

Electronic Supporting Information for:

**Self-assembling covalent organic frameworks functionalized
magnetic graphene hydrophilic biocomposite as an ultrasensitive
matrix for N-linked glycopeptide recognition**

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EXPERIMENTAL SECTION

Nano-Liquid chromatography tandem mass spectrometry (Nano-LC–MS/MS) analysis of glycopeptides.

Glycopeptides enriched from human serum were resuspended with 10 μ L solvent A (A: water with 0.1% formic acid; B: ACN with 0.1% formic acid) and then separated by nanoLC and analyzed by on-line electrospray tandem mass spectrometry. The experiments were performed on an EASY-nLC 1000 system (Thermo Fisher Scientific, Waltham, MA) connected to an Orbitrap Fusion mass spectrometer (Thermo Fisher Scientific, San Jose, CA) equipped with an online nano-electrospray ion source. 8 μ L peptide sample was loaded onto the trap column (Thermo Scientific Acclaim PepMap C18, 100 μ m x 2cm), with a flow of 10 μ L/min for 3 min and subsequently separated on the analytical column (Acclaim PepMap C18, 75 μ m x 25cm) with a linear gradient, from 2% B to 25% B in 105 min. The column was re-equilibrated at initial conditions for 15 min. The column flow rate was maintained at 300 nL/min and column temperature was maintained at 40 °C. The electrospray voltage of 2.0 kV versus the inlet of the mass spectrometer was used. The Orbitrap Fusion mass spectrometer was operated in the data-dependent mode to switch automatically between MS and MS/MS acquisition. Survey full-scan MS spectra (m/z 350-1500) were acquired in Orbitrap with a mass resolution of 120 000 at m/z 200. The AGC target was set to 1 000 000, and the maximum injection time was 50ms. MS/MS acquisition was performed in Orbitrap with 3 s cycle time, the resolution was 15 000 at m/z 200. The intensity threshold was 50 000, and the maximum injection time was 100 ms. The AGC target was set to 200 000, and the isolation window was 2 m/z . Ions with charge states 2+, 3+, and 4+ were sequentially fragmented by higher energy collisional dissociation (HCD) with a normalized collision energy (NCE) of 35%, fixed first mass was set at 110. In all cases, one microscan was recorded using dynamic exclusion of 30 seconds.

Database search. Tandem mass spectra were extracted by Proteome Discoverer software (Thermo Fisher Scientific, version 1.4.0.288). Charge state deconvolution and deisotoping were not performed. All MS/MS samples were analyzed using Mascot (Matrix Science, London, UK; version 2.3.2). The database was the Human UniProtKB/Swiss-Prot database (Release 2015-03-11, with 20199 sequences). Raw files generated by the Orbitrap Fusion were searched directly using a 10ppm precursor mass tolerance and a 50 mmu fragment mass tolerance. The enzyme specificity with trypsin was used. Up to two missed cleavages were allowed and peptides with at least 7 amino acids were retained. Carbamidomethyl on cysteine was set as a fixed modification. Oxidation on methionine and Deamidation on asparagine were set as variable modifications. Use the percolator algorithm to control peptide level false discovery rates (FDR) lower than 1%. The Asn modification that did not occur in the N-X-S/T (X≠P) sequon was eliminated to ensure the false positive rate below 1% for the identified glycosylation sites.

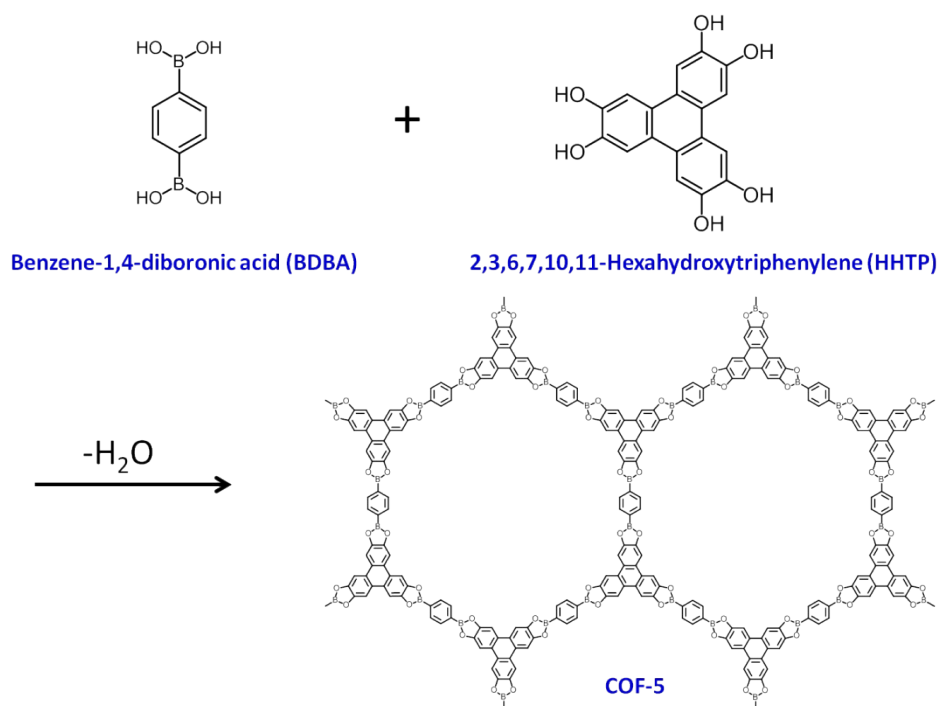


Fig. S1 Schematic representation for the direct construction of COF-5.

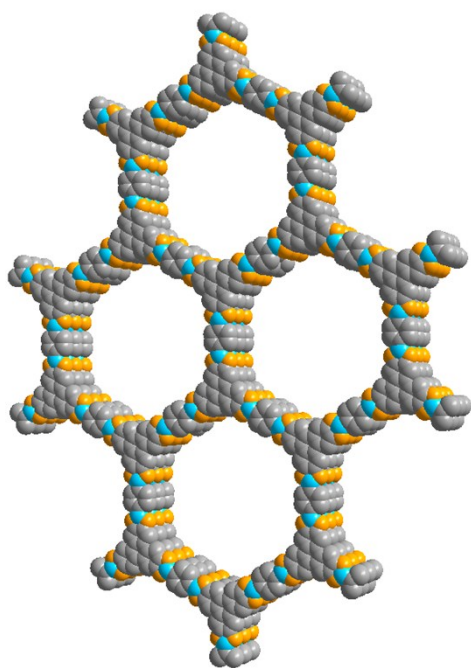


Fig. S2 The extended structures based on the eclipsed arrangements. C, gray; O, orange; B, blue. H atoms are omitted for clarity.

