

Supplementary Information

DNA tetrahedron-inspired magnetic hybrid with regular distribution of trypsin for ultra-fast digestion of proteins and glycoproteomics analysis

Jiahui Lai, Danni Wang, Xiuqin Sheng, Chuan-Fan Ding, Yinghua Yan**

*School of Materials Science and Chemical Engineering, Ningbo
University, Ningbo 315211, China*

*Corresponding author: Chuan-Fan Ding, Yinghua Yan

E-mail: dingchuanfan@nbu.edu.cn; yanyinghua@nbu.edu.cn °

Characterization

Deionized water was got through using Milli-Q IQ 7000. Scanning electron microscopy (SEM) was characterized by ZEISS Sigma 300 (Germany). Transmission electron microscopy (TEM) was characterized by FEI Tecnai F20 (USA). Fourier transform infrared spectra (FT-IR) was characterized by Thermo Fisher Scientific 10 infrared spectrometer analysis. Thermogravimetry Analysis was characterized by TA TGA 550 (USA). X-ray photoelectron spectroscopy (XPS) was characterized by Thermo Kalpha. AFM was characterized by Agilent 5500.

Standard protein pretreatment

According to the classic method, BSA was pretreated with DTT and IAA. BSA (50 mg) dispersed in NH_4HCO_3 (50 mM, 2.5 mL) was boiled and denatured at 100 °C for 10 min. Then a mixed solution of DTT (5 mM, 0.25 mL) and NH_4HCO_3 (50 mM, 2.25 mL) was added to it and shook at 56 °C for 1 h. Afterwards, a mixed solution of IAA (14 mM, 0.5 mL) and NH_4HCO_3 (50 mM, 4.5 mL) was added to it under dark conditions and alkylated at 37 °C for 1 h.

HRP was pretreated with DTT and IAA. HRP (50 mg) was dispersed to NH_4HCO_3 (50 mM, 2.5 mL), and then heating it at 100 °C for 10 min to denature it. Then a mixed solution of DTT (5 mM, 0.25 mL) and NH_4HCO_3 (50 mM, 2.25 mL) was added to it and shook at 56 °C for 1 h. Afterwards, a mixed solution of IAA (14 mM, 0.5 mL) and NH_4HCO_3 (50 mM, 4.5 mL) was added to it under dark conditions and alkylated at 37 °C for 1 h.

Practical sample pretreatment

The protein sample (0.4 mg) was completely dissolved in NH_4HCO_3 (50 mM, 20 μL) solution and heated at 100 °C for 10 min to denature it. Subsequently, a mixed solution of DTT (5 mM, 2 μL) and NH_4HCO_3 (50 mM, 18 μL) was added to it and shook at 56 °C for 1 h. Afterwards, a mixed

solution of IAA (14 mM, 4 μ L) and NH_4HCO_3 (50 mM, 36 μ L) was added to it under dark conditions and alkylated at 37 °C for 1 h.

MALDI-TOF MS analysis

All studies were carried out in reflected positive ion mode using MALDI-TOF MS (Bruker Auto Flex Max). An enhanced Nd:YAG laser (the detection frequency was 1000 Hz) operated the device. The mass charge ratio measurement range was 1000-3500 Da, the detector voltage was around 18 kV, and the launch tube acceleration voltage was 20 kV. The matrix used was DHB (20 mg/mL), dissolved in a mixed solution (ACN: H_2O : TFA = 20%: 79.9%: 0.1%, volume ratio).

Nano LC-MS/MS analysis.

Liquid phase A was a 0.1% formic acid aqueous solution, and B was a 0.1% formic acid acetonitrile solution (acetonitrile is 100%). Column Thermo Scientific analytical column (75 μ m $\times$ 25 cm, 5 μ m, 100 Å, C18) was equilibrated with 95% A liquid. The sample was loaded by the autosampler to the Thermo Scientific EASY trap column (100 μ m $\times$ 2cm, 5 μ m, 100Å, C18), and then separated by the chromatographic column. The relevant liquid gradient was as follows: from 0 min to 40 min, linear gradient of liquid B was from 5% to 28%; from 40 minutes to 42 minutes, the linear gradient of liquid B was from 28% to 90%; from 42 minutes to 60 minutes, the liquid B was maintained at 90%. The enzymatic hydrolysis products were desalted and separated by capillary high performance liquid chromatography, and then analyzed by mass spectrometry using Orbitrap Fusion Lumos mass spectrometer (Thermo Finnigan, San Jose, CA). Analysis time: 60min, detection method: positive ion, precursor ion scan range: 375-1800 m/z, primary mass spectrometry resolution: 120,000 at m/z 200, scan mode: Top-Speed, AGC target: 4e5, first-level Maximum IT: 50 ms, Number of scan ranges: 1, Dynamic exclusion: 40.0 s, Data Dependent Mode: Cycle Time,

Time between Master Scan: 3s. The mass-to-charge ratios of peptides and peptide fragments were collected according to the following method: the secondary spectrum scan (MS2 scan) was performed at the same time as each primary scan (Master scan), the scanning cycle was 3s, MS2 Activation Type: HCD, secondary mass spectrometry Resolution: 50,000 at m/z 200, Microscans: 1, Level 2 Maximum IT: 105 ms, AGC target: 1e5.

Database search and data analysis.

RAW files were utilized for MS analysis, library detection and quantification using the Proteome Discoverer 2.4 software (Thermo Scientific). The human database downloaded from the website "<http://www.uniprot.org>" was used for comparison. The library search involved uploading RAW files to SequestHT through software and selecting and searching the established databases. The following were the search parameters: peptides with charged numbers of 2⁺, 3⁺, and 4⁺; a maximum of 2 allowed missing cut sites; trypsin digestion; and monoisotopic mass. Additionally, fixed modifications to carbamidomethylation (C), dynamic modifications to oxidation (M), acethyl (protein N-term), deamidated [N], and phospho (S, T, Y) were considered. The maximum error was 0.05 Da for precursor ions and 10 ppm for precursor ions. Proteome Discoverer 2.4 outputs the results after high-confidence peptide screening based on the peptide identification results.

Preparation of Fe₃O₄

Magnetic Fe₃O₄ were synthesized through a solvothermal reduction method. Firstly, FeCl₃·6H₂O (2.16 g), diethylene glycol (60 mL), and ethylene glycol (20 mL) were added to a beaker (100 mL) and sonicated for 10 min until homogeneous, then NaAc (6.0 g) and sodium acrylates (6.0 g) were added, and mechanically stirred until this mixed solution was a homogeneous dark yellow solution, and then the mixed solution was transferred to a reactor (100 mL) and heated at 200 °C for 10 h. After the reaction, the mixed solution was cooled to room temperature, the product was separated with a magnet, washed three times with anhydrous ethanol, and the washed product was dried under vacuum. The Fe₃O₄ was finally obtained.

Preparation of poly (Fe₃O₄@DOPA/PEI)

Firstly, Fe₃O₄ (40 mg), Tris-HCl (10 mM, 160 mL), and L-DOPA (160 mg) were added to a beaker (500 mL) and sonicated for 10 min until homogeneous, then PEI (2 mL), CuSO₄ (212 mg), and H₂O₂ (1.6 mL) were added sequentially, sonicated to a homogeneous dark brown solution, and the mixed solution was subsequently mechanically stirred for 12 h at room temperature. After the reaction, the product was separated with a magnet, washed three times with deionized water, and the washed product was dried under vacuum. The poly (Fe₃O₄@DOPA/PEI) was finally obtained.

Preparation of AuNPs solution

HAuCl₄·4H₂O (20 mg·mL⁻¹, 2.5 mL) was added to deionized water (122 mL) for condensation reflux with continuous heating. Waiting for the heating to 181 °C, sodium citrate dihydrate (38.8 mM, 12.5 mL) was quickly added, and the color of the solution rapidly changed from light yellow to dark purple. After continuous heating at 181 °C for 15 min, the solution was cooled to room temperature and set aside. The AuNPs solution was finally obtained.

Preparation of Fe₃O₄@DOPA/PEI@Au

Firstly, poly (Fe₃O₄@DOPA/PEI) (90 mg) was added to AuNPs solution (60 mL) and sonicated for 30 min until homogeneous. Subsequently, the mixed solution was mechanically stirred at room temperature for 1 h. After the reaction, the product was separated with a magnet, washed three times with anhydrous ethanol, and the washed product was dried under vacuum. The Fe₃O₄@DOPA/PEI@Au was finally obtained.

Preparation of DNA tetrahedron (DNA TET)

Through the principle of base complementary pairing, four ssDNA were carefully designed, which were used to prepare DNA TET. First, each ssDNA was dissolved and diluted to a concentration of 100 $\mu\text{mol}\cdot\text{L}^{-1}$. Then, each ssDNA solution (1 μL) was added to TE buffer (96 μL). The final concentration of each ssDNA solution was 1 $\mu\text{mol}\cdot\text{L}^{-1}$. The above solutions were reacted at 95 °C for 10 min and then at 4 °C for 30 min. The product (DNA TET) was finally obtained.

Optimization of Trypsin amount in preparing Fe₃O₄@DOPA/PEI@Au@DNA TET-Trypsin

We optimized the amount of trypsin to achieve the maximum loading of trypsin on the precursor material, thereby improving the digestion and enrichment efficiency of Fe₃O₄@DOPA/PEI@Au@DNA TET-Trypsin. We evaluated the effects of trypsin solutions of different concentrations for digestion and enrichment performance of material. Firstly, we added trypsin solutions of different concentrations (0.5, 1.0, 1.5, 2.0 $\text{mg}\cdot\text{mL}^{-1}$, 0.1 mL) to the prepared Fe₃O₄@DOPA/PEI@Au@DNA TET and incubated overnight at 4 °C. Then, we washed the obtained mixed solution with PBS and freeze-dried to obtain the target product. Finally, we enriched the glycopeptides in the standard protein utilizing these target products obtained under different experimental conditions, and analyzed these eluates utilizing MALDI-TOF-MS. The spectra is

shown in Figure S9. By comparing the information provided by the MALDI-TOF mass spectra, we can see that when the concentration of trypsin solution is greater than $1.0 \text{ mg} \cdot \text{mL}^{-1}$, the enrichment performance of $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA}$ TET-Trypsin on glycopeptides does not change significantly.

Study on the loading capacity of trypsin on the $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA}$ TET

We quantified the loading capacity of trypsin on $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA}$ TET using the BCA protein assay kit. Firstly, we plotted a standard curve ($y=1.35783x+0.0075$, $R^2=0.99649$) by measuring the standard protein. Subsequently, we determined that the average absorbance value of the supernatant after the reaction between trypsin solution ($1.0 \text{ mg} \cdot \text{mL}^{-1}$, 0.1 mL) and $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA}$ TET was 1.2974. Based on the standard curve, we can calculate that the protein concentration of the supernatant at this time is $0.95 \text{ mg} \cdot \text{mL}^{-1}$. Therefore, after reacting trypsin solution ($1.0 \text{ mg} \cdot \text{mL}^{-1}$, 0.1 mL) with $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA}$ TET, approximately 0.005 mg of trypsin was successfully loaded onto precursor material.

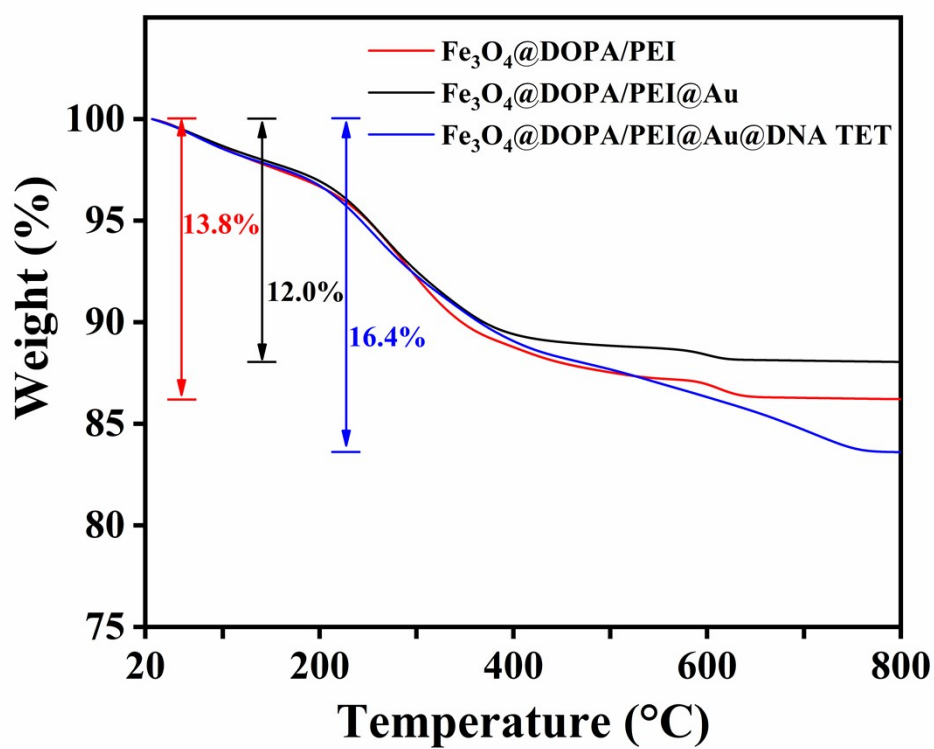


FIGURE S1. The TGA curves of $\text{Fe}_3\text{O}_4@\text{DOPA/PEI}$, $\text{Fe}_3\text{O}_4@\text{DOPA/PEI@Au}$, and $\text{Fe}_3\text{O}_4@\text{DOPA/PEI@Au@DNA TET}$.

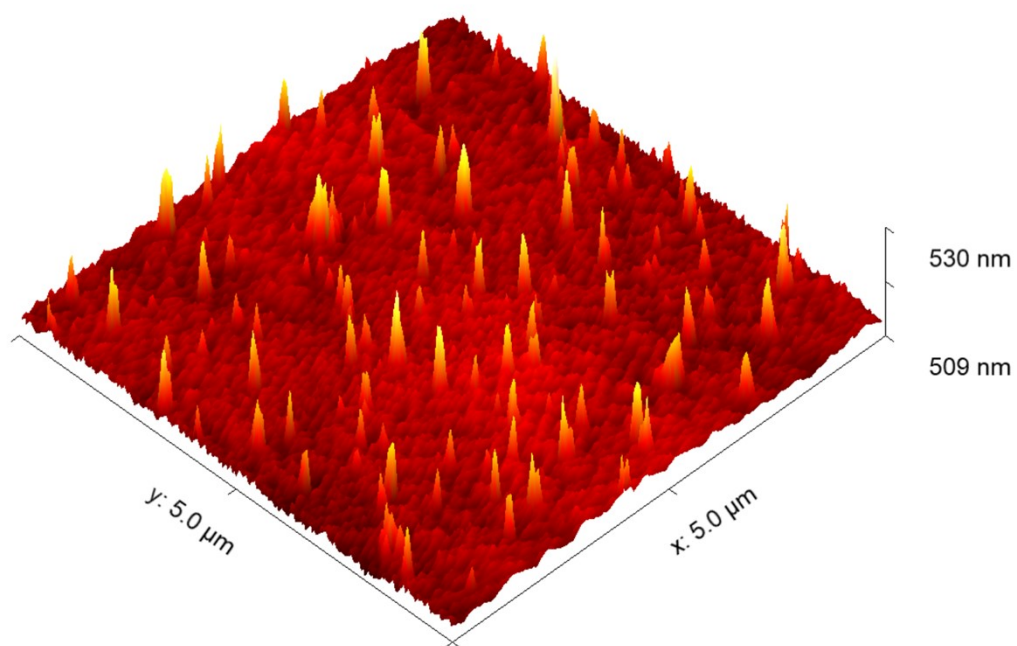


FIGURE S2. The AFM image of DNA tetrahedron (DNA TET).

