Stomach (2 h)	557	IAEKKAD	773.4283	Stomach (2 h)
543	ENLPEKADRQY	1476.684	Stomach (2 h)	558	IAEKKADAVTL	1157.666	Stomach (2 h)
544	ENLPEKADRQYE	1605.727	Stomach (2 h)	559	IIPMGILRPY	1171.679	Stomach (2 h)
545	FGSPPGQR	844.4192	Stomach (2 h)	560	IIPMGILRPYL	1284.763	Stomach (2 h)
546	FGSPPGQRD	959.4461	Stomach (2 h)	561	IIPMGILRPYLS	1371.795	Stomach (2 h)
547	FGSPPGQRDL	1072.53	Stomach (2 h)	562	ILRPYLS	860.512	Stomach (2 h)
548	FGSPPGQRDLL	1185.614	Stomach (2 h)	563	ILRPYLSW	1046.591	Stomach (2 h)
549	FKDSALGF	883.444	Stomach (2 h)	564	ILRPYLSWT	1147.639	Stomach (2 h)
550	FKSETKNL	965.5182	Stomach (2 h)	565	ILRPYLSWTE	1276.682	Stomach (2 h)
551	FKSETKNLL	1078.602	Stomach (2 h)	566	KGEADAL	702.3548	Stomach (2 h)
552	FVKETTVF	969.5172	Stomach (2 h)	567	KGEADALNL	929.4818	Stomach (2 h)
553	GRDPYKLRPVA	1270.715	Stomach (2 h)	568	KKADAVTL	844.5018	Stomach (2 h)
554	GTKESPQTHY	1146.531	Stomach (2 h)	569	KLGAPSIT	785.4647	Stomach (2 h)
555	GTKESPQTHYY	1309.594	Stomach (2 h)	570	LAQVPSH	750.4024	Stomach (2 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
571	LAVAPNH	720.3919	Stomach (2 h)	586	LTTLKNLRETAE	1387.767	Stomach (2 h)
572	LAVAPNHA	791.429	Stomach (2 h)	587	LYLGSRY	870.4599	Stomach (2 h)
573	LAVAPNHAVVSRSD	1434.758	Stomach (2 h)	588	MGILRPY	848.4578	Stomach (2 h)
574	LFKDSALGF	996.528	Stomach (2 h)	589	MGILRPYL	961.5419	Stomach (2 h)
575	LRIPSKVD	926.5549	Stomach (2 h)	590	MGILRPYLS	1048.574	Stomach (2 h)
576	LRIPSKVDS	1013.587	Stomach (2 h)	591	MGILRPYLSW	1234.653	Stomach (2 h)
577	LRIPSKVDSA	1084.624	Stomach (2 h)	592	MGILRPYLSWTE	1464.744	Stomach (2 h)
578	LRIPSKVDSAL	1197.708	Stomach (2 h)	593	PEKADRDQY	1120.515	Stomach (2 h)
579	LRIPSKVDSALY	1360.771	Stomach (2 h)	594	RAAHVKQVL	1020.619	Stomach (2 h)
580	LSKAQEKFGKNKSR	1619.911	Stomach (2 h)	595	RPYLSWTE	1050.513	Stomach (2 h)
581	LSKAQEKFGKNKSRS	1706.943	Stomach (2 h)	596	RSVDGKEDL	1017.509	Stomach (2 h)
582	LTTLKNL	801.496	Stomach (2 h)	597	RTAGWNIP	913.477	Stomach (2 h)
583	LTTLKNLR	957.5971	Stomach (2 h)	598	RTAGWNIPMG	1101.539	Stomach (2 h)
584	LTTLKNLRE	1086.64	Stomach (2 h)	599	RTAGWNIPMGL	1214.623	Stomach (2 h)
585	LTTLKNLRETA	1258.725	Stomach (2 h)	600	SLEPLQGA	813.4232	Stomach (2 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