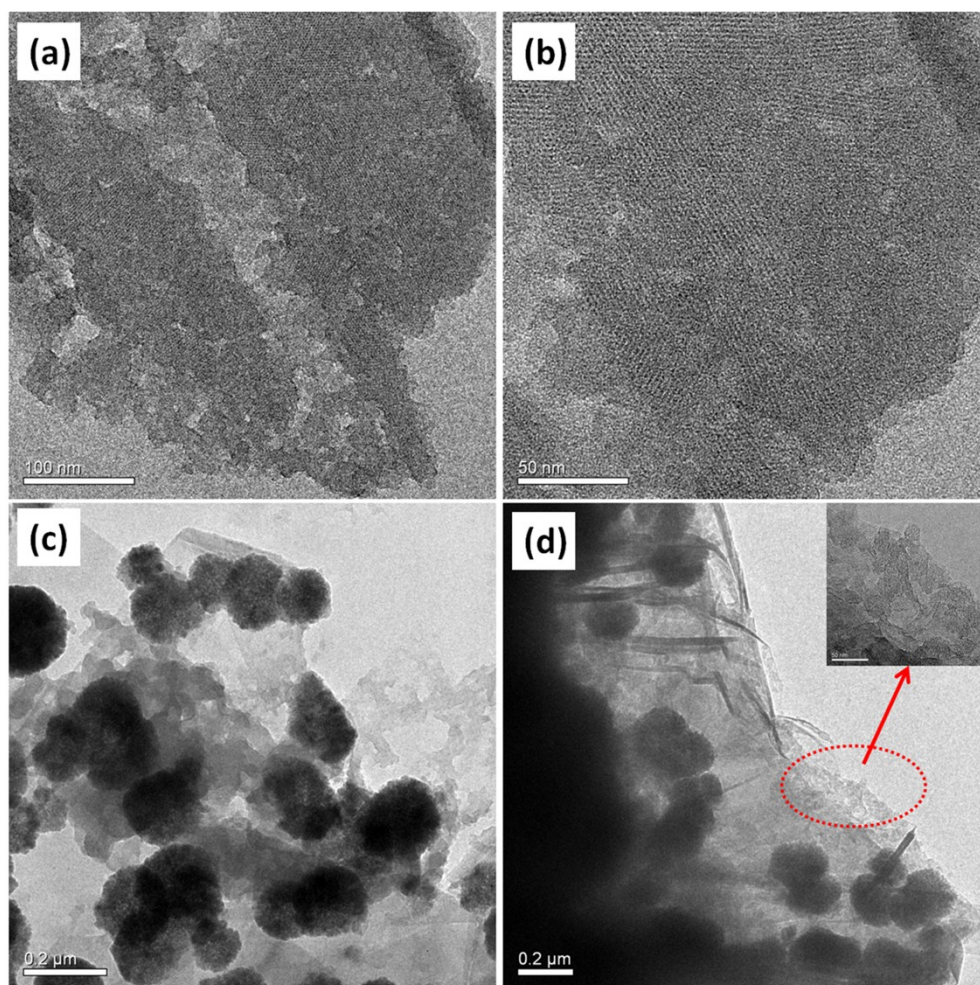


Fig. S3 The TEM images of (a, b) COF-5 materials, (c) MagG@COF-5-0.5C and (d) MagG@COF-5 biocomposite.

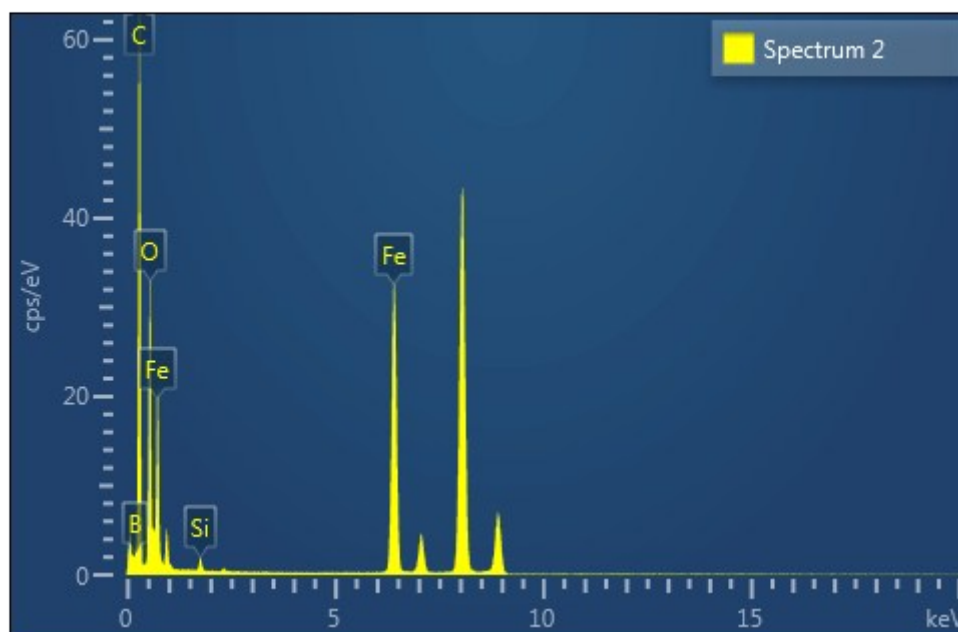


Fig. S4 The energy dispersive X-ray (EDX) spectrum data of MagG@COF-5 biocomposite.

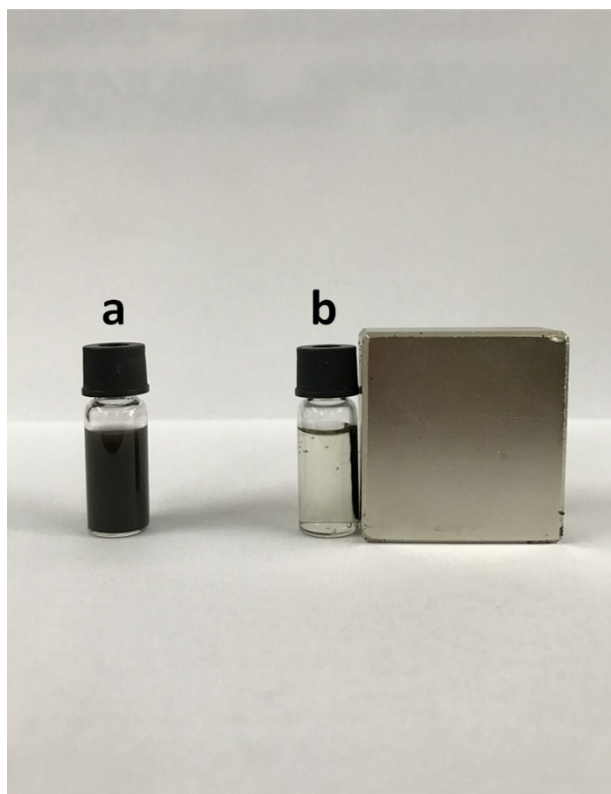


Fig. S5 The photos of the aqueous dispersion of MagG@COF-5 biocomposite: (a) before and (b) after separation by a magnet for 5 seconds.