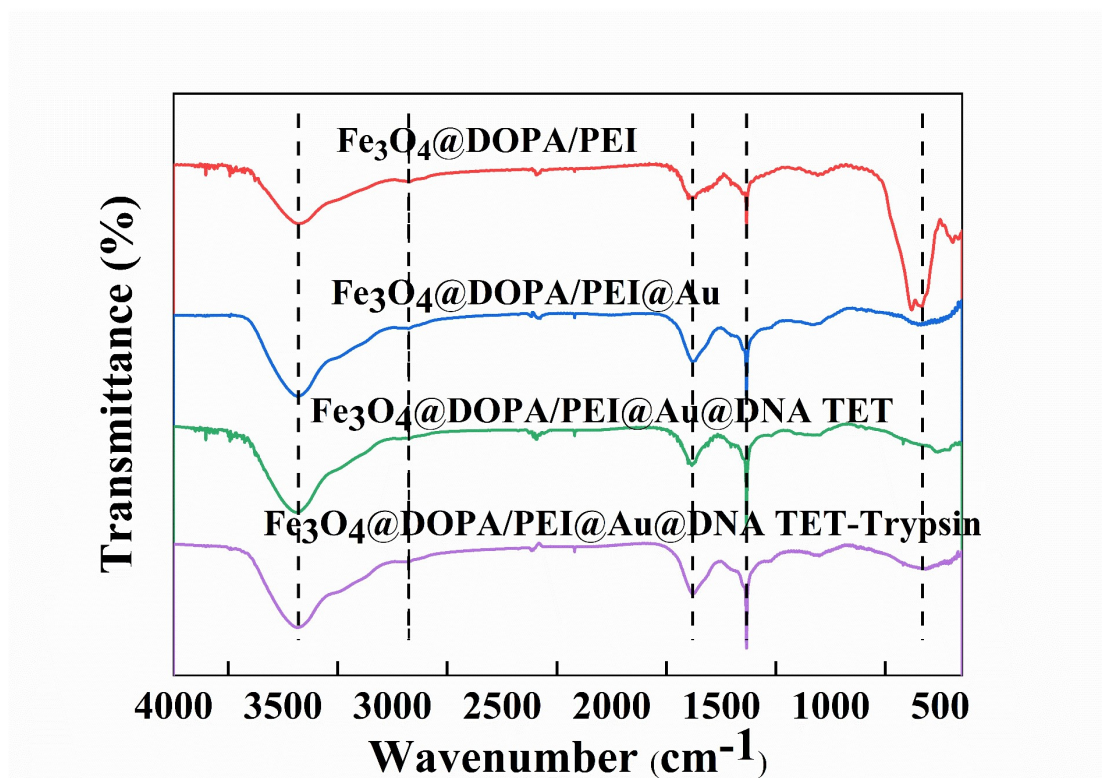


FIGURE S3. The FT-IR spectra of $\text{Fe}_3\text{O}_4@\text{DOPA/PEI}$, $\text{Fe}_3\text{O}_4@\text{DOPA/PEI@Au}$, $\text{Fe}_3\text{O}_4@\text{DOPA/PEI@Au@DNA TET}$, and $\text{Fe}_3\text{O}_4@\text{DOPA/PEI@Au@DNA TET-Trypsin}$.

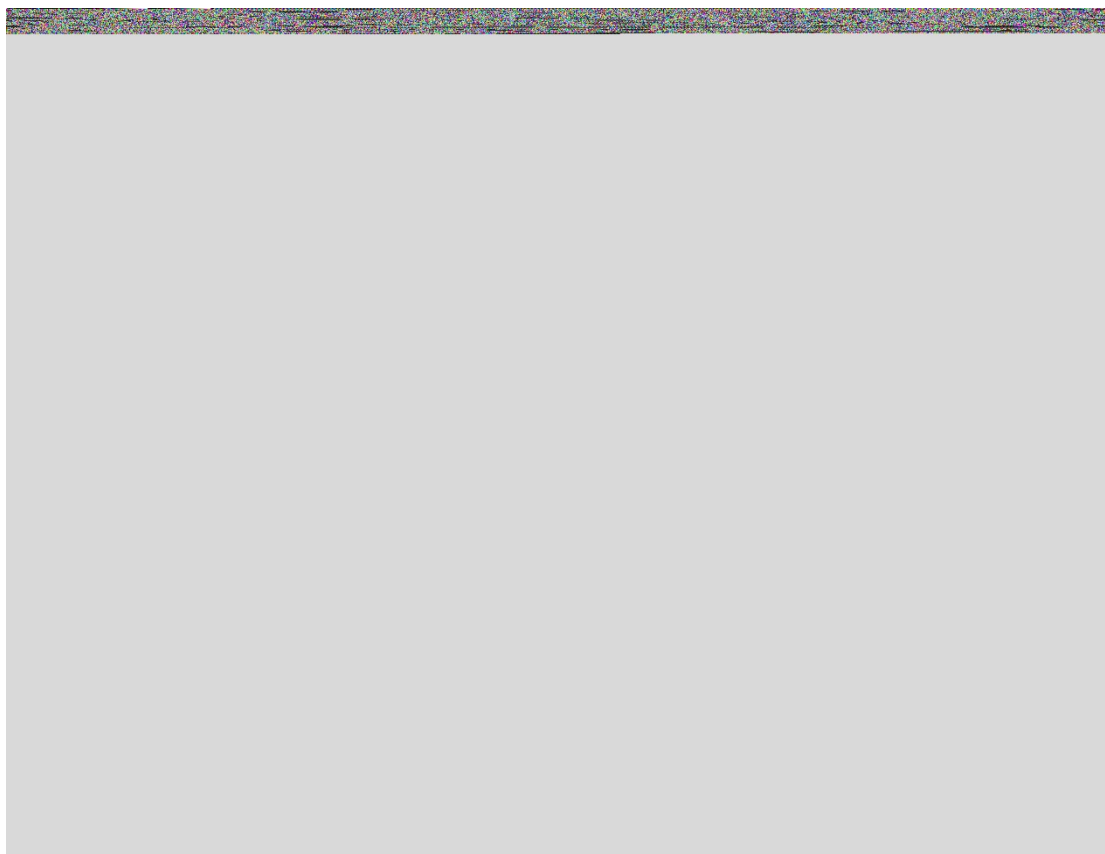


FIGURE S4. The XPS spectra of Fe₃O₄@DOPA/PEI@Au@DNA TET-Trypsin.

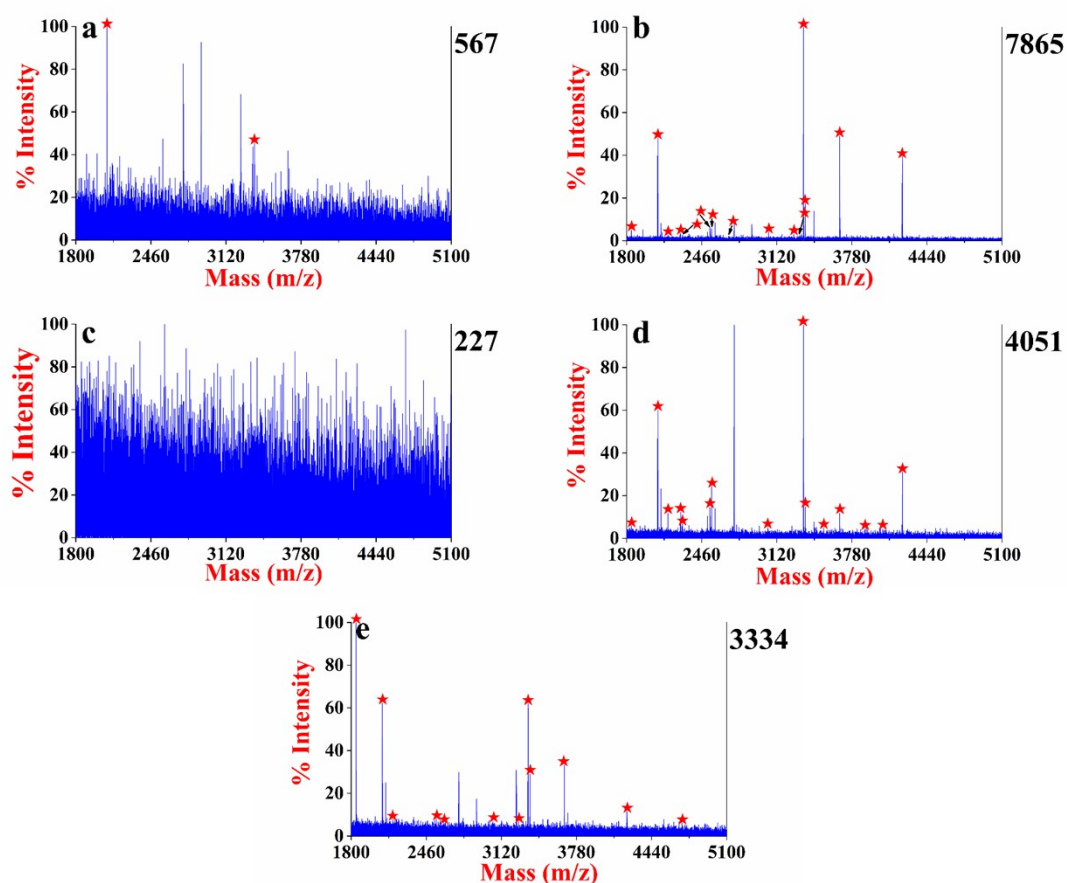


FIGURE S5. The MALDI-TOF mass spectra of glycopeptides from the HRP pretreatment solution after digested and enriched by $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA}$ TET-Trypsin prepared under different conditions. (a)EDC: NHS=3:1; (b)EDC: NHS =2:1; (c)EDC: NHS =1:1; (d)EDC: NHS =1:2; (e)EDC: NHS =1:3. Glycopeptide peaks are marked with "★".

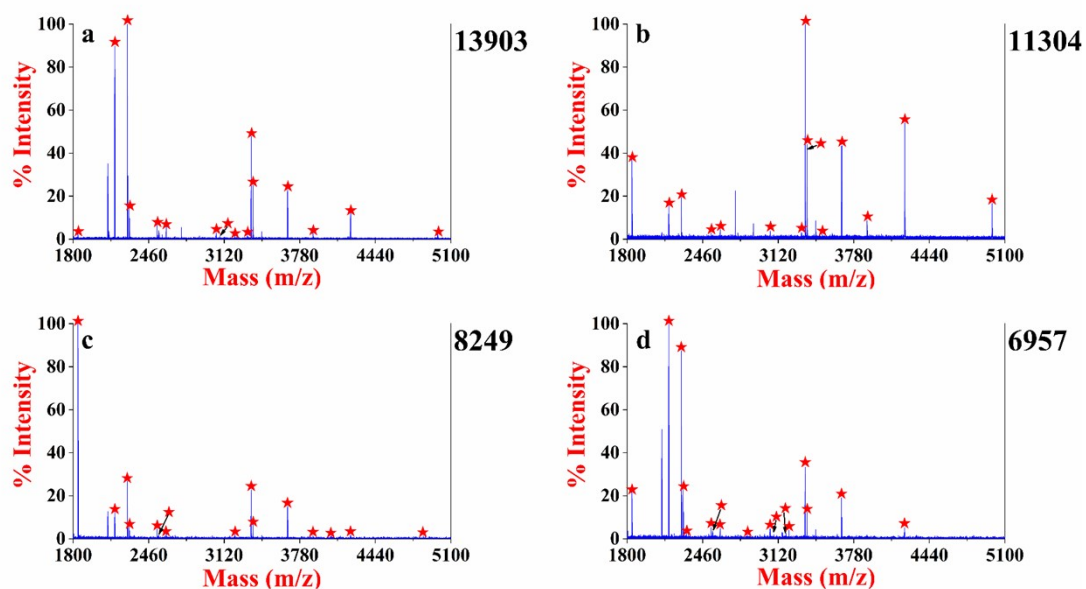


FIGURE S6. The MALDI-TOF mass spectra of glycopeptides from the HRP pretreatment solution after digested and enriched by $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA}$ TET-Trypsin at different elution times. (a)1 h; (b)30min; (c)10min; (d)5min. Glycopeptide peaks are marked with "★".

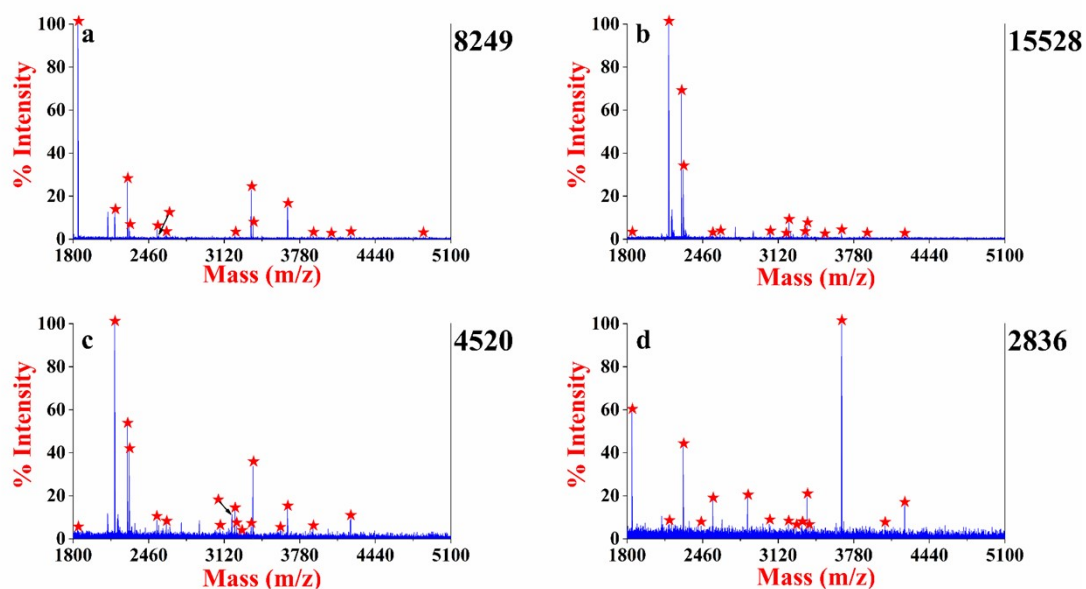


FIGURE S7. The MALDI-TOF mass spectra of glycopeptides from the HRP pretreatment solution after digested and enriched by $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA}$ TET-Trypsin at different enrichment times. (a)1 h; (b)30min; (c)10min; (d)5min. Glycopeptide peaks are marked with "★".