601	SPPGQRD	755.3562	Stomach (2 h)	616	WAKNLNRED	1144.563	Stomach (2 h)
602	TAEVKARY	1065.545	Stomach (2 h)	617	WIIPMGILRP	1194.695	Stomach (2 h)
603	TAGWNIPMGL	1058.522	Stomach (2 h)	618	WIIPMGILRPYL	1486.837	Stomach (2 h)
604	TAVDRTAGWNIPMGL	1600.803	Stomach (2 h)	619	WIIPMGILRPYLS	1573.869	Stomach (2 h)
605	TESLEPLQGA	1043.514	Stomach (2 h)	620	YAVAVVKKGGSNFQ	1409.767	Stomach (2 h)
606	TTLKNLRE	973.5556	Stomach (2 h)	621	YAVAVVKKGGSNFQL	1522.851	Stomach (2 h)
607	VAVVKKANEGLTW	1413.798	Stomach (2 h)	622	DQLQGRK	843.4563	Stomach (3 h)
608	VAVVKKANEGLTWNSL	1727.957	Stomach (2 h)	623	LAVAPNH	720.3919	Stomach (3 h)
609	VAVVKKGGSNFQL	1288.75	Stomach (2 h)	624	AAHVKQVLLHQQ	1370.778	Duodenum (0.5 h)
610	VKETTVF	822.4487	Stomach (2 h)	625	AAHVKQVLLHQQAL	1554.899	Duodenum (0.5 h)
611	VLKGEADAL	914.5073	Stomach (2 h)	626	AVAVVKKGGSNFQLDQLQG	1901.037	Duodenum (0.5 h)
612	VLKGEADALNL	1141.634	Stomach (2 h)	627	CAVGPEE	703.2847	Duodenum (0.5 h)
613	VVKKANEGLTW	1243.693	Stomach (2 h)	628	DCVLRPT	802.4007	Duodenum (0.5 h)
614	VVKKANEGLTWNSL	1557.852	Stomach (2 h)	629	DGGMVFEAGRDPY	1412.603	Duodenum (0.5 h)
615	WAKNLNR	900.4929	Stomach (2 h)	630	GDDQGLDK	846.3719	Duodenum (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
631	GGRPTYEEYLGTEY	1633.726	Duodenum (0.5 h)	646	LDCVLRPT	915.4848	Duodenum (0.5 h)
632	GGRPTYEEYLGTEYVT	1833.842	Duodenum (0.5 h)	647	LGGRPTYEEYLGTEY	1746.81	Duodenum (0.5 h)
633	GILRPYL	830.5014	Duodenum (0.5 h)	648	LGGRPTYEEYLGTEYVT	1946.926	Duodenum (0.5 h)
634	GILRPYLSWTESLEPL	1872.999	Duodenum (0.5 h)	649	LGGRPTYEEYLGTEYVTA	2017.963	Duodenum (0.5 h)
635	GRPTYEEYLGTEY	1576.705	Duodenum (0.5 h)	650	LGGRPTYEEYLGTEYVTAI	2131.047	Duodenum (0.5 h)
636	HLAQVPS	750.4024	Duodenum (0.5 h)	651	LGGRPTYEEYLGTEYVTA	2202.084	Duodenum (0.5 h)
637	KADAVTLDGGMVF	1338.649	Duodenum (0.5 h)	652	LRIPSKVDSAL	1197.708	Duodenum (0.5 h)
638	KANEGLT	731.3813	Duodenum (0.5 h)	653	LRPVAAEIY	1030.581	Duodenum (0.5 h)
639	KGSNFQLDQLQGR	1489.764	Duodenum (0.5 h)	654	LSWTESLEPLQGAVAK	1727.909	Duodenum (0.5 h)
640	KLRPVAAEIYGTKESPO	1886.026	Duodenum (0.5 h)	655	LSWTESLEPLQGAVAKF	1874.978	Duodenum (0.5 h)