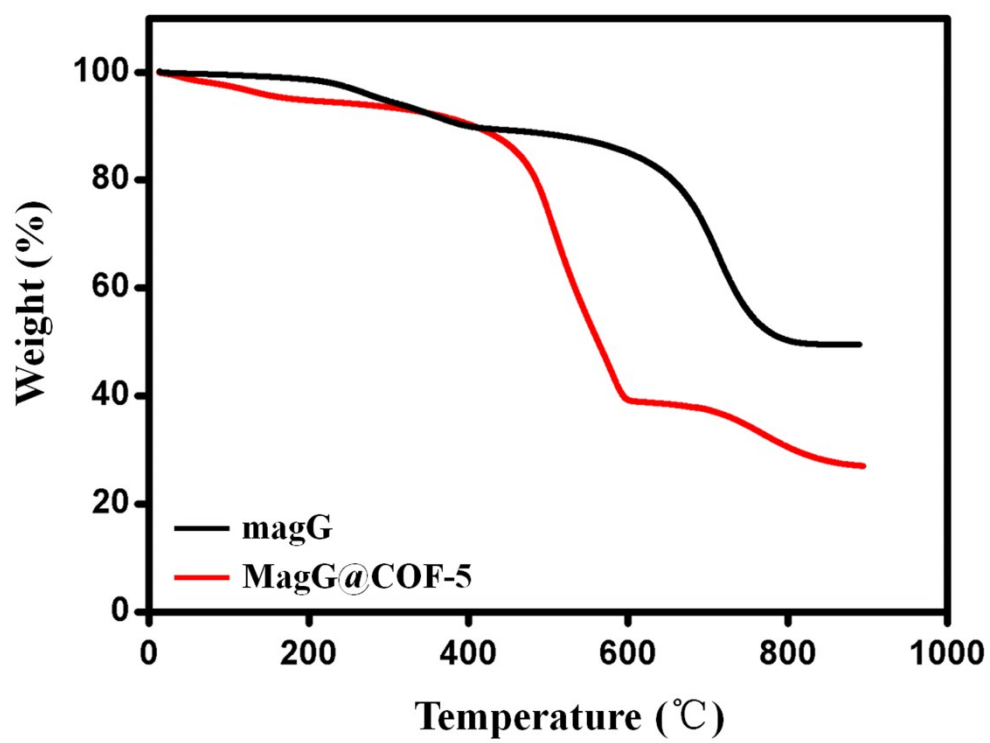


Fig. S6 TGA curves of magG and MagG@COF-5 biocomposite.

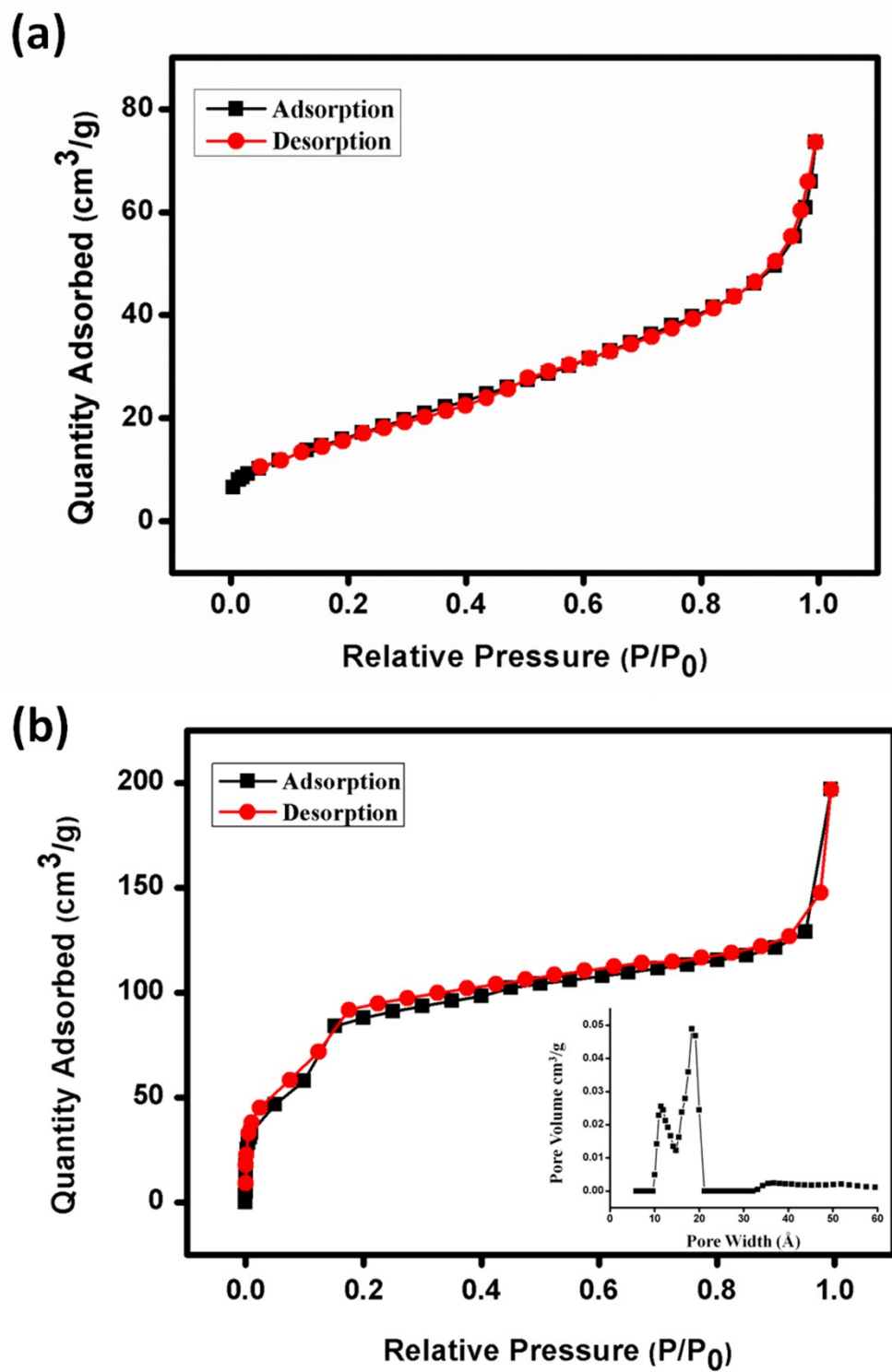


Fig. S7 N_2 adsorption/desorption isotherms of (a) magG and (b) MagG@COF-5 biocomposite.