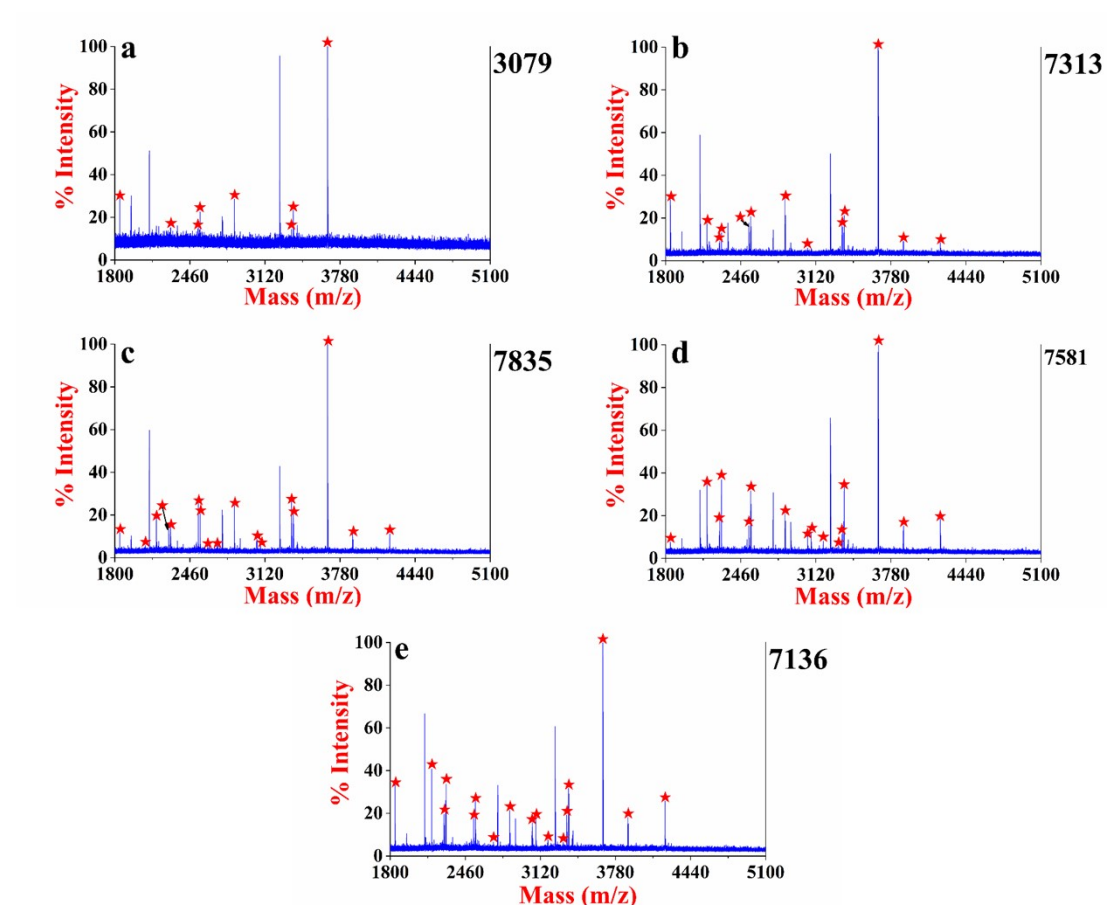


FIGURE S8. The MALDI-TOF mass spectra of glycopeptides from the HRP pretreatment solution after digested and enriched by $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA}$ TET-Trypsin with different amounts. (a) 0.1 mg; (b) 0.5 mg; (c) 1.0 mg; (d) 1.5 mg; (e) 2.0 mg. Glycopeptide peaks are marked with "★".

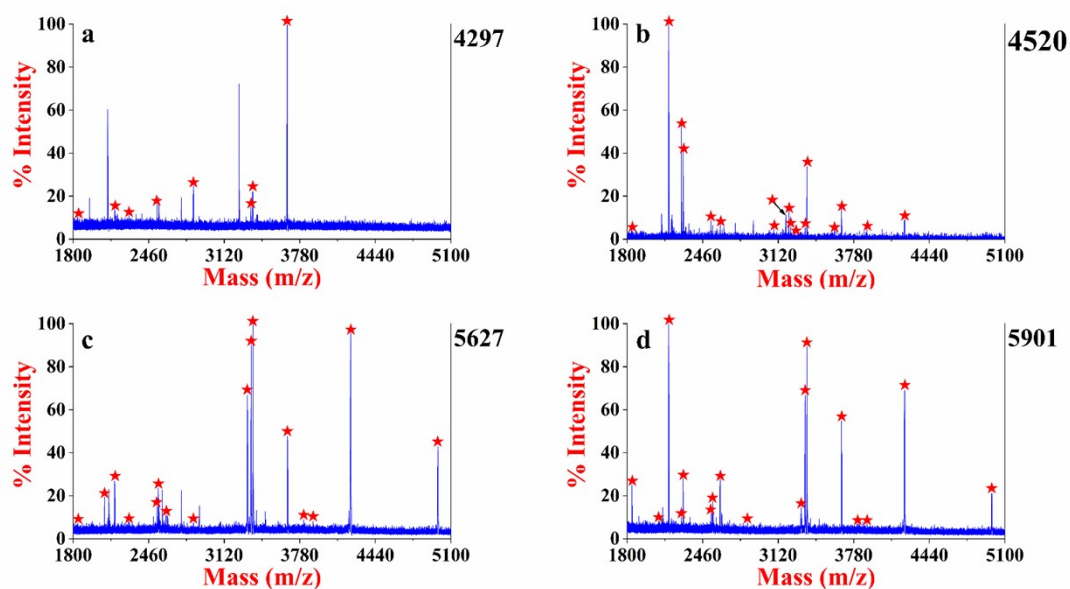


FIGURE S9. The MALDI-TOF mass spectra of glycopeptides from the HRP pretreatment solution after digested and enriched by $\text{Fe}_3\text{O}_4@\text{DOPA}/\text{PEI}@\text{Au}@\text{DNA TET}$ -Trypsin prepared under different concentrations of trypsin solution. (a) $0.5 \text{ mg} \cdot \text{mL}^{-1}$; (b) $1.0 \text{ mg} \cdot \text{mL}^{-1}$; (c) $1.5 \text{ mg} \cdot \text{mL}^{-1}$; (d) $2.0 \text{ mg} \cdot \text{mL}^{-1}$. Glycopeptide peaks are marked with "★".

Table S1. The detailed information of glycopeptides derived from HRP after digestion and enrichment by Fe₃O₄@DOPA/PEI@Au@DNA TET-Trypsin.

Number	m/z	Amino acid sequence	Glycan composition
1	1842.6	NVGLN#R	XylMan3FucGlcNAc2
2	2164.5	LYN#FSN#TGLP	Man3GlcNAc2Fuc
3	2276.2	SILLDN#TTSFR	XylMan2FucGlcNAc2
4	2291.2	SILLDN#TTSFR	XylMan2GlcNAc2
5	2533.6	SFAN#STQTFNFAFVEAMDR	FucGlcNAc
6	2545.8	SSPN#ATDTIPLVR	XylMan3GlcNAc2Fuc
7	2612.2	MGN#ITPLTGTQGQIR	XylMan3GlcNAc2
8	2702.0	GLIQSDQELFSSPN#ATDTIPLVR	GlcNAc
9	2853.7	GLIQSDQELFSSPN#ATDTIPLVR	FucGlcNAc
10	3049.7	SFAN#STQTFNFAFVEAMDR	XylMan2GlcNAc2
11	3085.5	GLCPLNGN#LSALVDFDLR	XylMan3FucGlcNAc2
12	3211.3	SFAN#STQTFNFAFVEAMDR	XylMan3GlcNAc2
13	3323.8	QLTPTFYDNSCPN#VSNIVR	XylMan3FucGlcNAc2
14	3355.2	SFAN#STQTFNFAFVEAMDR	XylMan3FucGlcNAc2
15	3369.7	SFAN#STQTFNFAFVEAM*DR	XylMan3FucGlcNAc2
16	3673.1	GLIQSDQELFSSPN#ATDTIPLVR	NQCRGLCPLNGN#LSALVDFDLR
17	3896.8	LHFHDCFVNGCDASILLDN#TTSFR	XylMan3FucGlcNAc2
18	4224.6	QLTPTFYDNSC(AAVESACPR)PN#VSNIVR	XylMan3FucGlcNAc2
19	4985.9	LYN#FSNTGLPDPTLN#TTYLQTLR	XylMan3FucGlcNAc2, XylMan3FucGlcNAc2

Table S2. The details of glycopeptides in normal control serum after digestion and enrichment by Fe₃O₄@DOPA/PEI@Au@DNA TET-Trypsin.

Number	Annotated Sequence	Modifications	Master protein	MH ⁺ [Da]
1	[R].TLNQSSDELQLS MGNAMFVKEQLSL LDRFTEDAK.[R]	1xDeamidated [N3]	P01011	3859.87284
2	[K].LQDRGPDVLTA TVSGKLPTQNITFQT ESSVAEQEAEFQSP K.[Y]	1xDeamidated [N21]	Q14624	4448.21036
3	[K].LATALSLSNKFV EGSHNSTVSLTTK.[N]	1xDeamidated [N17]	P04114	2606.36719
4	[R].FSLLGHASISCT VENETIGVWRPSP TCEK.[I]	2xCarbamidome thyl [C11; C28]; 1xDeamidated [N15]	P04003	3373.61914
5	[K].SLGNVNFTVSA EALESQELCGTEVP SVPEHGRK.[D]	1xCarbamidome thyl [C20]; 1xDeamidated [N6]	P01023	3542.70677
6	[K].MVSHHNLTTGA TLNEQWLLTTAK.[N]	1xDeamidated [N6]	P00739; P00738	2680.37631
7	[R].EEQFNSTFRVVS VLTVVHQDWLNGK .[E]	1xDeamidated [N5]	P01859	2933.4792
8	[R].EEQFNSTFRVVS VLTVVHQDWLNGK EYK.[C]	1xDeamidated [N5]	P01859	3353.68008
9	[K].AGLQAFFQVQE CNKSSSKDNIR.[G]	1xCarbamidome thyl [C12]; 1xDeamidated [N13]	P00450	2528.21981
10	[K].KQLVEIEKVVL HPNYSQVDIGLIK.[L]	1xDeamidated [N14]	P00738	2763.56549
11	[R].RNPPMGGNVVI FDTVITNQEEPYN HSGR.[F]	1xDeamidated [N25]	P02745	3270.55965
12	[R].IADAHLDRVEN TTVYYLVLDVQESD CSVLSR.[K]	1xCarbamidome thyl [C26]; 1xDeamidated	P04196	3581.74282

		[N11]		
13	[R].QQQHLLFGSNVT DCSGNFCLFRSETK. [D]	2xCarbamidome thyl [C13; C18]; 1xDeamidated [N9]	P02787	2961.32542
14	[R].DIPTNSPELEETL THTITKLNAENNAT FYFKIDNVK.[K]	1xDeamidated [N25]	P01042	4122.05536
15	[K].ADTHDEILEGLN FNLTEIPEAQIHEGF QELLR.[T]	1xDeamidated [N14]	P01009	3692.80786
16	[K].IITILEEEMNVSV CGLYTYGKPVPGH VTVSICR.[K]	2xCarbamidome thyl [C14; C32]; 1xDeamidated [N10]	P01023	3735.87945
17	[K].YLGNATAIFFLP DEGKLQHLENELTH DIITK.[F]	1xDeamidated [N4]	P01009	3541.82133
18	[R].RNPPMGGNVVI FDTVITNQEEPYN HSGR.[F]	1xDeamidated [N25]; 1xOxidation [M5]	P02745	3286.55456
19	[K].SKWNITMESYV VHTNYDEYAIFLT K.[K]	1xDeamidated [N4]	P02760	3053.4601
20	[K].TKPREEQFNSTF RVVSVLTVVHQDW LNGK.[E]	2xDeamidated [N9; N27]	P01859	3416.75973
21	[K].QLVEIEKVVLHP NYSQVDIGLIK.[L]	1xDeamidated [N13]	P00738	2635.47053
22	[K].MVSHHNLTTGA TLNQWLLTTAK. [N]	1xDeamidated [N6]; 1xOxidation [M1]	P00739; P00738	2696.37123
23	[K].TKPREEQFNSTF RVVSVLTVVHQDW LNGKEYK.[C]	1xDeamidated [N9]	P01859	3835.9766
24	[K].GCVLLSYLNET VTVSASLESVRGNR .[S]	1xCarbamidome thyl [C2]; 2xDeamidated [N9; N24]	P01023	2726.36654
25	[R].VNQNLVYESGS LNFSKLEIQSQVDS QHVGHSVLTAK.[G]	2xDeamidated [N4; N13]	P04114	3957.98286

26	[K].QLVEIEKVVVLP NYSQVDIGLIK.LK.[Q]	1xDeamidated [N13]	P00738	2876.64956
27	[K].NLFLNHSENAT AKDIAPTLTLYVGK K.[Q]	2xDeamidated [N5; N9]	P00739; P00738	2860.5091
28	[R].VNQNLVYESGS LNFSKLEIQSQVDS QHVGHSVLTAK.[G]	3xDeamidated [N2; N4; N13]	P04114	3958.96688
29	[R].EEQYNSTYRVV SVLTVLHQDWLNG KEYK.[C]	2xDeamidated [N5; N23]	P0DOX5	3400.66958
30	[K].SLGNVNFTVSA EALESQELCGTEVP SVPEHGR.[K]	1xCarbamidome thyl [C20]; 1xDeamidated [N6]	P01023	3414.6118
31	[K].THTNISESHRNA TFSAVGEASICEDD WNSGER.[F]	1xCarbamidome thyl [C23]; 2xDeamidated [N4; N11]	P01871	3520.48297
32	[R].EEQFNSTFRVVS VLTVVHQDWLNGK EYK.[C]	2xDeamidated [N5; N23]	P01859	3354.6641
33	[K].GLKFNLTTTSEA EIQSFQHLR.[T]	1xDeamidated [N5]	P01011	2699.37875
34	[R].ESKPLTAQQTTK LDAPTNLQFVNETD STVLVR.[W]	1xDeamidated [N23]	P02751	3545.83335
35	[K].TKPREEQFNSTF RVVSVLTVVHQDW LNGKEYK.[C]	2xDeamidated [N9; N27]	P01859	3836.96061
36	[K].TKPREEQFNSTF RVVSVLTVVHQDW LNGK.[E]	1xDeamidated [N9]	P01859	3415.77571
37	[K].TIDRLAGKPTHV NVSVVMAEVDGTC Y.[-]	1xCarbamidome thyl [C25]; 1xDeamidated [N13]	P01876	2833.38589
38	[R].TVDKSTGKPTL YNVSLVMSDTAGT CY.[-]	1xCarbamidome thyl [C25]; 1xDeamidated [N13]	P0DOX6; P01871	2809.32704
39	[K].SLGNVNFTVSA EALESQELCGTEVP	1xCarbamidome thyl [C20]; 2xDeamidated	P01023	3543.69078

	SVPEHGRK.[D]	[N4; N6]		
40	[R].GNEANYYSNAT TDEHGLVQFSINTT NVMGTS�TVR.[V]	2xDeamidated [N9; N23]	P01023	3806.745
41	[K].SVQEIQATFFYF TPNKTEDTIFLR.[E]	1xDeamidated [N15]	P02763; P19652	2896.44035
42	[R].TAGWNIPMGLL YNKINHCR.[F]	1xCarbamidome thyl [C18]; 1xDeamidated [N]	P02787	2259.11614
43	[R].LAGKPTHVNVS VVMAEVDGTCY.[-]	1xCarbamidome thyl [C21]; 1xDeamidated [N9]	P01876	2348.12609
44	[R].IQNILTEEPKVT QVYAENGTVLQGS TVASVYK.[G]	1xDeamidated [N18]	P27169	3480.81082
45	[R].STQDTVIALDAL SAYWIASHTTEERG LNVTLSSSTGR.[N]	1xDeamidated [N28]	P0C0L5	3864.92502
46	[K].ELHHLQEQNVS NAFLDKGEFYIGSK. [Y]	1xDeamidated [N9]	P00450	2904.41626
47	[K].VVLHPNYSQVD IGLIKŁK.[Q]	1xDeamidated [N6]	P00738	2037.17429
48	[R].KIVLDPSGSMNI YLVDGSDSIGASN FTGAK.[K]	1xDeamidated [N26]	P00751	3170.59257
49	[R].LSLHRPALEDLL LGSEANLTCTLTGL R.[D]	1xCarbamidome thyl [C21]; 1xDeamidated [N18]	P01876; P01877; P0DOX2	2964.58228
50	[R].KIVLDPSGSMNI YLVDGSDSIGASN FTGAKK.[C]	1xDeamidated [N26]	P00751	3298.68753
51	[R].GIISALLVPPETE EAKQVLFLDTVYG NCSTHFTVK.[T]	1xCarbamidome thyl [C28]; 1xDeamidated [N27]	P04114	3877.99322
52	[K].FNLТETSEAEIH QSFQHLLR.[T]	1xDeamidated [N2]	P01011	2401.17826
53	[K].LNAENNATFYF KIDNVKK.[A]	1xDeamidated [N6]	P01042	2130.0866