641	LAVAVVKK	826.564	Duodenum (0.5 h)	656	LTTLKNLR	957.5971	Duodenum (0.5 h)
642	LAVAVVKKKA	897.6011	Duodenum (0.5 h)	657	PSKVDSALYL	1091.586	Duodenum (0.5 h)
643	LAVAVVKKKANEGLTW	1597.919	Duodenum (0.5 h)	658	QLDQLQGR	956.5039	Duodenum (0.5 h)
644	LAVAVVKKKANEGLTWN	1711.962	Duodenum (0.5 h)	659	QLFGSPPGQR	1085.562	Duodenum (0.5 h)
645	LAVAVVKKKANEGLTWNS	1912.078	Duodenum (0.5 h)	660	QVLLHQQALFGK	1380.788	Duodenum (0.5 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
661	SAGWIIPMGILRPY	1588.844	Duodenum (0.5 h)	676	AVAPNHAVVS	963.5138	Duodenum (1 h)
662	SAGWIIPMGILRPYL	1701.928	Duodenum (0.5 h)	677	HLAVAPNH	857.4508	Duodenum (1 h)
663	SCAPGADPK	844.3749	Duodenum (0.5 h)	678	KLGGRPT	727.4341	Duodenum (1 h)
664	SFQLFGSPPGQR	1319.662	Duodenum (0.5 h)	679	LAQVPSH	750.4024	Duodenum (1 h)
665	SVDGKEDLIWKL	1401.75	Duodenum (0.5 h)	680	LAVAPNH	720.3919	Duodenum (1 h)
666	SVDGKEDLIWKLLSK	1729.961	Duodenum (0.5 h)	681	LAVAPNHA	791.429	Duodenum (1 h)
667	YAVAVVKK	876.5433	Duodenum (0.5 h)	682	RSVDGKED	904.425	Duodenum (1 h)
668	YAVAVVKKGSNF	1281.708	Duodenum (0.5 h)	683	AQVPSHA	708.3555	Duodenum (1 h)
669	YAVAVVKKGSNFQ	1409.767	Duodenum (0.5 h)	684	AVAPNHAV	777.4133	Duodenum (1 h)
670	YAVAVVKKGSNFQL	1522.851	Duodenum (0.5 h)	685	DCVLRPT	802.4007	Duodenum (1 h)
671	HLAQVPS	750.4024	Duodenum (1 h)	686	KANEGLT	731.3813	Duodenum (1 h)
672	VLRPTEG	770.4286	Duodenum (1 h)	687	KLGAPSIT	785.4647	Duodenum (1 h)
673	SCAPGADPK	844.3697	Duodenum (1 h)	688	QDGAGDV	660.2715	Duodenum (1 h)
674	LDCVLRPT	915.4848	Duodenum (1 h)	689	RETAEV	832.3926	Duodenum (1 h)
675	AVAPNHA	678.3449	Duodenum (1 h)	690	RIPSKVD	813.4708	Duodenum (1 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
691	SAGWIIPMGILRPY	1588.844	Duodenum (0.5 h)	706	APVDAFK	746.3963	Jejunum (0.5 h)
692	SAGWIIPMGILRPYL	1701.928	Duodenum (0.5 h)	707	APVDAFKECHLAQVPS	2344.211	Jejunum (0.5 h)
693	SCAPGADPK	844.3749	Duodenum (0.5 h)	708	AQEKFGK	806.4286	Jejunum (0.5 h)
694	AAHVKQVLLHQALFGK	1887.084	Jejunum (0.5 h)	709	AQVPSHAVVAR	1133.631	Jejunum (0.5 h)
695	AAHVKQVLLHQALFGK	2186.244	Jejunum (0.5 h)	710	AVAPNHAVVS	963.5138	Jejunum (0.5 h)
696	ADAVTLDGGMVF	1194.559	Jejunum (0.5 h)	711	AVAPNHAVVSR	1119.615	Jejunum (0.5 h)