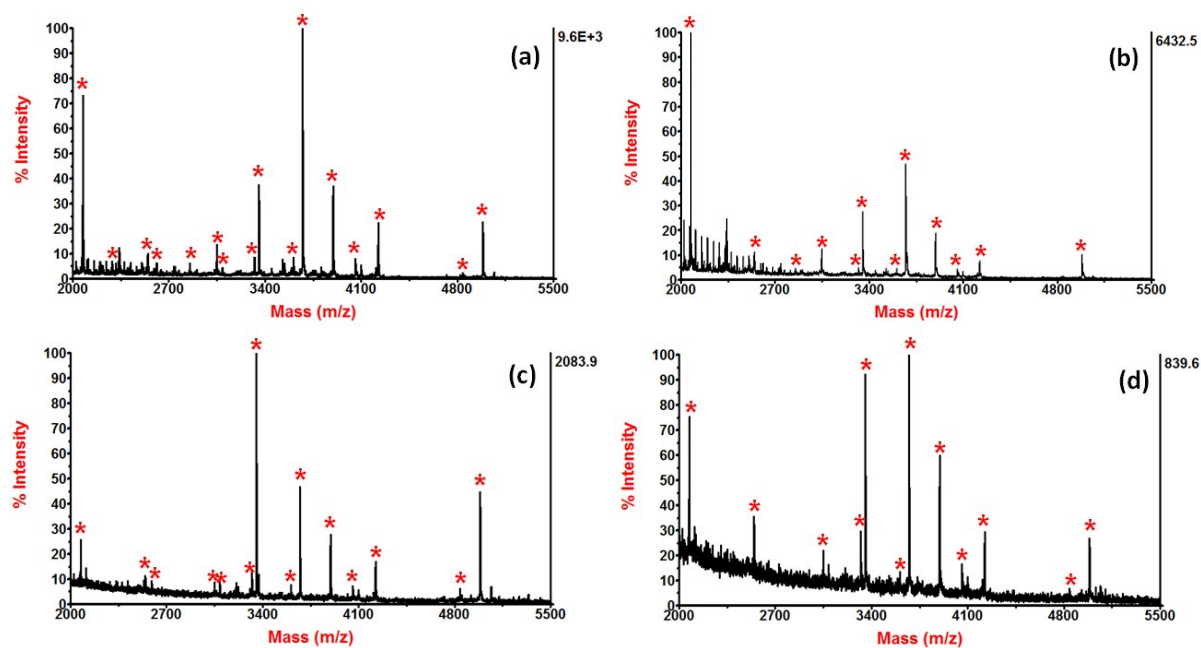


Fig. S8 MALDI-TOF-MS analysis of peptides derived from HRP enriched by MagG@COF-5 biocomposite with different buffer solution (a) ACN/H₂O/TFA (95:4:1, v/v/v); (b) ACN/H₂O/TFA (95:4.9:0.1, v/v/v); (c) ACN/H₂O/TFA (85:14:1, v/v/v) and (d) ACN/H₂O/TFA (85:14.9:0.1, v/v/v).

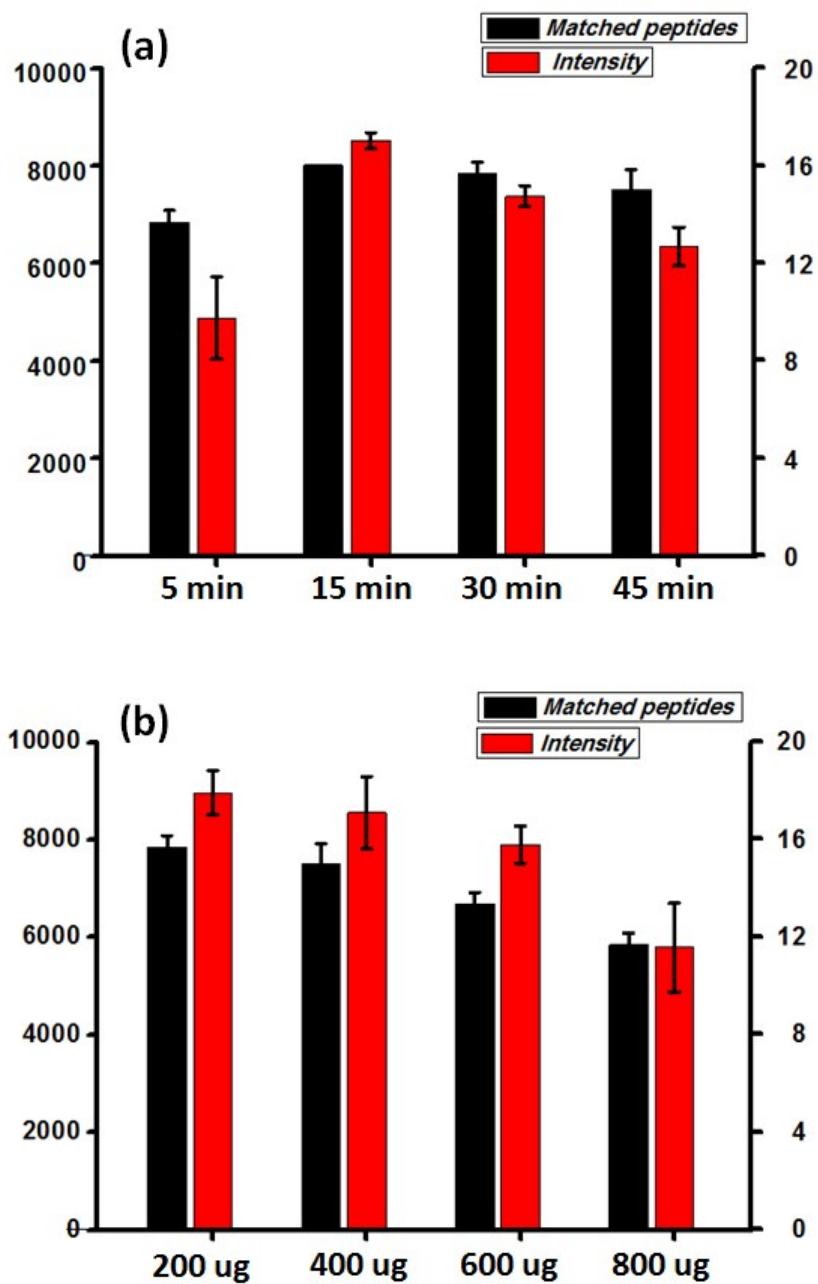


Fig. S9 The effect of enrichment time and different quantity of the MagG@COF-5 biocomposite in glycopeptides enriched from 10^{-6} M HRP tryptic digest (a, b) through three parallel tests.