54	[K].LPTQNITFQTES SVAEQEAEFQSPK.[Y]	1xDeamidated [N5]	Q14624	2810.33667
55	[K].SKPTVSSSMEFK YDFNSSMLYSTAK.[G]	1xDeamidated [N16]	P04114	2836.30557
56	[K].LGACNDTLQQL MEVFKFDTISEK.[T]	1xCarbamidome thyl [C4]; 1xDeamidated [N5]	P01008	2688.28953
57	[R].EEQFNSTFRVVS VLTVVHQDWLNGK .[E]	2xDeamidated [N5; N23]	P01859	2934.46321
58	[R].QQQHLFGSNVT DCSGNFCLFR.[S]	2xCarbamidome thyl [C13; C18]; 1xDeamidated [N9]	P02787	2516.10815
59	[R].NPPMGGNVVIF DTVITNQEEPYNH SGR.[F]	1xDeamidated [N24]	P02745	3114.45854
60	[K].KLVLSSSEKTVLT PATNHMGNTFTIP ANR.[E]	1xDeamidated [N20]	P01024	3140.67724
61	[R].LFGDKSLTFNET YQDISELVYGAK.[L]	1xDeamidated [N10]	P01008	2739.33997
62	[K].VVLHPNYSQVD IGLIK.[L]	1xDeamidated [N6]	P00738	1795.99526
63	[K].LDAPTNLQFVN ETDSTVLVR.[W]	1xDeamidated [N11]	P02751	2233.13467
64	[R].EEQYNSTYRVV SVLTVLHQDWLNG K.[E]	1xDeamidated [N5]	P0DOX5	2979.48468
65	[K].TVLTPATNHMG NVTFTIPANR.[E]	1xDeamidated [N12]	P01024	2256.14413
66	[K].VTQVYAENGTV LQGSTVASVYKGK. [L]	1xDeamidated [N8]	P27169	2500.29296
67	[K].AVLQLNEEGVD TAGSTGVTNLTSK PIILR.[F]	1xDeamidated [N21]	P08185	3110.69433
68	[R].MAGKPTHINVS VVMAEADGTCY.[-]	1xCarbamidome thyl [C21]; 1xDeamidated	P01877	2352.06686

		[N9]		
69	[K].QEPERNECFLQ HKDDNP NLPR.[L]	1xCarbamidome thyl [C8]; 1xDeamidated [N6]	P02768	2637.21104
70	[R].GFGVAIVGNYT AALPTEAALR.[T]	1xDeamidated [N9]	Q96PD5	2092.10733
71	[R].VLSNNSDANLE LINTWVAK.[N]	1xDeamidated [N]	P05155	2102.07642
72	[K].IITILEEEMNVSV CGLYTYGKPVPGH VTVSICRK.[Y]	2xCarbamidome thyl [C14; C32]; 1xDeamidated [N10]	P01023	3863.97441
73	[R].ITYSIVQTNCSK ENFLFLTPDCK.[S]	2xCarbamidome thyl [C10; C22]; 1xDeamidated [N9]	P01042	2779.33173
74	[K].TVLTPATNHMG NVTFTIPANREFKSE K.[G]	1xDeamidated [N12]	P01024	3004.51968
75	[R].TLNQSSDELQLS MGNAMFVK.[E]	1xDeamidated [N3]	P01011	2214.0417
76	[K].AALAAFNAQNN GSNFQLEEISR.[A]	1xDeamidated [N11]	P02765	2366.13713
77	[K].QVLFLDTVYGN CSTHFTVK.[T]	1xCarbamidome thyl [C12]; 1xDeamidated [N11]	P04114	2230.08488
78	[R].EEQYNSTYRVV SVLTVLHQDWLNG KEYK.[C]	1xDeamidated [N5]	P0DOX5	3399.68556
79	[R].ESVTDHVNLTIP LEKPLQNFTLCFR.[A]	1xCarbamidome thyl [C23]; 1xDeamidated [N19]	P02743	2972.51862
80	[R].VIDFNCTTSSVS SALANTK.[D]	1xCarbamidome thyl [C6]; 1xDeamidated [N5]	P04196	2015.95901
81	[R].GVNFNVSKVSS VEECQKR.[C]	1xCarbamidome thyl [C15]; 1xDeamidated [N5]	P03952	2068.01278
82	[K].LQAPLNYTEFQ	1xCarbamidome	P03952	2248.16822

	KPICLPSK.[G]	thyl [C15]; 1xDeamidated [N6]		
83	[R].QDQCIYNTTYL NVQR.[E]	1xCarbamidome thyl [C4]; 1xDeamidated [N7]	P02763	1916.8807
84	[R].QDQCIYNTTYL NVQRENGTISR.[Y]	1xCarbamidome thyl [C4]; 2xDeamidated [N12; N17]	P02763	2675.23658
85	[K].TLYETEVFSTDF SNISAAKQEINSHVE MQTK.[G]	1xDeamidated [N14]	P05543	3548.67373
86	[K].LGSFEGLVNLTF IHLQHNR.[L]	1xDeamidated [N9]	P51884	2196.15601
87	[K].ERSWPAVGNC SALR.[W]	1xCarbamidome thyl [C10]; 1xDeamidated [N9]	P02790	1690.79658
88	[K].SLGNVNFTVSA EALESQELCGTEVP SVPEHGR.[K]	1xCarbamidome thyl [C20]; 2xDeamidated [N4; N6]	P01023	3415.59582
89	[R].QLAHQSNSTNIF FSPVSIATAFAMLSL GTK.[A]	2xDeamidated [N7; N10]	P01009	3183.60308
90	[K].TVLTPATNHMG NVTFTIPANR.[E]	2xDeamidated [N8; N12]	P01024	2257.12814
91	[K].TVLTPATNHMG NVTFTIPANR.[E]	1xDeamidated [N12]; 1xOxidation [M10]	P01024	2272.13904
92	[K].LHINHNNLTESV GPLPK.[S]	1xDeamidated [N7]	P51884	1883.99739
93	[K].VGQLQLSHNLS LVILVPQNLK.[H]	1xDeamidated [N9]	P05155	2314.34929
94	[K].KLINDYVKNGT R.[G]	1xDeamidated [N9]	P01011	1421.7747
95	[K].GAFISNFSMTVD GKTFR.[S]	1xDeamidated [N6]	P19823	1878.90546
96	[R].TTLSGAPCQPW ASEATYRNVTAEQ AR.[N]	1xCarbamidome thyl [C8]; 1xDeamidated	P00748	2866.34245

		[N19]		
97	[R].QNQCFYNSSYL NVQR.[E]	1xCarbamidome thyl [C4]; 1xDeamidated [N7]	P19652	1921.84974
98	[R].GLNVTLSSTGR NGFK.[S]	2xDeamidated [N3; N12]	P0C0L5	1552.79656
99	[R].LTWEAGADHNS NISEYIVEFEGNK.[E]	3xDeamidated [N10; N12; N23]	O00533	2726.21041
100	[R].NHSCEPCQTLA VR.[S]	2xCarbamidome thyl [C4; C7]; 1xDeamidated [N1]	P00748	1572.68934
101	[R].FNSSYLQGTNQI TGR.[Y]	1xDeamidated [N2]	P04114	1686.80819
102	[R].GLTFQQNASSM CVPDQDTAIR.[V]	1xCarbamidome thyl [C12]; 1xDeamidated [N7]	P0DOX6; P01871	2340.05947
103	[K].VCQDCPLLAPL NDTR.[V]	2xCarbamidome thyl [C2; C5]; 1xDeamidated [N12]	P02765	1772.83058
104	[K].SLTFNETYQDIS ELVYGAK.[L]	1xDeamidated [N5]	P01008	2179.04412
105	[R].ANISHKDMQLG R.[L]	1xDeamidated [N2]	P01031	1370.68451
106	[K].LNAENNATFYF KIDNVK.[K]	1xDeamidated [N6]	P01042	2001.99163
107	[R].KVCQDCPLLAP LNDTR.[V]	2xCarbamidome thyl [C3; C6]; 1xDeamidated [N13]	P02765	1900.92554
108	[K].GCVLLSYLNET VTVSASLESVR.[G]	1xCarbamidome thyl [C2]; 1xDeamidated [N9]	P01023	2398.21702
109	[R].VYKPSAGNNSL YR.[D]	1xDeamidated [N8]	P02749	1469.73832
110	[K].VVNSTTGPEH LR.[N]	1xDeamidated [N3]	P07996	1367.69137
111	[R].LANLTQGEDQY	1xDeamidated	P10909	1684.81769

	YLR.[V]	[N3]		
112	[K].TKPREEQFNSTF R.[V]	1xDeamidated [N9]	P01859	1640.80271
113	[K].NLFLNHSENAT AK.[D]	2xDeamidated [N5; N9]	P00739; P00738	1460.7016
114	[K].KYNSQNSNNQ FVLYR.[I]	1xDeamidated [N6]	P01042	2003.95698
115	[K].YLGNATAIFFLP DEGK.[L]	1xDeamidated [N4]	P01009	1756.87923
116	[R].QQQHLLFGSNVT DCSGNFCLFR.[S]	2xCarbamidome thyl [C13; C18]; 2xDeamidated [N9; N16]	P02787	2517.09217
117	[K].EHEGAIYPDNTT DFQR.[A]	1xDeamidated [N10]	P00450	1893.82496
118	[K].IVLDPSGSMNIY LVLDGSDSIGASNF TGAKK.[C]	1xDeamidated [N25]	P00751	3170.59257
119	[R].LSHNELADSGIP GNSFNVSSLVELDL SYNK.[L]	2xDeamidated [N14; N17]	P51884	3221.54846
120	[K].TAFITNFTLTIDG VTYPGNVKEK.[E]	1xDeamidated [N6]	Q06033	2530.30755
121	[K].ALPQPQNVTSL GCTH.[-]	1xCarbamidome thyl [C14]; 1xDeamidated [N7]	P02790	1736.8636
122	[R].NTSISTAYMELS SLRSEDVAVYYCAR .[-]	1xCarbamidome thyl [C24]; 1xDeamidated [N1]	P0DP01	2989.35538
123	[K].FVEGSHNSTVSL TTK.[N]	1xDeamidated [N7]	P04114	1607.79114
124	[R].GNEANYYSNAT TDEHGLVQFSINTT NVMGTS�TVR.[V]	3xDeamidated [N5; N9; N23]	P01023	3807.72902
125	[R].EEQYNSTYRVV SVLTVLHQDWLNG K.[E]	2xDeamidated [N5; N23]	P0DOX5	2980.46869
126	[R].GCNDSVLAVA GFALR.[D]	1xCarbamidome thyl [C2]; 1xDeamidated [N3]	Q9UGM5	1665.7901

127	[K].STGKPTLYNVSL VMSDTAGTCY.[-]	1xCarbamidome thyl [C21]; 1xDeamidated [N9]	P0DOX6; P01871	2366.08904
128	[K].MDGASNVTCIN SR.[W]	1xCarbamidome thyl [C9]; 1xDeamidated [N6]	P08603	1425.60969
129	[R].FSDGLESNSSTQ FEVKK.[Y]	1xDeamidated [N8]	P0C0L5	1903.89198
130	[K].SPDVINGSPISQ K.[I]	1xDeamidated [N6]	P08603	1342.68489
131	[K].LGNWSAMPSCK .[A]	1xCarbamidome thyl [C10]; 1xDeamidated [N3]	P02749	1251.54966
132	[R].LQNNENNISCVE R.[G]	1xCarbamidome thyl [C10]; 1xDeamidated [N7]	Q03591	1590.71766
133	[K].ENLTAPGSDSA VFFEQGTTR.[I]	1xDeamidated [N2]	P00450	2127.98292
134	[K].TPLTANITK.[S]	1xDeamidated [N6]	P01877; P0DOX2	959.54079
135	[K].LNAENNATFYF K.[I]	1xDeamidated [N6]	P01042	1432.67432
136	[R].ALGFENATQAL GR.[A]	1xDeamidated [N6]	Q08380	1348.68555
137	[R].ITYSIVQTNSK. [E]	1xCarbamidome thyl [C10]; 1xDeamidated [N9]	P01042	1414.68826
138	[R].AQLLQGLGFNL TER.[S]	1xDeamidated [N10]	P08185	1560.83803
139	[R].SWPAVGNCSSA LR.[W]	1xCarbamidome thyl [C8]; 1xDeamidated [N7]	P02790	1405.65287
140	[R].GLNVTLSSSTGR.[N]	1xDeamidated [N3]	P0C0L5	1105.58478
141	[K].LINDYVKNTR. [G]	1xDeamidated [N8]	P01011	1293.67974
142	[K].FLNNGTCTAEG	1xCarbamidome thyl [C7];	P05156	1312.58379

	K.[F]	1xDeamidated [N4]		
143	[K].GAFISNFSMTVD GK.[T]	1xDeamidated [N6]	P19823	1474.68826
144	[K].AGAFLGLTNVA VMNLSGNCLR.[N]	1xCarbamidome thyl [C19]; 1xDeamidated [N14]	P35858	2179.09982
145	[R].EEQFNSTFR.[V]	1xDeamidated [N5]	P01859	1158.50619
146	[R].LSCAASGFTVSN HSMSWVR.[Q]	1xCarbamidome thyl [C3]; 1xDeamidated [N12]; 1xOxidation [M15]	P0DOX2	2113.94299
147	[R].NECFLQHK.[D]	1xCarbamidome thyl [C3]; 1xDeamidated [N1]	P02768	1076.48296
148	[R].ENISDPTSPLR.[T]	1xDeamidated [N2]	P01591	1229.60082
149	[R].EEQFNSTYR.[V]	1xDeamidated [N5]	P01861	1174.50111
150	[R].EEQYNSTYR.[V]	1xDeamidated [N5]	P0DOX5	1190.49602
151	[R].EEQYNSTFR.[V]	1xDeamidated [N5]	P01860	1174.50111
152	[K].NASEIEVPFVTR. [N]	1xDeamidated [N1]	P07093	1362.68997
153	[R].VIDFNCTTSSVS SALANTKDSPVLID FFEDTERYR.[K]	1xCarbamidome thyl [C6]; 1xDeamidated [N5]	P04196	3998.89642
154	[K].SLGPNSCSANGP GLYLIHGPNLYCYS DVEKLNAAK.[A]	2xCarbamidome thyl [C7; C24]; 1xDeamidated [N]	P02790	3780.79962
155	[K].SLGPNSCSANGP GLYLIHGPNLYCYS DVEKLNAAK.[A]	2xCarbamidome thyl [C7; C24]; 2xDeamidated [N5; N10]	P02790	3781.78364
156	[K].FNWYVDGVEV HNAKTKPREEQYNS	1xDeamidated [N23]	P0DOX5	3331.57668

