

Supporting information for

**Boronate affinity mesoporous silica nanoparticles based
selective enrichment for highly efficient analysis of
ginsenosides**

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SUPPLEMENTARY DATA

Table S1. Detail information of powder X-Ray diffraction patterns

Material	$d_{100}/\text{\AA}$	$a_0/\text{\AA}$
MSN	34.8	40.1
BA-MSN	34.3	39.5

The d_{100} numbers represent the d-spacing corresponding to the main (100) XRD peak.

The unit-cell size (a_0) is calculated from the d_{100} using the formula $a_0 = 2 d_{100}/3^{1/2}$.

Table S2. Comparison of different separation method for ginsenoside

Method	Separation media	Target	RSD	LOD	Recovery	Ref
HILIC-ELSD	Phenomenex Luna HILIC column	ginsenoside Rb1	3.1%	0.025mg/mL	97.2%	1
2D LC-ESI MS	TSKGEL Amide-80 column & Acquity BEH C18	Ginsenoside Rd	1.8%	1ng/mL	101.6%	2
RP HPLC-PDA	Waters Symmetry C18	ginsenosides Rg1	4%	0.01mg/mL	98.3%	3
GC-MS	HP-1 fused-silica capillary	ginsenoside Rb1	5%	3ng/mL	80%	4
Proposed method	BA-MSNs	Ginsenoside Re	4%	0.5 μ g/mL	85%	-

Reference

1. J. P. Qin, J. Feng, Y. H. Li, K. F. Mo, S. Y. Lu, *J. Pharm. Biomed.*, 2011, **56**, 836.
2. S. Y. Wang, L. Z. Qiao, X. Z. Shi, C. X. Hu, H. W. Kong, G. W. Xu, *Anal. Bioanal. Chem.*, 2015, **407**, 331.
3. A. J. Lau, S. O. Woo, H. L. Koh, *J. Chromatogr. A*, 2003, **1011**, 77.
4. J. F. Cui, I. Björkhem, P. Eneroth, *J Chromatogr B* 1997, **689**, 349.

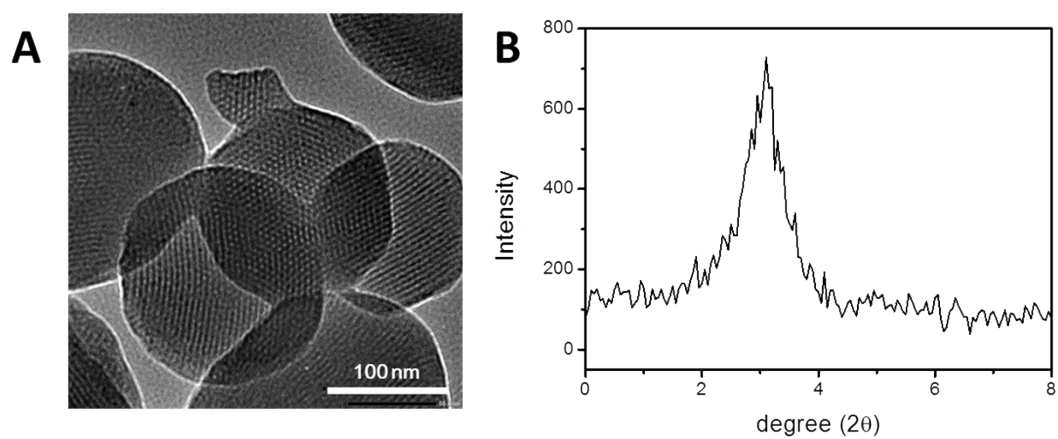


Figure S1. (A) TEM and (B) XRD of mesoporous silica nanoparticles.