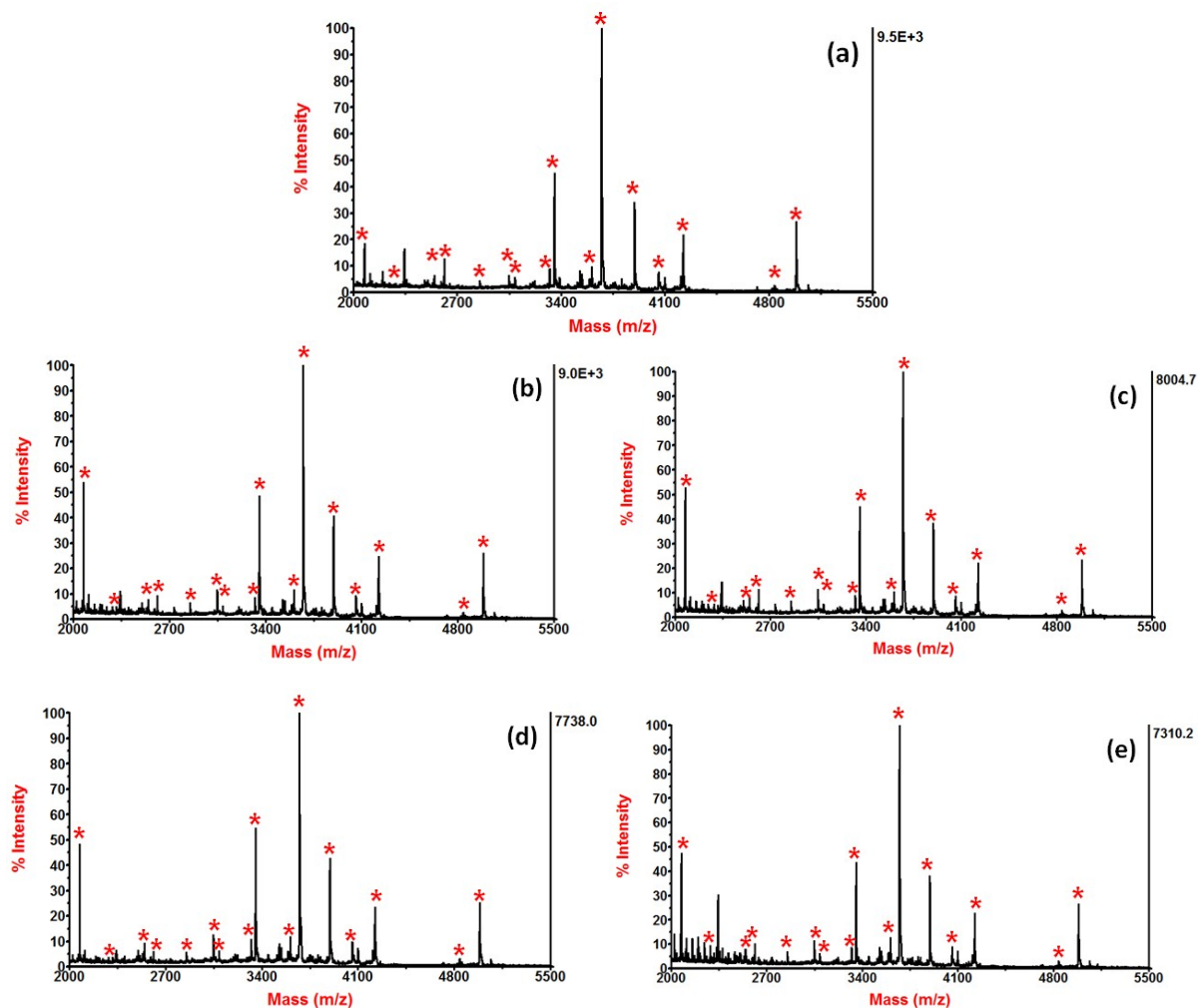


Fig. S10 MALDI-TOF-MS for the glycopeptides derived from HRP tryptic digest: (a) after treatment with MagG@COF-5 biocomposite used for the first time, (b) after enrichment with MagG@COF-5 recycled 2 times, (c) 3 times, (d) 4 times and (e) 5 times.

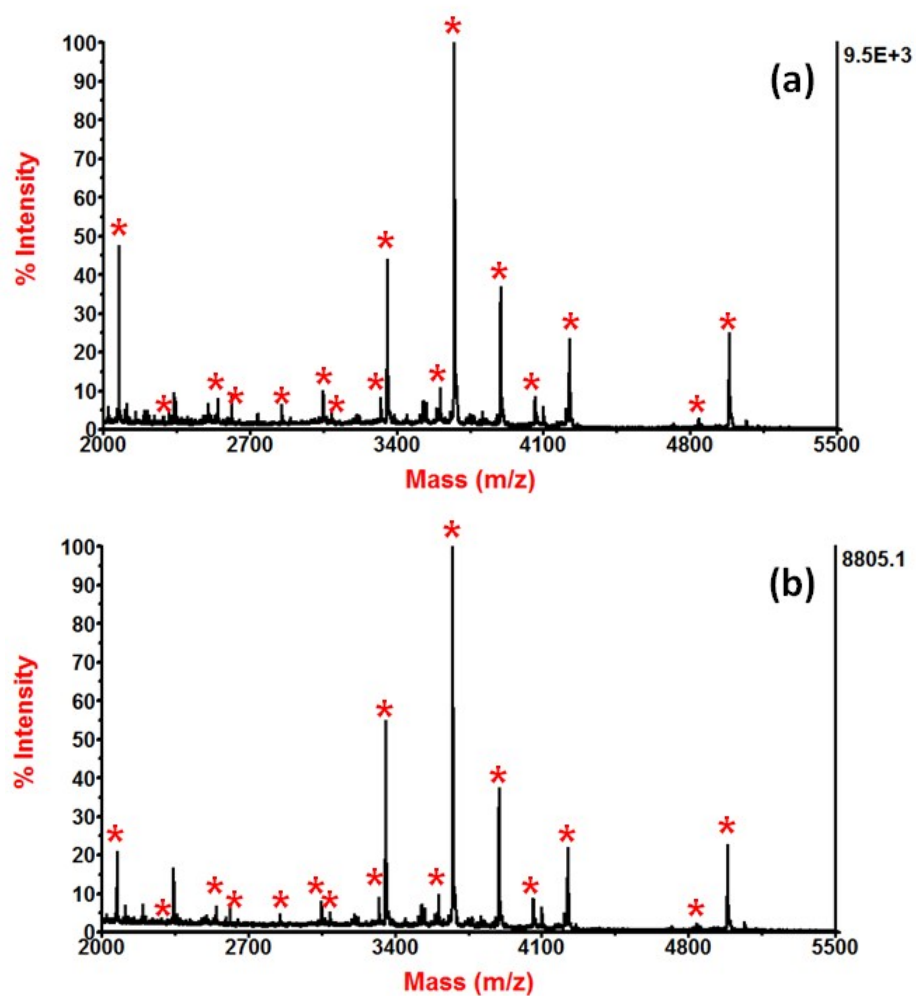


Fig. S11 MALDI-TOF-MS for the glycopeptides derived from HRP tryptic digest: (a) after treatment with the MagG@COF-5 biocomposite for the first time and (b) after enrichment with the MagG@COF-5 which had been placed for one month.

Table S1. Detailed information of the observed glycopeptides in HRP tryptic digest. **N#** denotes the N-linked glycosylation site.