	TYR.[V]			
157	[K].DIAPTLTLYVGK KQLVEIEKVVLHPN YSQVDIGLIK.[L]	1xDeamidated [N26]	P00738	4035.27803
158	[K].ILRQQQHFLFGSN VTDCSGNFCLFR.[S]	2xCarbamidome thyl [C16; C21]; 1xDeamidated [N12]	P02787; A5A6I6	2898.37739
159	[K].LVLSSSEKTVLTP ATNHMGNVFTTIPA NR.[E]	1xDeamidated [N19]	P01024	3012.58228
160	[R].ETYGEMADCCA KQEPERNECFLQHK DDNPNLPR.[L]	3xCarbamidome thyl [C9; C10; C20]; 1xDeamidated [N18]	P02768	4052.72661
161	[R].EGDHEFLEVPEA QEDVEATFPVHQPG NYSCSYR.[T]	1xCarbamidome thyl [C30]; 1xDeamidated [N27]	P04217	3837.66094
162	[K].TKPREEQYNST YRVVSVLTVLHQD WLNGKEYK.[C]	2xDeamidated [N9; N27]	P0DOX5	3882.96609
163	[R].LSLHRPALEDLL LGSEANLTCTLTGL R.[D]	1xCarbamidome thyl [C21]; 1xDeamidated [N18]	P01876; P0DOX2	2964.58228
164	[K].LCMGSGNLNCE PNNKEGYGYTGA FR.[C]	2xCarbamidome thyl [C2; C10]; 1xDeamidated [N]	P02787	2972.30117
165	[R].ADGTVNQIEGE ATPVNLTEPAKLEV K.[F]	1xDeamidated [N16]	P05090	2724.3938
166	[K].LVSANRLF GDK SLTFNETYQDISLV YGAK.[L]	1xDeamidated [N16]	P01008	3379.70563
167	[K].GVTSVSQIFHSP DLAIRDTFVNASR.[T]	1xDeamidated [N22]	P05155	2718.38457
168	[K].VVLHPNYSQVD IGLIKQK.[V]	1xDeamidated [N6]	P00738	2293.32783
169	[R].TAGWNIPMGLL YNKINHCRFDEFFS	2xCarbamidome thyl [C18; C28]; 1xDeamidated	P02787; A5A6I6	3817.75598

	EGCAPGSK.[K]	[N16]		
170	[R].FSDGLESNSSTQ FEVKKYVLPNFEVK .[I]	1xDeamidated [N8]	P0C0L5	2993.47786
171	[R].TAGWNIPMGLL YNKINHCRFDEFFS EGCAPGSK.[K]	2xCarbamidome thyl [C18; C28]; 2xDeamidated [N13; N16]	P02787; A5A6I6	3818.74
172	[K].LPETAERFQDN ASMWSDATAAIPSI EER.[L]	1xDeamidated [N11]	Q4WPQ8	3136.45278
173	[R].EEQYNSTYRVV SVLTVLHQDWLNG KEYKCK.[V]	1xCarbamidome thyl [C29]; 2xDeamidated [N5; N23]	P0DOX5	3688.79519
174	[R].QQQHLFGSNVT DCSGNFCLFRSETK. [D]	2xCarbamidome thyl [C13; C18]; 1xDeamidated [N9]	P02787; A5A6I6	2961.32542
175	[R].TAGWNIPMGLL YNKINHCR.[F]	1xCarbamidome thyl [C18]; 1xDeamidated [N]	P02787; A5A6I6	2259.11614
176	[K].GAVGVSMFMNG TSFGFVNCHLTSGN EKTAR.[R]	1xCarbamidome thyl [C19]; 2xDeamidated [N10; N18]; 1xOxidation [M8]	O15357	3210.46191
177	[K].YPPTVSMVEGQ GEKNVTFWGRPLPR .[C]	1xDeamidated [N15]	P27918	2846.42941
178	[R].WQSIPLCVEKIP CSQPPQIEHGTINSS R.[S]	2xCarbamidome thyl [C7; C13]; 1xDeamidated [N25]	P08603	3262.59834
179	[K].VLNQELRADGT VNQIEGEATPVNLT EPAKLEVK.[F]	1xDeamidated [N23]	P05090	3576.87555
180	[R].SRYPHKPEINST THPGADLQENFCR. [N]	1xCarbamidome thyl [C24]; 1xDeamidated [N10]	P00734	2955.38023
181	[R].TVDKSTGKPTL YNVSLVMSDTAGT	1xCarbamidome thyl [C25]; 1xDeamidated	P01871	2809.32704

	CY.[-]	[N13]		
182	[K].CGLVPVLAENY NKSDNCEDTPEAGY FAVAVVKK.[S]	2xCarbamidome thyl [C1; C17]; 2xDeamidated [N10; N12]	P02787	3659.72439
183	[R].AAINKWVSNKT EGR.[I]	1xDeamidated [N9]	P01008	1574.82853
184	[K].TVLTPATNHMG NVTFTIPANREFK.[S]	1xDeamidated [N12]	P01024	2660.3501
185	[K].TKPREEQYNST YRVVSVLTVLHQD WLNGKEYK.[C]	1xDeamidated [N9]	P0DOX5	3881.98208
186	[R].SPYYNVSDEISF HCYDGYTLR.[G]	1xCarbamidome thyl [C14]; 1xDeamidated [N5]	P00751	2587.1082
187	[R].GLNVTLSSTGR NGFKSHALQLNNR.[Q]	2xDeamidated [N3; N12]	P0C0L5	2586.33829
188	[R].NISDGFDPDN VDAALALPAHSYSG RER.[V]	1xDeamidated [N1]	P04004	3058.45008
189	[R].QQQHLFGSNVT DCSGNFCLFR.[S]	2xCarbamidome thyl [C13; C18]; 1xDeamidated [N9]	P02787; A5A6I6	2516.10815
190	[R].QQQHLFGSNVT DCSGNFCLFR.[S]	2xCarbamidome thyl [C13; C18]; 2xDeamidated [N9; N16]	P02787; A5A6I6	2517.09217
191	[R].AVNITSENLIDD VVSLIRGGTR.[K]	1xDeamidated [N3]	P02748	2343.25143
192	[R].GLTFQQNASM CVPDQDTAIR.[V]	1xCarbamidome thyl [C12]; 1xDeamidated [N7]	P01871	2340.05947
193	[K].AGEQVTYTCAT YYKMDGASNVTCI NSR.[W]	2xCarbamidome thyl [C9; C23]; 1xDeamidated [N20]	P08603	3061.3336
194	[R].VLYLAAYNCTL RPVSK.[K]	1xCarbamidome thyl [C9]; 1xDeamidated	Q9UGM5	1868.99388

		[N8]		
195	[R].FLCTGGVSPYA DPNTRGDSGGPLI VHKR.[S]	2xCarbamidome thyl [C3; C16]; 1xDeamidated [N14]	P00751	3132.49896
196	[K].NLFLNHSENAT AKDIAPTLTLYVGK. [K]	2xDeamidated [N5; N9]	P00739; P00738	2732.41414
197	[R].TAGWNIPMGLL YNKINHCR.[F]	1xCarbamidome thyl [C18]; 2xDeamidated [N5; N13]	P02787; A5A616	2260.10015
198	[K].CVEISCKSPDVI NGSPISQK.[I]	2xCarbamidome thyl [C1; C6]; 1xDeamidated [N13]	P08603	2219.06825
199	[R].AVNITSENLIDD VVSLIR.[G]	1xDeamidated [N3]	P02748	1972.05971
200	[R].VVGVPYQGNAT ALFILPSEGK.[M]	1xDeamidated [N9]	P05154	2161.15395
201	[K].NLFLNHSENAT AK.[D]	1xDeamidated [N5]	P00739; P00738	1459.71758
202	[R].KLPPGLLANFTL LR.[T]	1xDeamidated [N9]	P02750	1553.94138
203	[K].SVVAPATDGGL NLTSTFLR.[K]	1xDeamidated [N12]	P41222	1920.00728
204	[K].KLINDYVKNGT R.[G]	2xDeamidated [N4; N9]	P01011	1422.75872
205	[K].TKPREEQFNSTY R.[V]	1xDeamidated [N9]	P01861	1656.79762
206	[R].FSDGLESNSSTQ FEVK.[K]	1xDeamidated [N8]	P0C0L5	1775.79702
207	[K].TPLTANITK.[S]	1xDeamidated [N6]	P0DOX2	959.54079
208	[R].ISEENETTCYMG K.[W]	1xCarbamidome thyl [C9]; 1xDeamidated [N5]	P08603	1562.6349
209	[K].YTGNASALFILP DQDK.[M]	1xDeamidated [N4]	P01011	1753.86431
210	[K].NLSKLSMLSIRE NK.[I]	1xDeamidated [N1]	Q9UQ13	1633.89417

211	[K].NKSQKQNFNRG GGHR.[G]	3xDeamidated [N1; N6; N9]	Q00839	1659.79461
212	[K].ILRQQQHLFGSN VTDCSGNFCLFRSE TK.[D]	2xCarbamidome thyl [C16; C21]; 1xDeamidated [N12]	P02787	3343.59465
213	[K].ILRQQQHLFGSN VTDCSGNFCLFR.[S]	2xCarbamidome thyl [C16; C21]; 1xDeamidated [N12]	P02787	2898.37739
214	[K].HYTNSSQDVTV PCRVP PPPPCCHPR.[L]	3xCarbamidome thyl [C13; C21; C22]; 1xDeamidated [N4]	P0DOX2	2959.33963
215	[K].LCMGSGNLNCE PNNKEGYGYTGA FR.[C]	2xCarbamidome thyl [C2; C10]; 1xDeamidated [N]; 1xOxidation [M3]	P02787	2988.29609
216	[K].MVSHHNLTGTA TLNQWLLTTAK.[N]	2xDeamidated [N6; N15]; 1xOxidation [M1]	P00739; P00738	2697.35524
217	[R].QDQCIYNTTYL NVQRENGTISR.[Y]	1xCarbamidome thyl [C4]; 2xDeamidated [N7; N12]	P02763	2675.23658
218	[R].VVHAAKAALA AFNAQNNGSNFQLE EISR.[A]	1xDeamidated [N17]	P02765	2971.50206
219	[K].ILRQQQHLFGSN VTDCSGNFCLFR.[S]	2xCarbamidome thyl [C16; C21]; 2xDeamidated [N12; N19]	P02787	2899.36141
220	[K].MVSHHNLTGTA TLNQWLLTTAK.[N]	2xDeamidated [N6; N15]	P00739; P00738	2681.36033
221	[R].VYIHPFHLVIHN ESTCEQLAK.[A]	1xCarbamidome thyl [C16]; 1xDeamidated [N12]	Q9GLN8	2536.26531
222	[K].KLHINHNLTES VGPLPK.[S]	1xDeamidated [N8]	P51884	2012.09235
223	[R].GKLSFWEAGTT KAGYPYNYNTTAS	3xDeamidated	Q9IVZ8	3947.87225

	DQILIENAAGHR.[V]	[N18; N20; N31]		
224	[K].RHEEGHMLNCT CFGQGR.[G]	2xCarbamidome thyl [C10; C12]; 1xDeamidated [N9]	P02751	2089.87492
225	[R].TVDKSTGKPTL YNVSLVMSDTAGT CY.[-]	1xCarbamidome thyl [C25]; 1xDeamidated [N13]; 1xOxidation [M18]	P01871	2825.32195
226	[K].YNSQNQSNNQF VLYR.[I]	1xDeamidated [N]	P01042	1875.86202
227	[R].VLSAMINSNDD NGVLGNWSGTYT GGRDPR.[S]	1xDeamidated [N18]; 1xOxidation [M5]	Q08188	3154.44943
228	[R].TSPSLSPQVNGT PSRNYPATSMVSGL SSPR.[T]	2xDeamidated [N10; N16]	Q8WXI7	3077.48442
229	[R].CIQANYSLMEN GKIK.[V]	1xCarbamidome thyl [C1]; 2xDeamidated [N5; N11]	P05090	1770.84008
230	[R].GGSSGWSGGLA QNRSTITYR.[S]	1xDeamidated [N13]	P07357	2055.98426
231	[R].GLCVNASAVSR. [L]	1xCarbamidome thyl [C3]; 1xDeamidated [N5]	P17936	1134.55718
232	[R].QPPAPITMNQS LNPPNK.[T]	2xDeamidated [N10; N14]	E7F1U2	1945.96879

Table S3. The details of glycopeptides in TB serum after digestion and enrichment by Fe₃O₄@DOPA/PEI@Au@DNA TET-Trypsin.

Number	Annotated Sequence	Modifications	Master protein	MH+ [Da]
1	[R].VIDFNCTTSSVS SALANTKDSPVLID FFEDTERYR.[K]	1xCarbamidomethyl [C6]; 1xDeamidated [N5]	P04196	3998.89642
2	[R].TLNQSSDELQLS MGNAMFVKEQLSL LDRFTEDAKR.[L]	1xDeamidated [N3]	P01011	4015.97396
3	[R].RNPPMGGNVVI FDTVITNQEEPQYQN HSGR.[F]	1xDeamidated [N25]	P02745	3270.55965
4	[R].ETYGEMADCCA KQEPERNECFLQHK DDNPNLPR.[L]	3xCarbamidomethyl [C9; C10; C20]; 1xDeamidated [N18]	P02768	4052.72661
5	[R].NSITLTNLTPGT EYVVSIVALNGREE SPLLIGQQSTVSDVPR.[D]	1xDeamidated [N]	P02751	4469.34098
6	[R].QLAHQSNSTNIF FSPVSIATAFAMLSL GTK.[A]	1xDeamidated [N]	P01009	3182.61906
7	[R].IADAHLDRVEN TTVYYLVLDVQESD CSVLSR.[K]	1xCarbamidomethyl [C26]; 1xDeamidated [N11]	P04196	3581.74282
8	[K].LATALSLSNKFV EGSHNSTVSLTTK.[N]	1xDeamidated [N17]	P04114	2606.36719
9	[K].AGLQAFFQVQE CNKSSSKDNIR.[G]	1xCarbamidomethyl [C12]; 1xDeamidated [N13]	P00450	2528.21981
10	[R].NQALNLSLAYS FVTPLTSMVVTKPD DQEQSQVAEKPME GESR.[N]	1xDeamidated [N5]	Q14624	4639.25421
11	[K].ELHHLQEQNVS NAFLDKGEFYIGSK. [Y]	1xDeamidated [N9]	P00450	2904.41626

12	[K].YLGNATAIFFLP DEGKLQHLENELTH DIITK.[F]	1xDeamidated [N4]	P01009	3541.82133
13	[K].ADTHDEILEGLN FNLTEIPEAQIHEGF QELLR.[T]	1xDeamidated [N14]	P01009	3692.80786
14	[K].LGACNDTLQQL MEVFKFDTISEK.[T]	1xCarbamidome thyl [C4]; 1xDeamidated [N5]	P01008	2688.28953
15	[K].MVSHHNLTGA TLNEQWLLTTAK.[N]	1xDeamidated [N6]	P00739; P00738	2680.37631
16	[K].ILRQQQHFLFSN VTDCSGNFCLFR.[S]	2xCarbamidome thyl [C16; C21]; 1xDeamidated [N12]	P02787	2898.37739
17	[R].EEQYNSTYRVV SVLTVLHQDWLNG KEYK.[C]	2xDeamidated [N5; N23]	P01857; P0DOX5	3400.66958
18	[R].ESVTDHVNLTIP LEKPLQNFTLCFR.[A]	1xCarbamidome thyl [C23]; 1xDeamidated [N19]	P02743	2972.51862
19	[R].WVLTAAHCLLY PPWDKNFTENDLLV R.[I]	1xCarbamidome thyl [C8]; 1xDeamidated [N17]	P00734	3172.59245
20	[R].HGIQYFNNNTQ HSSLFMLNEVKR.[A]	1xDeamidated [N8]	P01042	2778.34166
21	[K].AALAAFNAQNN GSNFQLEEISR.[A]	1xDeamidated [N11]	P02765	2366.13713
22	[K].GVTSVSQIFHSP DLAIRDTFVNASR.[T]	1xDeamidated [N22]	P05155	2718.38457
23	[R].TEVSSNHVLIYL DKVSNQTLSLFFTV LQDVPVRDLKPAIV K.[V]	1xDeamidated [N17]	P01023	4627.53856
24	[K].LVSANRLFGDK SLTFNETYQDISELV YGAK.[L]	1xDeamidated [N16]	P01008	3379.70563
25	[R].LSLHRPAEDLL LGSEANLTCTLTGL	1xCarbamidome thyl [C21];	P01876;	2964.58228