697	ADAVTLDGGMVFEAGR	1607.761	Jejunum (0.5 h)	712	AVAVVKK	713.48	Jejunum (0.5 h)
698	ADAVTLDGGMVFEAGRDPY	1982.904	Jejunum (0.5 h)	713	AVTLDGGMVFEAGR	1924.935	Jejunum (0.5 h)
699	ADAVTLDGGMVFEAGRDPYK	2110.999	Jejunum (0.5 h)	714	CGLVPVLAENR	1169.623	Jejunum (0.5 h)
700	ADRDQYELL	1121.535	Jejunum (0.5 h)	715	DGGMVFEAGRDPYK	1540.698	Jejunum (0.5 h)
701	AEKKADA	731.3813	Jejunum (0.5 h)	716	DLLFKDSALGF	1224.639	Jejunum (0.5 h)
702	AFALECIR	921.4742	Jejunum (0.5 h)	717	DLLFKDSALGFLR	1493.824	Jejunum (0.5 h)
703	ALNLDGGYIYTAGK	1454.741	Jejunum (0.5 h)	718	DSALGFLR	877.4658	Jejunum (0.5 h)
704	ANEGLTWNSLK	1231.62	Jejunum (0.5 h)	719	DSALGFLRIPSK	1302.73	Jejunum (0.5 h)
705	ANEGLTWNSLKDKK	1602.837	Jejunum (0.5 h)	720	DSALGFLRIPSKVDSA	1951.042	Jejunum (0.5 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
721	DSALGFLRIPSKVDSALYL	2364.28	Jejunum (0.5 h)	736	GILRPYLSWTESLEPLQ	2427.316	Jejunum (0.5 h)
722	EAGRDPYK	934.4508	Jejunum (0.5 h)	737	GKCGLVPVLAENR	1354.739	Jejunum (0.5 h)
723	ECHLAQVPSHAVVAR	1615.825	Jejunum (0.5 h)	738	GLVPVLAENRK	1194.708	Jejunum (0.5 h)
724	EKYYGYTGAFR	1353.635	Jejunum (0.5 h)	739	GSNFQLDQLQGR	1361.669	Jejunum (0.5 h)
725	ENTNGESTADWAK	1421.606	Jejunum (0.5 h)	740	GSNFQLDQLQGRK	1489.764	Jejunum (0.5 h)
726	EPYFGYSGAFK	1264.576	Jejunum (0.5 h)	741	GSPPGQR	697.3507	Jejunum (0.5 h)
727	ESPQTHYY	1023.43	Jejunum (0.5 h)	742	HSSLDCVLRPTEGYLA	2384.289	Jejunum (0.5 h)
728	ESPQTHYYAVAVVK	1590.804	Jejunum (0.5 h)	743	IAEKKADA	844.4654	Jejunum (0.5 h)
729	ESPQTHYYAVAVVKK	1718.899	Jejunum (0.5 h)	744	IPSKVDSALY	1091.586	Jejunum (0.5 h)
730	ETTVFENLPEK	1305.645	Jejunum (0.5 h)	745	IPSKVDSALYLGSR	1504.825	Jejunum (0.5 h)
731	ETTVFENLPEKADR	1647.81	Jejunum (0.5 h)	746	ISQPEWFK	1033.523	Jejunum (0.5 h)
732	FGYSGAFK	875.4178	Jejunum (0.5 h)	747	KADAVTLD	831.4338	Jejunum (0.5 h)
733	GEADALNLDGGYIY	1469.667	Jejunum (0.5 h)	748	KADAVTLDGGMVF	1322.654	Jejunum (0.5 h)
734	GEADALNLDGGYIYTAGK	1826.869	Jejunum (0.5 h)	749	KADAVTLDGGMVFEAGR	1735.856	Jejunum (0.5 h)
735	GGMVFEAGRDPYK	1425.671	Jejunum (0.5 h)	750	KADAVTLDGGMVFEAGR	1866.878	Jejunum (0.5 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