No.	Observed m/z	Glycan composition	Amino acid sequence
1	2074	XylMan3GlcNAc2	PN#VSNIVR
2	2321	Man2GlcNAc2	MGN#ITPLGTQGQIR
3	2543	XylMan3FucGlcNAc2	SSPN#ATDTIPLVR
4	2612	XylMan3GlcNAc2	MGN#ITPLGTQGQIR
5	2850	FucGlcNAc	GLIQSDQELFSSPN#ATDTIPLVR
6	3048	XylMan2GlcNAc2	SFAN#STQTFNFAVEAMDR
7	3087	XylMan3FucGlcNAc2	GLCPLNGN#LSALVDFDLR
8	3323	XylMan3FucGlcNAc2	QLTPTFYDNSCPN#VSNIVR
9	3354	XylMan3FucGlcNAc2	SFAN#STQTFNFAVEAMDR
10	3606	XylMan3FucGlcNAc2	NQCRGLCPLNGN#LSALVDFDLR
11	3672	XylMan3FucGlcNAc2	GLIQSDQELFSSPN#ATDTIPLVR
12	3894	XylMan3FucGlcNAc2	LHFHDCFVNGCDASILDN#TTSFR
13	4057	XylMan3GlcNAc2	QLTPTFYDNSC(AAVESACPR)PN#VSNIVR-H2O
14	4222	XylMan3FucGlcNAc2	QLTPTFYDNSC(AAVESACPR)PN#VSNIVR
15	4839	XylMan3FucGlcNAc2, XylMan3GlcNAc2	LYN#FSNTGLPDPTLN#TTYLQTLR
16	4984	XylMan3FucGlcNAc2, XylMan3FucGlcNAc2	LYN#FSNTGLPDPTLN#TTYLQTLR

Table S2. Data of the identified N-linked glycopeptides enriched by MagG@COF-5 biocomposite. **N#** denotes the N-linked glycosylation site.

No.	Sequence	Protein Group Accessions	Charge	MH+ [Da]
1	GNEANYYS N# ATTDEHGLVQFS N# TTNVMGTS ^T TVR	P01023	3	3806.74729
2	LAGKPTHV N# VSVVMAEVDG Tc Y	P01876	3	2348.12802
3	STGKPTLY N# VSLVMSDTAG Tc Y	P04220	2	2366.09062
4	QQQHLFGS N# VTDeSGNFcLFR	P02787	3	2516.10770
5	ITYSIVQT N# cSK	P01042	2	1414.68926
6	NLFLNHSE N# ATAK	P00739	2	1459.71697
7	FEVDSPVY N# ATWSASLK	P04114	2	1914.91203
8	ELHHLQE N# VSNAFLDKGEFYIGSK	P00450	3	2904.41745
9	ELHHLQE N# VSNAFLDK	P00450	3	2022.98893
10	TKPREEQY N# STYR	P01857	4	1672.79238
11	TKPREEQ F# N#STFR	P01859	3	1640.80277
12	LGSFEGLV N# LTFIHLQHNR	P51884	3	2196.15781
13	KLPPGLLA N# FTLLR	P02750	2	1553.94109
14	NAHGEEK E# N#LTAR	Q06033	3	1469.69657
15	VGQLQLSH N# LSLVILVPQNLK	P05155	3	2314.34671
16	VYKPSAG N# nSLYR	P02749	3	1470.72086
17	TVLTPTA n# HMGN # VTFTIPANR	P01024	3	2257.13065
18	YPHKPEI N# STTHPGADLQENFcR	P00734	3	2712.24405
19	FSDGLES N# SSTQFEVK	P0C0L5	2	1775.79778
20	FSDGLES N# SSTQFEVKK	P0C0L5	3	1903.89292
21	VYKPSAG N# NSLYR	P02749	3	1469.73905
22	KLHINH N# N#LTESVGPLPK	P51884	4	2012.09072
23	VTQVYA E# N#GTVLQGSTVASVYK	P27169	2	2315.18022
24	LPPGLLA N# FTLLR	P02750	2	1425.85100
25	LLDLSG N# N#LTHLPK	P40197	3	1535.84305
26	NPPMG Gn# VVIFDTVITNQEEPQ N# HSGR	P02745	3	3115.46335
27	EHEAQS N# ASLDVFLGH Tn# VEELMK	P00736	4	2700.26601
28	SLPNFP N# TSATAN # ATGGR	Q6UXB8	2	1777.84221
29	QLAHQS N# ST n# IFFSPVSIATAFA m# LSLGTK	P01009	3	3199.61936
30	DIVEYY N# DS N# GSHVLQGR	P25311	3	2067.92447
31	GLTFQQ N# ASSMcVPDQD Ta IR	P01871	2	2356.05132
32	QVFPGL n# YcTSGAYS N# ASSTDSASY Yp L Tg DTR	P04114	3	3552.56797
33	SWPAVG N# cSSALR	P02790	2	1405.65459
34	ALPQPQ N# VTSLLG c# TH	P02790	2	1736.86467
35	GLTFQQ N# ASSMcVPDQD Ta IR	P01871	2	2340.06182
36	LSDLS I# N#STEcLHVHcR	P05156	4	2041.94497
37	LQNNEN N# IScVER	Q03591	2	1590.71868
38	FVEGSH N# STVSL Tt TK	P04114	2	1607.79314
39	QLAHQS N# ST n# IFFSPVSIATAFA m# LSLGTK	P01009	3	3182.62937
40	KKEDAL N# ETR	P10909	3	1204.61679
41	YAEDKF N# ETTEK	P43652	2	1475.65666
42	LHINH N# N#LTESVGPLPK	P51884	4	1884.00039
43	IYSGIL N# LSDITK	P03952	2	1437.78362