	R.[D]	1xDeamidated [N18]	P0DOX2	
26	[K].ALGISPFHEHAE VVFTANDSGPRR.[Y]	1xDeamidated [N18]	P02766	2608.29027
27	[R].ANLSSQALQMS LDYGFVTPLTSMIS R.[G]	1xDeamidated [N2]	P19827	2831.39539
28	[R].RNPPMGGNVVI FDTVITNQEEPYN HSGR.[F]	2xDeamidated [N18; N25]	P02745	3271.54366
29	[K].TKPREEQYNST YRVVSVLTVLHQD WLNGKEYK.[C]	2xDeamidated [N9; N27]	P01857; P0DOX5	3882.96609
30	[K].YTGNASALFILP DQDKMEEVEAMLL PETLKR.[W]	1xDeamidated [N4]	P01011	3523.76988
31	[K].THTNISESHRNA TFSAVGEASICEDD WNSGER.[F]	1xCarbamidome thyl [C23]; 2xDeamidated [N4; N11]	P01871	3520.48297
32	[R].ESVTDHVNLTIP LEKPLQNFSLCFR.[A]	1xCarbamidome thyl [C23]; 2xDeamidated [N8; N19]	P02743	2973.50263
33	[K].MLNTSSLLEQL NEQFNWVSR.[L]	1xDeamidated [N3]	P10909	2410.17074
34	[R].DTAVFECLPQH AMFGNDTITCTTHG NWTKLPECR.[E]	3xCarbamidome thyl [C7; C21; C33]; 2xDeamidated [N16; N26]	P02749	4009.7612
35	[K].KLVLSSEKTVLT PATNHMGNVFTIP ANR.[E]	1xDeamidated [N20]	P01024	3140.67724
36	[R].SRYPHKPEINST THPGADLQENFCR.[N]	1xCarbamidome thyl [C24]; 1xDeamidated [N10]	P00734	2955.38023
37	[K].SVQEIQATFFYF TPNKTEDTIFLR.[E]	1xDeamidated [N15]	P02763; P19652	2896.44035
38	[R].NPPMGGNVVIF DTVITNQEEPYNH SGR.[F]	1xDeamidated [N24]	P02745	3114.45854

39	[K].NPVGLIGAENA TGETDPSHSK.[F]	1xDeamidated [N10]	P43251	2094.99382
40	[K].QEPERNECFLQ HKDDNPNLPR.[L]	1xCarbamidome thyl [C8]; 1xDeamidated [N6]	P02768	2637.21104
41	[R].LFGDKSLTFNET YQDISELVYGAK.[L]	1xDeamidated [N10]	P01008	2739.33997
42	[R].IQNILTEEPKVT QVYAENGTVLQGS TVASVYKGK.[L]	1xDeamidated [N18]	P27169	3665.92725
43	[R].HGIQYFNNNTQ HSSLFMLNEVK.[R]	1xDeamidated [N8]	P01042	2622.24055
44	[K].GCVLLSYLNET VTVSASLESVRGNR .[S]	1xCarbamidome thyl [C2]; 2xDeamidated [N9; N24]	P01023	2726.36654
45	[K].LKELPGVCNET MMALWEECKPCLK .[Q]	3xCarbamidome thyl [C8; C19; C22]; 1xDeamidated [N9]	P10909	2937.36873
46	[R].QQQHLFGSNVT DCSGNFCLFR.[S]	2xCarbamidome thyl [C13; C18]; 1xDeamidated [N9]	P02787	2516.10815
47	[R].GIISALLVPPETE EAKQVLFLDTVYG NCSTHFTVK.[T]	1xCarbamidome thyl [C28]; 1xDeamidated [N27]	P04114	3877.99322
48	[R].ADGTVNQIEGE ATPVNLTEPAKLEV K.[F]	1xDeamidated [N16]	P05090	2724.3938
49	[K].SLGNVNFTVSA EALSEQELCGTEVP SVPEHGRK.[D]	1xCarbamidome thyl [C20]; 1xDeamidated [N6]	P01023	3542.70677
50	[R].EGDHEFLEVPEA QEDVEATFPVHQPG NYSCSYR.[T]	1xCarbamidome thyl [C30]; 1xDeamidated [N27]	P04217	3837.66094
51	[K].LPTQNITFQTES SVAEQEAEFQSPK.[Y]	1xDeamidated [N5]	Q14624	2810.33667

52	[R].GLTFQQNASSM CVPDQDTAIR.[V]	1xCarbamidome thyl [C12]; 1xDeamidated [N7]	P01871	2340.05947
53	[K].LVLSEKTVLTP ATNHMGNVFTIPA NR.[E]	1xDeamidated [N19]	P01024	3012.58228
54	[K].KLHINHNNLTES VGPLPK.[S]	1xDeamidated [N8]	P51884	2012.09235
55	[K].SLTFNETYQDIS ELVYGAKLQPLDFK ENAEQSR.[A]	1xDeamidated [N5]	P01008	3834.87085
56	[R].NKGCSSTSVLL TLDNNVVNGSSPAI R.[T]	1xCarbamidome thyl [C4]; 1xDeamidated [N20]	P07996	2791.38906
57	[R].FGCEIENNRSSG AFWK.[Y]	1xCarbamidome thyl [C3]; 1xDeamidated [N8]	P25311	1902.84392
58	[R].NISDGFDPDN VDAALALPAHSYSG RER.[V]	1xDeamidated [N1]	P04004	3058.45008
59	[R].EEQYNSTYRVV SVLTVLHQDWLNG K.[E]	1xDeamidated [N5]	P01857; P0DOX5	2979.48468
60	[R].AAINKWVSNKT EGR.[I]	1xDeamidated [N9]	P01008	1574.82853
61	[R].LSVDKDQYVEP ENVTIQCDSGYGVV GPQSITCSGNR.[T]	2xCarbamidome thyl [C18; C32]; 2xDeamidated [N13; N35]	P04003	3973.80662
62	[K].LNAENNATFYF KIDNVKK.[A]	1xDeamidated [N6]	P01042	2130.0866
63	[K].SLGNVNFTVSA EALESQELCGTEVP SVPEHGRK.[D]	1xCarbamidome thyl [C20]; 2xDeamidated [N4; N6]	P01023	3543.69078
64	[R].QDQCIYNTTYL NVQRENGTISR.[Y]	1xCarbamidome thyl [C4]; 2xDeamidated [N7; N12]	P02763	2675.23658
65	[K].VCQDCPLLAPL NDTRVVHAAK.[A]	2xCarbamidome thyl [C2; C5]; 1xDeamidated	P02765	2378.19551

		[N12]		
66	[R].AVNITSENLIDD VVSLIR.[G]	1xDeamidated [N3]	P02748	1972.05971
67	[K].GAVGVSFMFNG TSFGFVNCHLTSGN EKTAR.[R]	1xCarbamidome thyl [C19]; 2xDeamidated [N10; N18]; 1xOxidation [M8]	O15357	3210.46191
68	[K].RHEEGHMLNCT CFGQGR.[G]	2xCarbamidome thyl [C10; C12]; 1xDeamidated [N9]	P02751	2089.87492
69	[R].GVNFNVSKVSS VEECQKR.[C]	1xCarbamidome thyl [C15]; 1xDeamidated [N5]	P03952	2068.01278
70	[K].SLGPNSCSANGP GLYLIHGPNLYCYS DVEKLNAAK.[A]	2xCarbamidome thyl [C7; C24]; 3xDeamidated [N5; N10; N21]	P02790	3782.76765
71	[R].ESKPLTAQQTTK LDAPTNLQFVNETD STVLVR.[W]	1xDeamidated [N23]	P02751	3545.83335
72	[K].ALGISPFHEHAE VVFTANDSGPR.[R]	1xDeamidated [N18]	P02766	2452.18916
73	[K].YTGNASALFILP DQDKMEEVEAMLL PETLK.[R]	1xDeamidated [N4]	P01011	3367.66877
74	[R].SPYYNVSDSISF HCYDGYTLR.[G]	1xCarbamidome thyl [C14]; 1xDeamidated [N5]	P00751	2587.1082
75	[R].DIPTNSPELEETL THTITKLNAENNAT FYFK.[I]	1xDeamidated [N25]	P01042	3552.73805
76	[K].AGEQVTYTCAT YYKMDGASNVTCI NSR.[W]	2xCarbamidome thyl [C9; C23]; 1xDeamidated [N20]	P08603	3061.3336
77	[K].QVLFLDTVYGN CSTHFTVK.[T]	1xCarbamidome thyl [C12]; 1xDeamidated [N11]	P04114	2230.08488

78	[R].HGIQYFNNNTQ HSSLFMLNEVKR.[A]	2xDeamidated [N8; N9]	P01042	2779.32567
79	[R].SQILEGLGFNLT ELSESDVHR.[G]	1xDeamidated [N10]	P29622	2345.16195
80	[R].LAGKPTHVNVS VVMAEVDGTCY.[-]	1xCarbamidome thyl [C21]; 1xDeamidated [N9]	P01876	2348.12609
81	[R].SIPACVPWSPYL FQPNDTCIVSGWGR .[E]	2xCarbamidome thyl [C5; C19]; 1xDeamidated [N16]	P05156	3008.40696
82	[K].ELHHLQEQNVS NAFLDKGEFYIGSK YK.[K]	1xDeamidated [N9]	P00450	3195.57455
83	[K].QEPERNECFLQ HKDDNPNLPR.[L]	1xCarbamidome thyl [C8]; 2xDeamidated [N6; N16]	P02768	2638.19505
84	[R].KLPPGLLANFTL LR.[T]	1xDeamidated [N9]	P02750	1553.94138
85	[K].KYNSQNSNNQ FVLYR.[I]	1xDeamidated [N6]	P01042	2003.95698
86	[R].QDQCIYNTTYL NVQR.[E]	1xCarbamidome thyl [C4]; 1xDeamidated [N7]	P02763	1916.8807
87	[R].NHSCEPCQTLA VR.[S]	2xCarbamidome thyl [C4; C7]; 1xDeamidated [N1]	P00748	1572.68934
88	[R].EEQYNSTYRVV SVLTVLHQDWLNG K.[E]	2xDeamidated [N5; N23]	P01857; P0DOX5	2980.46869
89	[K].KAFITNFSMIID GMTYPGIIKEK.[A]	1xDeamidated [N6]	Q14624	2618.36084
90	[K].TVLTPATNHMG NVTFTIPANR.[E]	1xDeamidated [N12]	P01024	2256.14413
91	[R].GLNVTLSSTGR NGFK.[S]	2xDeamidated [N3; N12]	P0C0L5	1552.79656
92	[R].SKAAGMLMRQ DNASALNSLGIFIW AWPDGPANMPERL	3xDeamidated [N12; N17; N32]	Q5PC41	4559.25199

	SQLAK.[A]			
93	[K].AALAAFNAQNN GSNFQLEEISR.[A]	2xDeamidated [N]	P02765	2367.12114
94	[R].NSITLTNLTPGT EYVVSIVALNGREE SPLLIGQQSTVSDVP R.[D]	2xDeamidated [N7; N22]	P02751	4470.32499
95	[R].GFGVAIVGNYT AALPTEAALR.[T]	1xDeamidated [N9]	Q96PD5	2092.10733
96	[R].NECFLQHKDDN PNLPR.[L]	1xCarbamidome thyl [C3]; 1xDeamidated [N1]	P02768	1997.9134
97	[K].LDAPTNLQFVN ETDSTVLVR.[W]	1xDeamidated [N11]	P02751	2233.13467
98	[R].QNQCFYNSSYL NVQRENGTVSR.[Y]	1xCarbamidome thyl [C4]; 2xDeamidated [N12; N17]	P19652	2666.18997
99	[R].NGTGHGNSTHH GPEYMR.[C]	2xDeamidated [N1; N7]	P02790	1853.76198
100	[R].VLSNNSDANLE LINTWVAK.[N]	1xDeamidated [N4]	P05155	2102.07642
101	[K].SLGPNSCSANGP GLYLIHGPNLYCYS DVEKLNAAK.[A]	2xCarbamidome thyl [C7; C24]; 2xDeamidated [N5; N10]	P02790	3781.78364
102	[R].RNPPMGGNVVI FDTVITNQEEPYN HSGR.[F]	1xDeamidated [N25]; 1xOxidation [M5]	P02745	3286.55456
103	[R].TTLSGAPCQPW ASEATYRNVTAEQ AR.[N]	1xCarbamidome thyl [C8]; 1xDeamidated [N19]	P00748	2866.34245
104	[K].LHINHNNLTESV GPLPK.[S]	1xDeamidated [N7]	P51884	1883.99739
105	[R].VIDFNCTTSSVS SALANTK.[D]	1xCarbamidome thyl [C6]; 1xDeamidated [N5]	P04196	2015.95901
106	[K].KLINDYVKNGT R.[G]	1xDeamidated [N9]	P01011	1421.7747

107	[K].TVLTPATNHMG NVTFTIPANREFKSE K.[G]	1xDeamidated [N12]	P01024	3004.51968
108	[K].SLTFNETYQDIS ELVYGAK.[L]	1xDeamidated [N5]	P01008	2179.04412
109	[K].LGACNDTLQQL MEVFK.[F]	1xCarbamidome thyl [C4]; 1xDeamidated [N5]	P01008	1867.89285
110	[R].TVDKSTGKPTL YNVSLVMSDTAGT CY.[-]	1xCarbamidome thyl [C25]; 1xDeamidated [N13]	P01871	2809.32704
111	[R].QNQCFYNSSYL NVQR.[E]	1xCarbamidome thyl [C4]; 1xDeamidated [N7]	P19652	1921.84974
112	[R].QLAHQSNSTNIF FSPVSIATAFAMLSL GTK.[A]	2xDeamidated [N7; N10]; 1xOxidation [M24]	P01009	3199.59799
113	[R].NISDGFDPDN VDAALALPAHSYSG R.[E]	1xDeamidated [N1]	P04004	2773.30638
114	[K].GAFISNFSMTVD GKTFR.[S]	1xDeamidated [N6]	P19823	1878.90546
115	[K].YNSQNQSNNQF VLYR.[I]	1xDeamidated [N]	P01042	1875.86202
116	[K].VVLHPNYSQVD IGLIK.[L]	1xDeamidated [N6]	P00738	1795.99526
117	[K].TVLTPATNHMG NVTFTIPANR.[E]	2xDeamidated [N8; N12]	P01024	2257.12814
118	[K].VGQLQLSHNLS LVILVPQNLK.[H]	1xDeamidated [N9]	P05155	2314.34929
119	[K].EHEGAIYPDNTT DFQR.[A]	1xDeamidated [N10]	P00450	1893.82496
120	[K].VVNSTTGPEH LR.[N]	1xDeamidated [N3]	P07996	1367.69137
121	[K].ENLTAPGSDSA VFFEQGTTR.[I]	1xDeamidated [N2]	P00450	2127.98292
122	[K].KLINDYVKNGT R.[G]	2xDeamidated [N4; N9]	P01011	1422.75872