751	KDAVTLDGGMVFEAGR	2126.994	Jejunum (0.5 h)	766	LAQVPSHAVVAR	1246.715	Jejunum (0.5 h)
752	KDAVTLDGGMVFEAGR	2239.094	Jejunum (0.5 h)	767	LAVAPNHAVVSR	1232.699	Jejunum (0.5 h)
753	KANEGLT	731.3813	Jejunum (0.5 h)	768	LAVAVVK	698.4691	Jejunum (0.5 h)
754	KANEGLTWNSLK	1359.715	Jejunum (0.5 h)	769	LAVAVVKK	826.564	Jejunum (0.5 h)
755	KANEGLTWNSLKDKK	1730.932	Jejunum (0.5 h)	770	LFGSPPGQR	957.5032	Jejunum (0.5 h)
756	KGSNFQLDQLQGR	1489.764	Jejunum (0.5 h)	771	LGAPSITCVR	1015.548	Jejunum (0.5 h)
757	KGSNFQLDQLQGRK	1617.859	Jejunum (0.5 h)	772	LGAPSITCVRR	1171.65	Jejunum (0.5 h)
758	KLGRPTCVRR	1299.745	Jejunum (0.5 h)	773	LGGRPTYEEY	1183.551	Jejunum (0.5 h)
759	KLGRPT	727.4341	Jejunum (0.5 h)	774	LGGRPTYEEYLGTE	1583.747	Jejunum (0.5 h)
760	KLRPVAAEIY	1158.676	Jejunum (0.5 h)	775	LGGRPTYEEYLGTEY	1746.81	Jejunum (0.5 h)
761	KLRPVAAEIYGTK	1444.84	Jejunum (0.5 h)	776	LGGRPTYEEYLGTEYV	2557.306	Jejunum (0.5 h)
762	KPVTEAQ	771.4127	Jejunum (0.5 h)	777	LGGRPTYEEYLGTEYV	2685.401	Jejunum (0.5 h)
763	KPVTEAQSCHLAVAPNHA	2313.201	Jejunum (0.5 h)	778	LGTEYVTAIANLK	1391.766	Jejunum (0.5 h)
764	KSCHTGLGR	957.4814	Jejunum (0.5 h)	779	LGTEYVTAIANLKK	1519.861	Jejunum (0.5 h)
765	KVDSALYLGSR	1207.656	Jejunum (0.5 h)	780	LLCLDGTR	889.4691	Jejunum (0.5 h)

(continue)

No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
781	LLSKAQEKFGKNK	1489.862	Jejunum (0.5 h)	796	SFQLFGSPPGQR	1319.662	Jejunum (0.5 h)
782	LRIPSKVDSALYLGSR	1774.01	Jejunum (0.5 h)	797	SKVDSALYLGSR	1294.688	Jejunum (0.5 h)
783	LRPVAAEIY	1030.581	Jejunum (0.5 h)	798	SVDGKEDLIWK	1288.666	Jejunum (0.5 h)
784	LRPVAAEIYGTK	1316.745	Jejunum (0.5 h)	799	TESLEPLQGAVAK	1341.714	Jejunum (0.5 h)
785	LSWTESLEPLQGAVAK	1727.909	Jejunum (0.5 h)	800	VDSALYLGSR	1079.561	Jejunum (0.5 h)
786	NLLFNDNTECLAK	1493.718	Jejunum (0.5 h)	801	VLKGEADAL	914.5073	Jejunum (0.5 h)
787	NSLKDKK	831.4814	Jejunum (0.5 h)	802	VLKGEADALNLDGGY	2167.116	Jejunum (0.5 h)
788	PSKVDSALYL	1091.586	Jejunum (0.5 h)	803	VTAIANLK	828.5069	Jejunum (0.5 h)
789	QVLLHQQALFGK	1380.788	Jejunum (0.5 h)	804	VTAIANLKK	956.6018	Jejunum (0.5 h)
790	QVLLHQQALFGKNGK	1679.947	Jejunum (0.5 h)	805	VVWCAVGPEEQK	1343.654	Jejunum (0.5 h)
791	RAFALECIR	1077.575	Jejunum (0.5 h)	806	WCTISQPEWFK	1423.66	Jejunum (0.5 h)
