

Supporting Information

Guanidyl-Functionalized Graphene as a Bi-function Adsorbent for Selective Enrichment of Phosphopeptides

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Experimental Section

Materials

Graphene oxide (GO) was purchased from Nanjing XFNANO Materials Tech Co., Ltd. Thionyl chloride (SOCl₂) was purchased from Sinopharm Chemical Reagent Co., Ltd. 1,6-hexanediamine, O-methylisourea hemisulfate (OMIU), and 2,5-dihydroxybenzoic acid (DHB) were purchased from J&K Scientific Ltd. N,N-dimethylformamide (DMF) was from J. T. Baker. Trypsin, α -casein, β -casein, and acetonitrile (ACN) were purchased from Sigma-Aldrich. Ammonium bicarbonate (NH₄HCO₃) was purchased from Fluka. Trifluoroacetic acid (TFA) was obtained from Acros Organics. Acetic acid (HAc), phosphoric acid (H₃PO₄) and sodium hydroxide (NaOH) were from Beijing Chemical Works. Water used as solvent was from Wahaha Group Co., Ltd. Non-fat milk was purchased from a local supermarket.

Preparation of amino- and guanidyl-functionalized graphene

Synthesis of amino-functionalized graphene (AFG)

120 mg GO, 30 mL SOCl₂, and 1.5 mL DMF were mixed in a 100 mL flask. After sonification for sufficient dispersion of GO in solvent, the mixture was stirred at 70°C for 24 h. The acylated intermediate was separated from solution by centrifugation, washed with dry DMF and directly transferred into 30 mL 1,6-hexanediamine heated at 55°C. The mixture was stirred at 55°C for 48 h and then diluted with 25 mL ethanol. The product was washed with water and ethanol for several times and then dried under vacuum at room temperature.

Synthesis of guanidyl-functionalized graphene (GFG)

50 mg AFG, 3.0781 g OMIU, and 50 mL water were mixed in a 100 mL flask. After sonification for dispersion of AFG and dissolution of OMIU, the pH was adjusted to 11 by NaOH. The mixture was stirred at 60°C for 24 h and the reaction was terminated by adding

1.65 mL TFA. After washing by water and ethanol, the product was dried under vacuum at room temperature.

Preparation of α -casein, β -casein, and non-fat milk digests

α -casein and β -casein were dissolved in 50 mM NH_4HCO_3 solution. After addition of trypsin with 1:40 (w/w) enzyme-to-protein ratio, the proteins were digested at 37°C for 20 h.

30 μL non-fat milk was diluted with 970 μL 50 mM NH_4HCO_3 solution. The diluent was centrifuged under 14,000 rpm for 25 min. After denaturation of the supernatant at 100°C for 5 min, 30 μg trypsin was added and the mixture was allowed for digestion at 37°C for 20 h.

Enrichment of phosphopeptides

In general, the tryptic digests were diluted with loading buffer (buffer A: 50% ACN, 0.1 M HAC; or buffer B: 66.6% ACN, 0.05% TFA) to a certain concentration, and the adsorbent (AFG or GFG) was dispersed in water by sonification. 50 μL suspension of adsorbent (1 mg/mL-4 mg/mL) was added to 200 μL peptide diluent, and the mixture was gently shaken by a rotator. After centrifugation and removal of the supernatant, the adsorbent was washed by 100 μL loading buffer for three times and mixed with 5 μL matrix solution of ACN/H₂O 1:1 (v/v) containing 30 mg/mL DHB and 5% H₃PO₄.

Instrumentation

Infrared spectrum was measured with KBr pellet by Bruker Tensor 27 FT-IR. Absorbance spectrum was acquired with resolution of 4 cm^{-1} and subtraction of background of air. Average spectrum of 32-time measurements was recorded.

Elemental composition was analyzed by Vario EL from Elementar Analysensysteme GmbH to acquire the content of C, H, N in weight

For TEM analyses, drops of dispersion of materials were deposited on 200 mesh copper grids equipped with micro-grids and coated with ultrathin carbon film and dried in air. The morphology of samples were observed by FEI Tecnai G2 T20 TEM operated at 120 kV.

MALDI-TOF MS spectra were acquired by a Bruker Daltonics ultraflex TOF mass spectrometer. Reflection mode was adopted and voltage parameters were set as followed in our work: ion source 1, 25.00 kV; ion source 2, 22.15 kV; lens, 10.55 kV; reflector, 26.30 kV; reflector 2, 14.10 kV. The laser frequency was set on 20 Hz. 1 μL of mixture of materials with peptides adsorbed and matrix solution was directly deposited on a plate and dried in air for MALDI-TOF MS analysis.

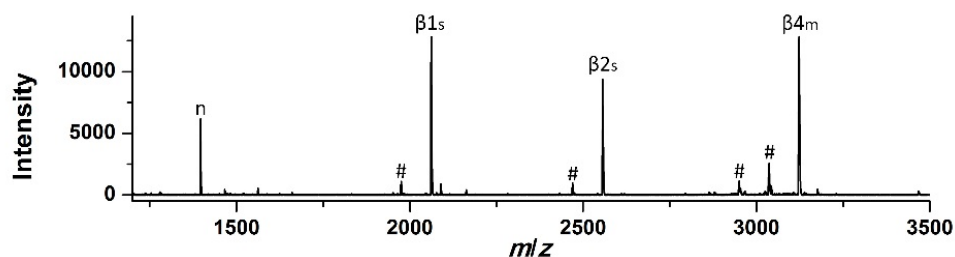
Figure and Table

Figure S1 Enrichment of phosphopeptides from digest of β -casein at 4×10^{-7} M by AFG. Composition of loading buffer is 60% ACN with 0.1 M HAc. (s: monophosphopeptide; m: multiphosphopeptide; #: dephosphorylated residue of phosphopeptides; n: non-phosphorylated peptide).

Table S1 Detailed informations of phosphopeptides enriched from digests of α -casein, β -casein and non-fat milk

No.	<i>m/z</i>	phosphorylation sites	Sequences
α 1	1237.0	1	TVDMEpSTEVF
α 2	1661.0	1	VPQLEIVPNpSAEER
α 3	1927.2	2	DIGpSEpSTEDQAMEDIK
α 4	1943.1	2	DIGpSEpSTEDQAoMEDIK
α 5	1952.3	1	YKVPQLEIVPNpSAEER
α 6	2618.1	4	NTMEHVpSpSpSEESIIpSQETYK
α 7	2635.1	4	NToMEHVpSpSpSEESIIpSQETYK
α 8	2677.3	3	VNELpSKDIGpSEpSTEDQAMEDIK
α 9	2703.3	5	pyroQMEAEpSIpSpSpSEEIVPNpSVEQK
α 10	2719.5	5	QMEAEpSIpSpSpSEEIVPNpSVEQK
α 11	2735.3	5	QoMEAEpSIpSpSpSEEIVPNpSVEQK
α 12	2935.0	3	EKVNELpSKDIGpSEpSTEDQAMEDIK
α 13	3007.8	4	NANEEYpSIGpSpSpSEESAEEVATEEVK
α 14	3087.7	5	NANEEYpSIGpSpSpSEEpSAEVATEEVK
β 1	2061.6	1	FQpSEEQQTEDELQDK
β 2	2555.9	1	FQpSEEQQTEDELQDKIHFP
β 3	2965.8	4	ELEELNVPGEIVEpSLpSpSpSEESITR
β 4	3122.1	4	RELEELNVPGEIVEpSLpSpSpSEESITR