123	[R].QQQHLFGSNVT DCSGNFCLFR.[S]	2xCarbamidome thyl [C13; C18]; 2xDeamidated [N9; N16]	P02787	2517.09217
124	[R].KVCQDCPLLAP LNDTR.[V]	2xCarbamidome thyl [C3; C6]; 1xDeamidated [N13]	P02765	1900.92554
125	[R].VYKPSAGNNSL YR.[D]	1xDeamidated [N8]	P02749	1469.73832
126	[K].YTGNASALFILP DQDK.[M]	1xDeamidated [N4]	P01011	1753.86431
127	[R].QDQCIYNTTYL NVQRENGTISR.[Y]	1xCarbamidome thyl [C4]; 3xDeamidated [N7; N12; N17]	P02763	2676.2206
128	[-].MDPMELNNVSIEP DGDSCSGDSIQDSY TGMENSDK.[D]	1xAcetyl [N- Term]; 1xCarbamidome thyl [C18]; 1xDeamidated [N8]; 2xOxidation [M4; M30]	Q8R1S9	3912.51982
129	[K].TELFSSSCPGGI MLNETGQGYQR.[F]	1xCarbamidome thyl [C8]; 1xDeamidated [N15]	O95445	2533.13336
130	[R].TAGWNVPIGTL RPFLNWTGPPEPIEA AVAR.[F]	1xDeamidated [N16]	P02788	3231.69494
131	[K].VTQVYAENGTV LQGSTVASVYKGK. [L]	1xDeamidated [N8]	P27169	2500.29296
132	[K].TVLTPATNHMG NVTFTIPANR.[E]	1xDeamidated [N12]; 1xOxidation [M10]	P01024	2272.13904
133	[K].INVTGSAAAIAD VGNSPTSSTTR.[I]	1xDeamidated [N2]	Q9L5A4	2191.08369
134	[K].QEPERNECFLQ HK.[D]	1xCarbamidome thyl [C8]; 1xDeamidated [N6]	P02768; P02770	1715.78059
135	[K].ALPQPQNVTSL	1xCarbamidome thyl [C14];	P02790	1736.8636

	GCTH.[-]	1xDeamidated [N7]		
136	[K].VSNQTLSLFFTV LQDVPVR.[D]	1xDeamidated [N3]	P01023	2164.16485
137	[K].TKPREEQFNSTF R.[V]	1xDeamidated [N9]	P01859	1640.80271
138	[K].QLDEEIKELNES NSQMEADMIK.[L]	2xDeamidated [N10; N13]; 1xOxidation [M16]	Q9UL68	2612.15897
139	[K].SYNETKIKFDKY K.[A]	1xDeamidated [N3]	P04114	1664.85301
140	[K].TKPREEQFNSTY R.[V]	1xDeamidated [N9]	P01861	1656.79762
141	[R].LANLTQGEDQY YLR.[V]	1xDeamidated [N3]	P10909	1684.81769
142	[K].SLGNVNFTVSA EALESQELCGTEVP SVPEHGR.[K]	1xCarbamidome thyl [C20]; 2xDeamidated [N4; N6]	P01023	3415.59582
143	[K].TKPREEQYNST YR.[V]	1xDeamidated [N9]	P01857; P0DOX5	1672.79254
144	[R].GAFFPLTERNW SLPNR.[A]	1xDeamidated [N10]	P55058	1905.96061
145	[K].NLFLNHSENAT AK.[D]	2xDeamidated [N5; N9]	P00739; P00738	1460.7016
146	[K].YLGNATAIFFLP DEGK.[L]	1xDeamidated [N4]	P01009	1756.87923
147	[R].YAEDKFNETTE K.[S]	1xDeamidated [N7]	P43652	1475.65365
148	[K].NPSTNVSVVVF DSTKDVEDAHSGLL KGNSR.[Q]	3xDeamidated [N1; N5; N28]	Q3T906	3175.53896
149	[K].SPDVINGSPISQ K.[I]	1xDeamidated [N6]	P08603	1342.68489
150	[K].LISNCSKFYGNR .[F]	1xCarbamidome thyl [C5]; 1xDeamidated [N4]	P05156	1459.69983
151	[K].MDGASNVTCIN SR.[W]	1xCarbamidome thyl [C9]; 1xDeamidated	P08603	1425.60969

		[N6]		
152	[R].TAGWNIPMGLLYNKINHCR.[F]	1xCarbamidomethyl [C18]; 1xDeamidated [N16]	P02787	2259.11614
153	[K].VCQDCPLLAPLNDTR.[V]	2xCarbamidomethyl [C2; C5]; 1xDeamidated [N12]	P02765	1772.83058
154	[K].CVEISCKSPDVINGSPISQK.[I]	2xCarbamidomethyl [C1; C6]; 1xDeamidated [N13]	P08603	2219.06825
155	[K].TVLTPATNHMGNVTF TIPANREFK.[S]	1xDeamidated [N12]	P01024	2660.3501
156	[R].GCNDSVDLAVAGFALR.[D]	1xCarbamidomethyl [C2]; 1xDeamidated [N3]	Q9UGM5	1665.7901
157	[K].ENKTLQEEIKDLIDQLGEGGR.[S]	1xDeamidated [N2]	Q9H6N6	2386.20962
158	[R].FNSSYLQGTNQITGR.[Y]	1xDeamidated [N2]	P04114	1686.80819
159	[R].DQCIVDDITYNVNDTFHK.[R]	1xCarbamidomethyl [C3]; 1xDeamidated [N13]	P02751	2197.97064
160	[R].LQNNENNISCVEER.[G]	1xCarbamidomethyl [C10]; 1xDeamidated [N7]	P36980; Q03591	1590.71766
161	[K].NLSMPLLPADFHK.[E]	1xDeamidated [N1]	P05546	1483.76136
162	[R].VVGVPYQGNATALFILPSEGK.[M]	1xDeamidated [N9]	P05154	2161.15395
163	[K].KPGNNEGSGAPSPLSKSSPATTVTSPNSTPAK.[T]	3xDeamidated [N4; N5; N27]	O60641	3069.48586
164	[K].AGAFLGLTNVAVMNLSGNCLR.[N]	1xCarbamidomethyl [C19]; 1xDeamidated [N14]	P35858	2179.09982
165	[K].STGKPTLYNVSL	1xCarbamidomethyl	P01871	2366.08904

	VMSDTAGTCY.[-]	thyl [C21]; 1xDeamidated [N9]		
166	[R].FSDGLESNSSTQ FEVKK.[Y]	1xDeamidated [N8]	P0C0L5	1903.89198
167	[K].YDFNSSMLYST AK.[G]	1xDeamidated [N4]	P04114	1527.66719
168	[R].SWPAVGNCSSA LR.[W]	1xCarbamidome thyl [C8]; 1xDeamidated [N7]	P02790	1405.65287
169	[K].LINDYVKNCTR. [G]	1xDeamidated [N8]	P01011	1293.67974
170	[R].ISEENETTCYMG K.[W]	1xCarbamidome thyl [C9]; 1xDeamidated [N5]	P08603	1562.6349
171	[R].GLNVTLSSTGR.[N]	1xDeamidated [N3]	P0C0L5	1105.58478
172	[R].FSDGLESNSSTQ FEVK.[K]	1xDeamidated [N8]	P0C0L5	1775.79702
173	[K].TPLTANITK.[S]	1xDeamidated [N6]	P0DOX2	959.54079
174	[R].AQLLQGLGFNL TER.[S]	1xDeamidated [N10]	P08185	1560.83803
175	[K].VSNVSCQASVS R.[M]	1xCarbamidome thyl [C6]; 1xDeamidated [N3]	P55058	1294.60559
176	[K].LNAENNATFYF K.[I]	1xDeamidated [N6]	P01042	1432.67432
177	[K].GAFISNFSMTVD GK.[T]	1xDeamidated [N6]	P19823	1474.68826
178	[K].LGNWSAMPSCK .[A]	1xCarbamidome thyl [C10]; 1xDeamidated [N3]	P02749	1251.54966
179	[K].TMFPNLTDVR.[E]	1xDeamidated [N5]	Q8SQ74	1194.58234
180	[K].GIHFNATKLSDS SAK.[L]	1xDeamidated [N5]	C0HLG3	1576.79656
181	[K].WSDIWNATK.[Y]	1xDeamidated	P06276	1121.5262

	]	[N6]		
182	[R].EEQFNSTYR.[V]	1xDeamidated [N5]	P01861	1174.50111
183	[K].FLNNGTCTAEG K.[F]	1xCarbamidome thyl [C7]; 1xDeamidated [N4]	P05156	1312.58379
184	[R].EEQYNSTYR.[V]	1xDeamidated [N5]	P01857; P0DOX5	1190.49602
185	[K].LPPGLLANFTLL R.[T]	1xDeamidated [N8]	P02750	1425.84641
186	[R].EEQFNSTFR.[V]	1xDeamidated [N5]	P01859	1158.50619
187	[K].KQLVEIEKVVL HPNYSQVDIGLIK.[L]	1xDeamidated [N14]	P00738	2763.56549
188	[K].FNWYVDGVEV HNAKTKPREEQYNS TYR.[V]	1xDeamidated [N23]	P0DOX5	3331.57668
189	[K].NLFLNHSENAT AKDIAPTLTLYVGK K.[Q]	2xDeamidated [N5; N9]	P00739; P00738	2860.5091
190	[K].ILRQQQHLFGSN VTDCSGNFCLFR.[S]	2xCarbamidome thyl [C16; C21]; 1xDeamidated [N12]	P02787; A5A6I6	2898.37739
191	[K].MVSHHNLTTGA TLNQWLLTTAK.[N]	1xDeamidated [N6]; 1xOxidation [M1]	P00739; P00738	2696.37123
192	[R].EEQYNSTYRVV SVLTVLHQDWLNG KEYK.[C]	1xDeamidated [N5]	P0DOX5	3399.68556
193	[R].STQDTVIALDAL SAYWIASHTTEERG LNVTLSSSTGR.[N]	1xDeamidated [N28]	P0C0L4; P0C0L5	3864.92502
194	[R].TAGWNIPMGLL YNKINHCR.[F]	1xCarbamidome thyl [C18]; 1xDeamidated [N]	P02787; A5A6I6	2259.11614
195	[K].TKPREEQYNST YRVVSVLTVLHQD WLNGKEYK.[C]	2xDeamidated [N9; N27]	P0DOX5	3882.96609

196	[R].FSLLGHASISCT VENETIGVWRPSPP TCEK.[I]	2xCarbamidome thyl [C11; C28]; 1xDeamidated [N15]	P04003	3373.61914
197	[R].EEQYNSTYRVV SVLTVLHQDWLNG KEYK.[C]	2xDeamidated [N5; N23]	P0DOX5	3400.66958
198	[K].YPPTVSMVEGQ GEKNVTFWGRPLPR .[C]	1xDeamidated [N15]	P27918	2846.42941
199	[R].TLNQSSDELQLS MGNAMFVKEQLSL LDRFTEDAK.[R]	2xDeamidated [N3; N15]	P01011	3860.85686
200	[K].MVSHHNLTGA TLINEQWLLTTAKN LFLNHSENATAK.[D]	3xDeamidated [N6; N15; N25]	P00739; P00738	4122.06007
201	[K].QLVEIEKVVLHP NYSQVDIGLIK.[L]	1xDeamidated [N13]	P00738	2635.47053
202	[K].VVLHPNYSQVD IGLIK.[Q]	1xDeamidated [N6]	P00738	2037.17429
203	[K].SLGPNSCSANGP GLYLIHGPNLYCYS DVEKLNAAK.[A]	2xCarbamidome thyl [C7; C24]; 1xDeamidated [N]	P02790	3780.79962
204	[K].SLGPNSCSANGP GLYLIHGPNLYCYS DVEKLNAAK.[A]	2xCarbamidome thyl [C7; C24]; 2xDeamidated [N10; N]	P02790	3781.78364
205	[K].TPENYPNAGLT MNYCRNPDADKGP WCFTTDPSVR.[W]	2xCarbamidome thyl [C15; C26]; 1xDeamidated [N13]	P00747	3945.72653
206	[K].MVSHHNLTGA TLINEQWLLTTAK.[N]	2xDeamidated [N6; N15]	P00739; P00738	2681.36033
207	[K].TIDRLAGKPTHV NVSVMMAEVDGTC Y.[-]	1xCarbamidome thyl [C25]; 1xDeamidated [N13]	P01876	2833.38589
208	[K].TKPREEQYNST YRVVSVLTVLHQD WLNGK.[E]	2xDeamidated [N9; N27]	P0DOX5	3462.76521
209	[K].ILRQQHFLFGSN VTDCSGNFCLFRSE	2xCarbamidome thyl [C16; C21];	P02787; A5A6I6	3343.59465

	TK.[D]	1xDeamidated [N12]		
210	[R].LSLHRPALEDLL LGSEANLTCTLTGL R.[D]	1xCarbamidome thyl [C21]; 1xDeamidated [N18]	P01876; P01877; P0DOX2	2964.58228
211	[R].WQSIPLCVEKIP CSQPPQIEHGTINSS R.[S]	2xCarbamidome thyl [C7; C13]; 1xDeamidated [N25]	P08603	3262.59834
212	[R].EEQFNSTFRVVS VLTVVHQDWLNGK EYK.[C]	2xDeamidated [N5; N23]	P01859	3354.6641
213	[K].TKPREEQFNSTF RVVSVLTVVHQDW LNGKEYK.[C]	1xDeamidated [N9]	P01859	3835.9766
214	[R].KVCQDCPLLAP LNDTRVVHAAK.[A]	2xCarbamidome thyl [C3; C6]; 1xDeamidated [N13]	P02765	2506.29047
215	[R].VYIHPFHLVIHN ESTCEQLAK.[A]	1xCarbamidome thyl [C16]; 1xDeamidated [N12]	Q9GLN8	2536.26531
216	[K].NLFLNHSENAT AKDIAPTLTLYVGK. [K]	2xDeamidated [N5; N9]	P00739; P00738	2732.41414
217	[K].CGLVPVLAENY NKSDNCEDTPEAGY FAVAVVKK.[S]	2xCarbamidome thyl [C1; C17]; 2xDeamidated [N10; N12]	P02787	3659.72439
218	[R].ENGTVSRYEGG REHVAHLLFLR.[D]	1xDeamidated [N2]	P19652	2541.29569
219	[K].LGSFEGLVNLTF IHLQHNR.[L]	1xDeamidated [N9]	P51884	2196.15601
220	[R].TVDKSTGKPTL YNVSLVMSDTAGT CY.[-]	1xCarbamidome thyl [C25]; 1xDeamidated [N13]; 1xOxidation [M18]	P01871	2825.32195
221	[R].IQNILTEEPKVT QVYAENGTVLQGS TVASVYK.[G]	2xDeamidated [N3; N18]	P27169	3481.79484