792	SAGWIIPM	873.4418	Jejunum (0.5 h)	807	YFGYSGAFK	1038.481	Jejunum (0.5 h)
793	SAGWIIPMGILRPY	1572.849	Jejunum (0.5 h)	808	YTRVVWCAVGPEEQK	1763.867	Jejunum (0.5 h)
794	SAGWIIPMGILRPYLSWTE	3282.748	Jejunum (0.5 h)	809	YVTAIANLK	991.5702	Jejunum (0.5 h)
795	SDRAAHVK	882.4671	Jejunum (0.5 h)	810	YVTAIANLKK	1119.665	Jejunum (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
811	YYGYTGAFR	1096.498	Jejunum (0.5 h)	826	TAGWNIPMGL	1058.522	Ileum (0.5 h)
812	DSALGFLR	877.4658	Jejunum (1 h)	827	NLNREDFR	1062.521	Ileum (0.5 h)
813	SFQLFGSPPGQR	1319.662	Jejunum (1 h)	828	YYGYTGAFR	1096.498	Ileum (0.5 h)
814	NTNGEST	721.2879	Jejunum (1 h)	829	AVAPNHAVVSR	1119.615	Ileum (0.5 h)
815	EDVGDVA	703.3025	Ileum (0.5 h)	830	LGGRPTYEEY	1183.551	Ileum (0.5 h)
816	KPVTEAQ	771.4127	Ileum (0.5 h)	831	ANEGLTWNSLK	1231.62	Ileum (0.5 h)
817	VGPEEQK	785.3919	Ileum (0.5 h)	832	LGGRPTYEEYL	1296.635	Ileum (0.5 h)
818	DDQGLDK	789.3505	Ileum (0.5 h)	833	KANEGLTWNSLK	1359.715	Ileum (0.5 h)
819	QQALFGK	790.4337	Ileum (0.5 h)	834	QVLLHQALFGK	1380.788	Ileum (0.5 h)
820	LAVAVVKK	826.564	Ileum (0.5 h)	835	ENTNGESTADWAK	1421.606	Ileum (0.5 h)
821	NSLKDKK	831.4814	Ileum (0.5 h)	836	ANEGLTWNSLKDK	1474.742	Ileum (0.5 h)
822	GDDQGLDK	846.3719	Ileum (0.5 h)	837	KANEGLTWNSLKDK	1602.837	Ileum (0.5 h)
823	TAGWNIPM	904.4113	Ileum (0.5 h)	838	GGRPTYEEYLGTEY	1633.726	Ileum (0.5 h)
824	YGYTGAFR	933.4345	Ileum (0.5 h)	839	KANEGLTWNSLKDKK	1730.932	Ileum (0.5 h)
825	VTAIANLKK	956.6018	Ileum (0.5 h)	840	LGGRPTYEEYLGTEY	1746.81	Ileum (0.5 h)

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No.	Peptide sequence	Mr. calc	Sample	No.	Peptide sequence	Mr. calc	Sample
841	NTNGEST	721.2879	Ileum (0.5 h)	845	ANEGLTWNSLK	1231.62	Ileum (0.5 h)
842	DDQGLDK	789.3505	Ileum (0.5 h)	846	NLDGGYIYTAGK	1270.619	Ileum (0.5 h)
843	LAVAVVKK	826.564	Ileum (0.5 h)	847	QVLLHQQALFGK	1380.788	Ileum (0.5 h)
844	GDDQGLDK	846.3719	Ileum (0.5 h)	848	KGSNFQLDQLQGR	1489.764	Ileum (0.5 h)

Note: The 848 peptides were marked from No.1 to No.848. The peptides in the content of stomach, duodenum, jejunum and ileum were marked as No.1 – No.623,

No.624 – No.693, No.694 – No.814, No.815 – No.848, respectively.