44	DKMFSQ N #DTR	P02750	2	1242.54021
45	DFYVDE N #TTVR	P29622	2	1359.60820
46	EHEAQ S #ASLDVFLGHTNVEELMK	P00736	4	2699.25893
47	QVHFFV N #ASDVDNVK	Q96IY4	3	1719.83219
48	LYLG S n N #LTALHPALFQ N #LSK	P22792	3	2317.21616
49	LQNN E n N #IScVER	Q03591	2	1591.70220
50	MVSH H n N #LTTGATLINEQWLLTTAKNLFL N #HSEnATAK	P00739	4	4122.05434
51	LYLG S nLTALHPALFQ N #LSK	P22792	3	2316.22519
52	ADGTVnQIEGEATPV N #LTEPAK	P05090	2	2256.09258
53	ADGTVnQIEGEATPV N #LTEPAKLEVK	P05090	3	2725.37949
54	MVSH H n N #LTTGATLInEQWLLTTAK	P00738	2	2681.38042
55	IIVPLnNRE N #ISDPTSPLR	P01591	3	2150.14945
56	GAFIS N #FSmTVDGK	P19823	2	1490.68535
57	MDGAS N #VTcInSR	P08603	2	1426.59453
58	WDPEV N #cSMAQIQLcPPPPQIPNSH N #MTTLNRYR	P08603	3	4011.81772
59	SLGNV N #FTVSAEALESQELcGTEVPSVPEHGR	P01023	3	3414.60349
60	SLGNV N #FTVSAEALESQELcGTEVPSVPEHGRK	P01023	4	3542.71401
61	LQAPL N #YTEFQKPlcLPSK	P03952	3	2248.17112
62	MVSH H n N #LTTGATLINEQWLLTTAK	P00739	3	2680.37802
63	VVLHP N #YSQVDIGLIK	P00738	2	1795.99626
64	VVLHP N #YSQVDIGLIKLK	P00738	4	2037.17568
65	MVSH H n N #LTTGATLINEQWLLTTAK	P00739	3	2680.37802
66	TPLT A n N #ITK	P01877	2	959.54119
67	SPDV I n N #GSPISQK	P08603	2	1342.68486
68	KEDAL N #ETR	P10909	2	1076.52178
69	GAFIS N #FSMTVDGK	P19823	2	1474.68901
70	LNAEN N #ATFYFK	P01042	2	1432.67497
71	HSHNN N #SSDLHPHK	P04196	3	1624.72062
72	DIEN F n N #STQK	P43652	2	1196.54350
73	ALGF E n N #ATQALGR	Q08380	2	1348.68584
74	LVP H M N #VSAVEK	Q96KN2	2	1324.69304
75	GNEAnYY S n N #ATTDEHGLVQFSInTTNVMGTSLTVR	P01023	4	3807.74624
76	NLFL N #HSE N #ATAK	P00739	2	1460.70220
77	EEQY N #STYRVVSVLTVLHQDWLnGK	P01857	3	2980.48056
78	EEQ F n N #STFRVVSVLTVVHQDWLnGK	P01859	4	2934.45913
79	EGY S n N #ISYIVVnHQGISSR	P49908	3	2125.03672
80	ISE N #ETTcYMGK	P08603	2	1562.63616
81	VIDF N #cTTSSVSSALAnTK	P04196	2	2016.94768
82	VIDF N #cTTSSVSSALANTK	P04196	2	2015.96050
83	AELS N #HTRPVILVPGcLGNQLEAK	P04180	4	2617.37490
84	EPGS N #VTMSVDAEcVPMVR	Q08380	2	2078.92827
85	EEQY N #STYR	P01857	2	1190.49687
86	EEQY N #STFR	P01860	2	1174.50139
87	EEQ F n N #STFR	P01859	2	1158.50517
88	EEQ F n N #STFRVVSVLTVVHQDWLnGK	P01859	3	2933.48636
89	EDAL N #ETR	P10909	2	948.42705
90	SLTF N #ETYQDISELVYGAK	P01008	2	2179.04741
91	LPTQ N #ITFQTESSVAEQEAEFQSPK	Q14624	3	2810.34079
92	GVNF N #VSK	P03952	2	865.44042

93	TMFPN#LTDVR	P06681	2	1194.58452
94	MFSQN#DTR	P02750	2	999.42046
95	DTFVN#ASR	P05155	2	910.42638
96	EGYSN#ISYIVVNHQGISSR	P49908	2	2124.03105
97	SLGnVN#FTVSAEALeSQCLeGTEVPSVPEHGR	P01023	3	3415.61826
98	VLSN#nSDANLELINTWVAK	P05155	3	2103.07999
99	VNQnLVYESGSLN#FSK	P04114	2	1800.86443
100	THTN#HSESHPN#ATFSAVGEASLeEDDWNsGER	P01871	4	3520.49917
101	YFYN#GTSMaCETfQYGGcMGnGNNFVTEK	P02760	3	3365.33200
102	YDFN#SSmLYSTAK	P04114	2	1543.66203
103	EWDN#TTTEcR	P20851	2	1312.51103
104	FLNN#GTcTAEgK	P05156	2	1312.58391
105	YFYN#GTSMaCETfQYGGcMnGnNFVTEK	P02760	3	3366.33396
106	YLGn#ATAIFFLPDEgK	P01009	2	1756.87981
107	YLGn#ATAIFFLPDEgKLQHLEnELTHDIITK	P01009	4	3541.83047
108	YTGn#ASALFILPDQDK	P01011	2	1753.86626
109	YTGn#ASALFILPDQDKMEeVEAMLLPETLKR	P01011	3	3523.77353
110	WVSN#KTEGR	P01008	2	1077.53240
111	AFEN#VTDLQWLILDHnLLENSK	P51884	3	2613.32279
112	DFVN#ASSK	P05546	2	868.40453
113	DFVN#ASSKYeITTIHnLFR	P05546	3	2256.12558
114	IYSN#HSALESALiPLQAPLK	P55058	3	2279.26535
115	VTQn#LTlIEESLTSEFIHDIDR	P36955	3	2574.29312
116	VLSN#nSDANLELINTWVAK	P05155	3	2102.07419
117	LQn#LTLPTN#ASIK	Q13201	2	1414.77971
118	FLn#GTcTAEgK	P05156	2	1313.56828
119	YKN#nSDISSTR	P01871	2	1286.58574
120	GLN#VTLsSTGRnGfK	P0C0L5	3	1552.79605
121	LGn#WSAmPScK	P02749	2	1267.54546
122	TLN#QSSDELQLSmGNAMFVK	P01011	2	2230.03887
123	VSN#VScQASVSR	P55058	2	1294.60747
124	LGn#WSAMPScK	P02749	2	1251.55022
125	VSN#QTLsLFFTVLQDVPVR	P01023	2	2164.16582
126	GLN#VTLsSTGR	P0C0L5	2	1105.58550
127	YKN#NSDISSTR	P01871	2	1285.60198
128	TLN#QSSDELQLSMGNAMFVK	P01011	2	2214.04375
129	LAN#LTQGEDQYYLR	P10909	2	1684.81902
130	GLN#LTEDTYKPR	Q08380	2	1407.71453
131	AVN#ITSENLIddVvSLIR	P02748	2	1972.05730
132	HAN#WTLTPLK	P27169	2	1181.63225
133	VVN#STTGpGEHLR	P07996	2	1367.69072
134	VLN#FTTK	P13671	2	823.45421
135	FLN#DTMAVYEAK	P29622	2	1402.65788
136	RNPPMGnVViFDTVITnQEEpYQn#HSGR	P02745	4	3270.56167
137	NPPMGnVViFDTVITnQEEpYQn#HSGR	P02745	3	3114.46579
138	AVLQLNEEGVDTAGSTGVTLN#LTSKPIILR	P08185	3	3110.68210
139	LnAENn#ATFYFK	P01042	2	1433.65911
140	FN#SSYLQGTnQITGR	P04114	2	1687.79399
141	HN#STGcLR	P10909	2	945.41954