222	[R].EYQTRQDQCIY NTTYLNVQRENGTI SR.[Y]	1xCarbamidome thyl [C9]; 2xDeamidated [N17; N22]	P02763	3352.54987
223	[R].GLNVTLSSTGR NGFK.[S]	2xDeamidated [N3; N12]	P0C0L4; P0C0L5	1552.79656
224	[R].QQQHLFGSNVT DCSGNFCLFR.[S]	2xCarbamidome thyl [C13; C18]; 1xDeamidated [N9]	P02787; A5A6I6	2516.10815
225	[R].VLSNNSDANLE LINTWVAK.[N]	1xDeamidated [N]	P05155	2102.07642
226	[K].SLGPNSCSANGP GLYLIHGPNLYCYS DVEK.[L]	2xCarbamidome thyl [C7; C24]; 2xDeamidated [N5; N10]	P02790	3284.48746
227	[R].VLYLAAYNCTL RPVSK.[K]	1xCarbamidome thyl [C9]; 1xDeamidated [N8]	Q9UGM5	1868.99388
228	[K].ERSWPAVGNC SALR.[W]	1xCarbamidome thyl [C10]; 1xDeamidated [N9]	P02790	1690.79658
229	[R].ITYSIVQTNCSK ENFLFLTPDCK.[S]	2xCarbamidome thyl [C10; C22]; 1xDeamidated [N9]	P01042	2779.33173
230	[R].LLDTQRDGLQN YEALLGLTNLSGR. S]	2xDeamidated [N11; N20]	Q8IWX7	2662.36825
231	[K].IPASKIVQYDMG NLSGEFPLEVTLAV K.[E]	1xDeamidated [N13]	Q2NHZ0	2920.53762
232	[R].ANISHKDMQLG R.[L]	1xDeamidated [N2]	P01031	1370.68451
233	[K].FVEGSHNSTVSL TTK.[N]	1xDeamidated [N7]	P04114	1607.79114
234	[R].IIVPLNNRENISD PTSPLR.[T]	1xDeamidated [N10]	P01591	2149.16116
235	[K].SVVAPATDGGL NLTSTFLR.[K]	1xDeamidated [N12]	P41222	1920.00728
236	[K].QSVPAHFVALN GSK.[L]	1xDeamidated [N11]	Q863A0	1455.75905

237	[R].VLSNNSDANLE LINTWVAK.[N]	2xDeamidated [N4; N5]	P05155	2103.06044
238	[R].FSDGLESNSSTQ FEVKK.[Y]	1xDeamidated [N8]	P0C0L4; P0C0L5	1903.89198
239	[K].VETNISKTSWIR SSMAASGKR.[V]	1xDeamidated [N4]	Q15849	2310.18705
240	[K].LCMGSGNLNCE PNNKEGYGYTGA FR.[C]	2xCarbamidome thyl [C2; C10]; 2xDeamidated [N8; N13]	P02787	2973.28519
241	[K].TKPREEQYNST YR.[V]	1xDeamidated [N9]	P0DOX5	1672.79254
242	[R].ALGFENATQAL GR.[A]	1xDeamidated [N6]	Q08380	1348.68555
243	[R].ITYSIVQTNSK. [E]	1xCarbamidome thyl [C10]; 1xDeamidated [N9]	P01042	1414.68826
244	[R].FSDGLESNSSTQ FEVK.[K]	1xDeamidated [N8]	P0C0L4; P0C0L5	1775.79702
245	[K].TPLTANITK.[S]	1xDeamidated [N6]	P01877; P0DOX2	959.54079
246	[R].FEVDSPVYNAT WSASLK.[N]	1xDeamidated [N9]	P04114	1914.91199
247	[R].ENISDPTSPLR.[T]	1xDeamidated [N2]	P01591	1229.60082
248	[R].GLNVTLSSTGR.[N]	1xDeamidated [N3]	P0C0L4; P0C0L5	1105.58478
249	[R].VNQNLVYESGS LNFSK.[L]	1xDeamidated [N13]	P04114	1799.88102
250	[R].EEQYNSTYR.[V]	1xDeamidated [N5]	P0DOX5	1190.49602
251	[R].EEQYNSTFR.[V]	1xDeamidated [N5]	P01860	1174.50111
252	[R].LSLHRPALEDLL LGSEANLTCTLTGL RDASGATFTWTPSS GK.[S]	1xCarbamidome thyl [C21]; 1xDeamidated [N18]	P0DOX2	4458.26095
253	[R].VNQNLVYESGS LNFSKLEIQSQVDS QHVGHSVLTAK.[G]	1xDeamidated [N13]	P04114	3956.99885
254	[R].NEEYNKSVQEIQ	2xDeamidated	P02763; P19652	3674.7537

	ATFFYFTPKNKTEDI FLR.[E]	[N5; N21]		
255	[R].QLAHQSNSTNIF FSPVSIATAFAMLSL GTK.[A]	1xDeamidated [N7]	P01009	3182.61906
256	[K].AREDIFMETLK DIVEYYNDSNGSHV LQGR.[F]	1xDeamidated [N21]	P25311	3400.61141
257	[R].SDHGSSISCQPP AEIPGYLPADTVHL AVEFFNLTHLPANL LQGASK.[L]	1xCarbamidome thyl [C9]; 1xDeamidated [N32]	P02750	4900.42506
258	[R].TLNQSSDELQLS MGNAMFVKEQLSL LDRFTEDAK.[R]	1xDeamidated [N3]	P01011	3859.87284
259	[R].SNHVS RTEVSSN HVLIIYLDKVS NQTL SLFFT V LQDVPVR.[D]	1xDeamidated [N23]	P01023	4443.33069
260	[R].QLAHQSNSTNIF FSPVSIATAFAMLSL GTK.[A]	1xDeamidated [N7]; 1xOxidation [M24]	P01009	3198.61398
261	[K].TKPREEQYNST YRVVSVLTVLHQD WLNGKEYK.[C]	1xDeamidated [N9]	P0DOX5	3881.98208
262	[R].SEGSSVNLSPPL EQCV PDRGQQYQG R.[L]	1xCarbamidome thyl [C15]; 1xDeamidated [N7]	P00734	2889.34317
263	[K].MVSHHNLTTGA TLINEQWLLTTAKN LFLNHSENATAK.[D]	3xDeamidated [N15; N25; N29]	P00739; P00738	4122.06007
264	[R].QLAHQSNSTNIF FSPVSIATAFAMLSL GTK.[A]	2xDeamidated [N7; N10]	P01009	3183.60308
265	[R].DKICDLLVANN HFAHFFAPQNL TN MNK.[N]	1xCarbamidome thyl [C4]; 1xDeamidated [N21]	P19827	3173.52953
266	[K].QLVEIEKVVLHP NYSQVDIGLIK LK.[Q]	1xDeamidated [N13]	P00738	2876.64956

267	[R].EEQYNSTYRVV SVLTVLHQDWLNG K.[E]	2xDeamidated [N5; N23]	P0DOX5	2980.46869
268	[K].TKPREEQFNSTF RVVSVLTVVHQDW LNGKEYK.[C]	2xDeamidated [N9; N27]	P01859	3836.96061
269	[R].EEQYNSTYRVV SVLTVLHQDWLNG K.[E]	1xDeamidated [N5]	P0DOX5	2979.48468
270	[K].IITILEEEMNVSV CGLYTYGKPVPGH VTVSICR.[K]	2xCarbamidome thyl [C14; C32]; 1xDeamidated [N10]	P01023	3735.87945
271	[K].HYTNSSQDVTV PCRVPPPPCCHPR.[L]	3xCarbamidome thyl [C13; C21; C22]; 1xDeamidated [N4]	P0DOX2	2959.33963
272	[R].VDHRGLTFQQN ASSMCVPDQDTAIR .[V]	1xCarbamidome thyl [C16]; 1xDeamidated [N11]	P01871	2847.31485
273	[R].KSRPANHCVYF YGDEISFSCHETSR.[F]	2xCarbamidome thyl [C8; C20]; 1xDeamidated [N6]	P04003	3048.33631
274	[R].TAGWNIPMGLL YNKINHCR.[F]	1xCarbamidome thyl [C18]; 1xDeamidated [N]	P02787	2259.11614
275	[K].MLNTSSLLEQL NEQFNWVSR.[L]	1xDeamidated [N3]; 1xOxidation [M1]	P10909	2426.16565
276	[K].AREDIFMETLK DIVEYYNDSNGSHV LQGR.[F]	2xDeamidated [N18; N21]	P25311	3401.59543
277	[R].VNQNLVYESGS LNFSKLEIQSQVDS QHVGHSVLTA.[G]	2xDeamidated [N4; N13]	P04114	3957.98286
278	[K].NCGVNCSGDVF TALIGEIASPNYPKP YPENSR.[C]	2xCarbamidome thyl [C2; C6]; 1xDeamidated [N5]	P09871	3527.62059
279	[R].FEDGVLPDPYP RNISDGFDPDNPV	2xDeamidated	P04004	4177.92614

	DAALALPAHSYSGR .[E]	[N13; N24]		
280	[R].DIPTNSPELEETL THTITKLNAENNAT FYFKIDNVKK.[A]	1xDeamidated [N25]	P01042	4250.15032
281	[R].GNEANYYSNAT TDEHGLVQFSINTT NVMGTSLTVR.[V]	2xDeamidated [N9; N23]	P01023	3806.745
282	[R].NISDGFDPDN VDAALALPAHSYSG R.[E]	2xDeamidated [N1; N12]	P04004	2774.29039
283	[K].NLFLNHSENAT AKDIAPTLTLYVGK K.[Q]	3xDeamidated [N1; N5; N9]	P00739; P00738	2861.49312
284	[K].ALQAVYSMMS WPDDVPPEGWNRT R.[H]	1xDeamidated [N21]	P00751	2807.2916
285	[K].LVLSEKTVLTP ATNHMGNVFTIPA NR.[E]	2xDeamidated [N15; N19]	P01024	3013.5663
286	[R].ETYGEMADCCA KQEPERNECFLQHK .[D]	3xCarbamidome thyl [C9; C10; C20]; 1xDeamidated [N18]	P02768	3131.29617
287	[R].FSGSNSGNTATL TISR.[VA]	1xDeamidated [N8]	P80748	1613.77655
288	[R].AVNITSENLIDD VVSLIRGGTR.[K]	1xDeamidated [N3]	P02748	2343.25143
289	[K].HYTNSSQDVT PCR.[V]	1xCarbamidome thyl [C13]; 1xDeamidated [N4]	P0DOX2	1664.73331
290	[K].KLVLSSEKTVLT PATNHMGNVFTIP ANR.[E]	2xDeamidated [N16; N20]	P01024	3141.66126
291	[R].VIDFNCTTSSVS SALANTKDSPVLID FFEDTER.[Y]	1xCarbamidome thyl [C6]; 1xDeamidated [N5]	P04196	3679.73198
292	[K].NLFLNHSENAT AK.[D]	1xDeamidated [N5]	P00739; P00738	1459.71758
293	[R].IQNILTEEPKVT QVYAENGTVLQGS	1xDeamidated [N18]	P27169	3480.81082

	TVASVYK.[G]			
294	[K].NNATVHEQVGG PSLTSDLQAQSK.[G]	1xDeamidated [N2]	P04004	2382.15317
295	[R].VFHIHNESWVL LTPK.[A]	1xDeamidated [N6]	O75882	1820.96938
296	[K].AGLQAFFQVQE CNKSSSK.[D]	1xCarbamidome thyl [C12]; 1xDeamidated [N13]	P00450	2029.96477
297	[K].QVFPGLNYCTS GAYSNASSTDSASY YPLTGDTR.[L]	1xCarbamidome thyl [C9]; 1xDeamidated [N16]	P04114	3551.55435
298	[K].QSVPAHFVALN GSK.[L]	1xDeamidated [N11]	P06681	1455.75905
299	[R].CIQANYSLMEN GKIK.[V]	1xCarbamidome thyl [C1]; 2xDeamidated [N5; N11]	P05090	1770.84008
300	[K].TQKDVISNTSDV IGTCEAADVAQK.[V]	1xCarbamidome thyl [C16]; 1xDeamidated [N8]	Q9H2G2	2551.2192
301	[K].TVLTPATNHMG NVTFTIPANREFK.[S]	2xDeamidated [N8; N12]	P01024	2661.33411
302	[K].AALAAFNAQNN GSNFQLEEISR.[A]	2xDeamidated [N10; N11]	P02765	2367.12114
303	[R].CSDGWSFDATT LDDNGTMLFFK.[G]	1xCarbamidome thyl [C1]; 1xDeamidated [N15]	P02790	2529.05847
304	[K].CGLVPVLAENY NKSDNCEDTPEAGY FAVAVVK.[K]	2xCarbamidome thyl [C1; C17]; 2xDeamidated [N12; N16]	P02787	3531.62943
305	[R].GLTFQQNASM CVPDQDTAIR.[V]	1xCarbamidome thyl [C12]; 1xDeamidated [N7]; 1xOxidation [M11]	P01871	2356.05439
306	[K].IYPGVDFGGEEL NVTQVK.[G]	1xDeamidated [N13]	P03952	1984.99024

307	[K].HANWTLTPLK.[S]	1xDeamidated [N3]	P27169	1181.63133
308	[K].VNGTFQANFPL GPATHGGTYR.[C]	1xDeamidated [N2]	Q14954	2206.06759
309	[R].WEYCNLTR.[C]	1xCarbamidome thyl [C4]; 1xDeamidated [N5]	P08519	1142.49352
310	[K].TMFPNLTDVR.[E]	1xDeamidated [N5]	P06681	1194.58234
311	[R].KLNGCTKEEFF AYR.[R]	1xCarbamidome thyl [C5]; 1xDeamidated [N3]	P07720	1763.84213

Table S3. Sequences of four single-stranded DNA used to synthesize DNA tetrahedron.

No.	DNA single-stranded sequence (5' to 3')	5' modification
N1	TACACAGATCATAGTAGGTAAGTTATCGAAC	5' SH C6
N2	ATCTGTGTACAGACGACGAATCCCTATCGGA	5' SH C6
N3	TTACAGTCTGCTTCGTCGTCGAGTTCGATAA	5' SH C6
N4	TCAGTCGGTCAGACTGTAAACTACCTACTAGATCCGATA GG	5' COOH

Table S4. The comparison table of Fe₃O₄@DOPA/PEI@Au@DNA TET-Trypsin and other previous materials in references.

NO.	Materials used	LOD	Selectivity	Total required time	Standard protein	Ref.
1	SCOF-2	0.010 fmol·μL ⁻¹	BSA: HRP= 5000: 1	20.5 h	HRP, IgG	[1]
2	CCM	0.250 fmol·μL ⁻¹	BSA: IgG= 1000: 1	18 h	IgG	[2]
3	3D-IL-COF-1 @CS hydrogel	0.100 fmol·μL ⁻¹	BSA: HRP= 2000: 1	19 h	HRP	[3]
4	mM-TiO ₂ @Cys	1.00 fmol·μL ⁻¹	-	19 h	IgG	[4]
5	Fe ₃ O ₄ @DOPA/PEI@A u @DNA TET-Trypsin	0.001 amol·μL ⁻¹	BSA: HRP= 5000: 1	20 min	HRP	This work

‘-’ not given.

- [1] S. S. Yang, Y. H. Jiang, S. J. Jiang, L. H. Liu, S. Liu, H. Zhang, Z. Y. Gu, Hydrophilic Sulfonate Covalent Organic Frameworks for Serum Glycopeptide Profiling, *Int. J. Mol. Sci.*, 2025, **26**, 1957. DOI: 10.3390/ijms26051957.
- [2] G. Wang, L. W. Zhang, A. Sugawara, Y. Hsu, T. A. Asoh, H. Uyama, Development of Citric-Acid-Modified Cellulose Monolith for Enriching Glycopeptides, *Anal. Chem.*, 2025, **97**, 1125-1134. DOI: 10.1021/acs.analchem.4c03857.
- [3] X. Q. Zhang, D. N. Wang, J. H. Shao, H. J. Mao, C.F. Ding, Y. H. Yan, Hydrophilic materials based on hydrogel polymers for enrichment of glycopeptides from serum of colorectal cancer patients, *J. SEP. SCI.*, 2024, **47**, 2400310. DOI: 10.1002/jssc.202400310.
- [4] Y. F. Lin, C. R. Du, H. M. Ying, Y. F. Zhou, F. F. Kong, H. L. Zhao, M. B. Lan, Multiply-mesoporous hydrophilic titanium dioxide nanohybrid for the highly-performed enrichment of N-glycopeptides from human serum, *ANAL. CHIM. ACTA*, 2024, **1287**, 342058. DOI: 10.1016/j.aca.2023.342058.