142	QnQcFY N SSYLVNQVR	P19652	2	1922.83660
143	FN SSYLQGTNITGR	P04114	2	1686.80913
144	EN LTAPGSDSAVFQEQTTR	P00450	2	2127.98491
145	FN LTETSEAEIHQSFAQHLLR	P01011	3	2401.17954
146	AN LSSQALQMSLDYGFVTPLTMSIR	P19827	3	2831.40726
147	EN ISDPTSPLR	P01591	2	1229.60112
148	EN ETEIIK	P06276	2	976.48277
149	DN NSIITR	P06276	2	933.46587
150	DN YTDLVAIQNK	P14151	2	1394.68218
151	LVLSSSEKTVLTPATNHMG N VTFTIPANR	P01024	4	3012.57754
152	LSLHRPALEDLLGSE AN LTcTLTGRL	P01877	3	2964.58554
153	ALGISPFHEHAEEVFT AN DSGPR	P02766	4	2452.21230
154	LSHNELADSGIPGNS FN VSSLVELDLSYNK	P51884	3	3220.56736
155	GGETAQSDAPQWEQLnn KN LSMPLLPADFBK	P05546	4	3439.62563
156	ADGTVNQIEGEATPV N LTEPAKLEVK	P05090	3	2724.39713
157	ADGTVNQIEGEATPV N LTEPAK	P05090	2	2255.10430
158	DVQIIVFPEDGIHGF N FTR	P43251	3	2205.10160
159	FQSPAGTEALFELH N ISVADS AN YScvYVDLKPPFGGSAPSER	P04217	4	4630.18813
160	SVQEIQATFFYFTP N KTEDTIFLR	P02763	3	2896.44193
161	ADTHDEILEGLN FN LTEIPEAQIHEGFQELLR	P01009	4	3692.81020
162	LQAILGVPWKDK N cTSR	P01019	4	1987.04213
163	VNQNLVYESGSL N FSK	P04114	2	1799.88213
164	IYPGVDFGGEEL N VTFTVK	P03952	3	1984.98117
165	LGTSLSGGHVL MN GTCLK	P80108	3	1715.89780
166	TVLTPATNHMG N VTFTIPANR	P01024	3	2256.14518
167	FNPGAESVVL SN STLTK	Q13201	2	1663.85381
168	SVVAPATDGGL N LTSTFLR	P41222	2	1920.00615
169	AALAAFNAQN N GSNFQLEEISR	P02765	2	2366.13994
170	QSVPAHFVAL N GSK	P06681	2	1455.75810
171	LDAPTNLQFV N ETDSTVLVR	P02751	3	2233.13907
172	SLDFNTLVD N ISVDPETGDLWVGcHPnGMK	P27169	3	3332.53232
173	SRYPHKPEI N STTHPGADLQENFcR	P00734	4	2955.37246
174	EHEGAIYPD N TTDFQR	P00450	2	1893.82207
175	TVIRPFYLT N SSGVD	Q08380	2	1669.84490
176	DIVEYYNDS N GSHVLQGR	P25311	3	2066.93991
177	VVAEGFDFAnGINISPDGK	P27169	2	1950.94780
178	SQILEGLGF N LTELSSESDVHR	P29622	3	2345.16520
179	IIVPLNNRE N ISDPTSPLR	P01591	2	2149.15923
180	AQLLQGLGF N LTER	P08185	2	1560.83916
181	nGTGHG N STHHGPEYmR	P02790	2	1869.75981
182	nGTGHG N STHHGPEYMR	P02790	2	1853.76348
183	nLFL N HSE N ATAK	P00739	2	1461.68584
184	nN ATVHEQVGGPSLTSDLQAQSK	P04004	3	2383.13816
185	N TTcQDLQIEVTVK	P0C0L5	2	1649.80498
186	N HScSEGQISIFR	O75882	2	1535.69121
187	N GTGHGNSTHHGPEYMR	P02790	4	1852.77736
188	N NSDISSTR	P01871	2	994.44310
189	N YTLTGR	P10643	2	825.41008
190	N LSMPLLPADFBK	P05546	3	1483.76236

191	RNPPmGGNVVIFDVTITNQEOPYQ#HSGR	P02745	3	3286.55344
192	NPPmGGNVVIFDVTITNQEOPYQ#HSGR	P02745	3	3130.46299
193	TVLTPATNHmGN#VTFTIPANR	P01024	3	2272.14054
194	mVSHH#LTTGATLINEQWLLTTAKnLFLnHSEnATAK	P00739	4	4139.06264
195	mVSHH#LTTGATLINEQWLLTTAK	P00738	4	2696.37441
196	mL#TSSLLEQLNEQFNWVSR	P10909	2	2426.17241
197	VYLQGLIDcYLFGN#SSTVLEDSK	P35542	3	2622.27024
198	LKELPGVc#ETMMALWEEcKPcLK	P10909	3	2937.36490
199	WVLTAAHcLLYPPWDK#FTENDLLVR	P00734	4	3172.60244
200	TELFSSScPGGIML#ETGQGYQR	O95445	2	2533.14018
201	DTAVFEcLPQHAmFG#DTITcTTHGnWTKLPecR	P02749	4	4009.75918
202	DTAVFEcLPQHAmFG#DTITcTTHGnWTK	P02749	4	3354.45376
203	DTAVFEcLPQHAmFG#DTITcTTHGnWTK	P02749	3	3370.45603
204	DAGVvcT#ETR	Q08380	2	1222.53752
205	ELPGVc#ETMMALWEEcKPcLK	P10909	3	2696.19449
206	LTDITcGVGN#MSAN#ASDQER	P06681	2	2140.91362
207	LTDITcGVGN#mSAnASDQER	P06681	2	2156.90996
208	AATcINPL#GSVcERPAN#HSAK	O75882	3	2369.09653
209	QDQcIYN#TTYLVQREnGTISR	P02763	3	2675.23862
210	QDQcIYN#TTYLVQR	P02763	2	1917.88152
211	QNQcFY#SSYLVQR	P19652	2	1921.85088
212	LGAc#DTLQQLMEVFK	P01008	2	1867.89372
213	LGAc#DTLQQLMEVFKDITSEK	P01008	3	2688.29697
214	DKIcDLLVANNHFAHFFAPQ#LTnMNK	P19827	4	3174.53896
215	DKIcDLLVANNHFAHFFAPQ#LTnMNK	P19827	4	3173.53042
216	LGHePDPVLVNGEFSSSGPV#VSDK	P20851	3	2612.21994
217	LGHePDPVLVnGEFSSSGPV#VSDK	P20851	3	2613.22062
218	VTAcHSSQP#ATLYK	P05543	3	1677.79428
219	KVScPIMPeS#ATVPDGEccPR	P07996	3	2536.07645
220	GIc#SSDVR	O75882	2	1008.44145
221	IPcSQPPQIEHGTIN#SSR	P08603	2	2021.97270
222	KVcQDcPLLAPL#DTR	P02765	2	1900.92681
223	GcVLLSYL#ETVTVSASLESVR	P01023	3	2398.21653
224	FcRD#YTDLVAIQNK	P14151	3	1857.88144
225	IcDLLVANNHFAHFFAPQ#LTnMNK	P19827	3	2930.41312
226	VcQDcPLLAPL#DTR	P02765	2	1772.83183
227	FSLLGHASIScTVEN#ETIGVWRPSPPTcEK	P04003	4	3373.61855
228	LEPVHLQLQcMSQEQLAQVAA#ATK	Q96PD5	3	2808.40537
229	cIQAN#YSLMEEnGK	P05090	2	1529.66448
230	cIQAN#YSLMEEnGKIK	P05090	3	1770.83768
231	cIQAN#YSLmENGK	P05090	2	1544.67436
232	cSDGWSFDATTLDD#GTMLFFK	P02790	2	2529.06328