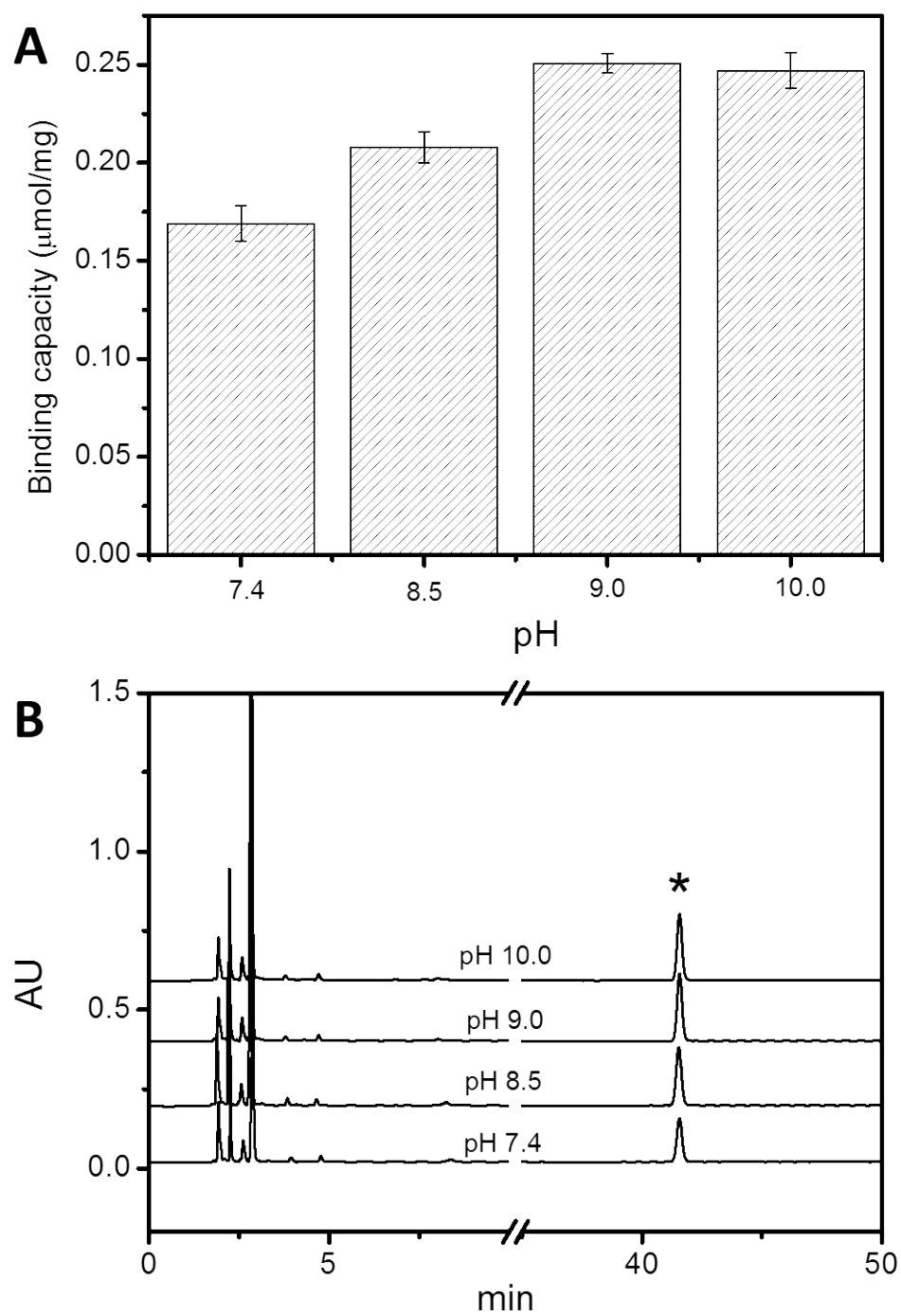


Figure S2. Binding performances of BA-MSNs towards ginsenoside Re at different pH of loading buffer. (A) binding capacity and (B) corresponding chromatograms. * indicates ginsenoside Re.

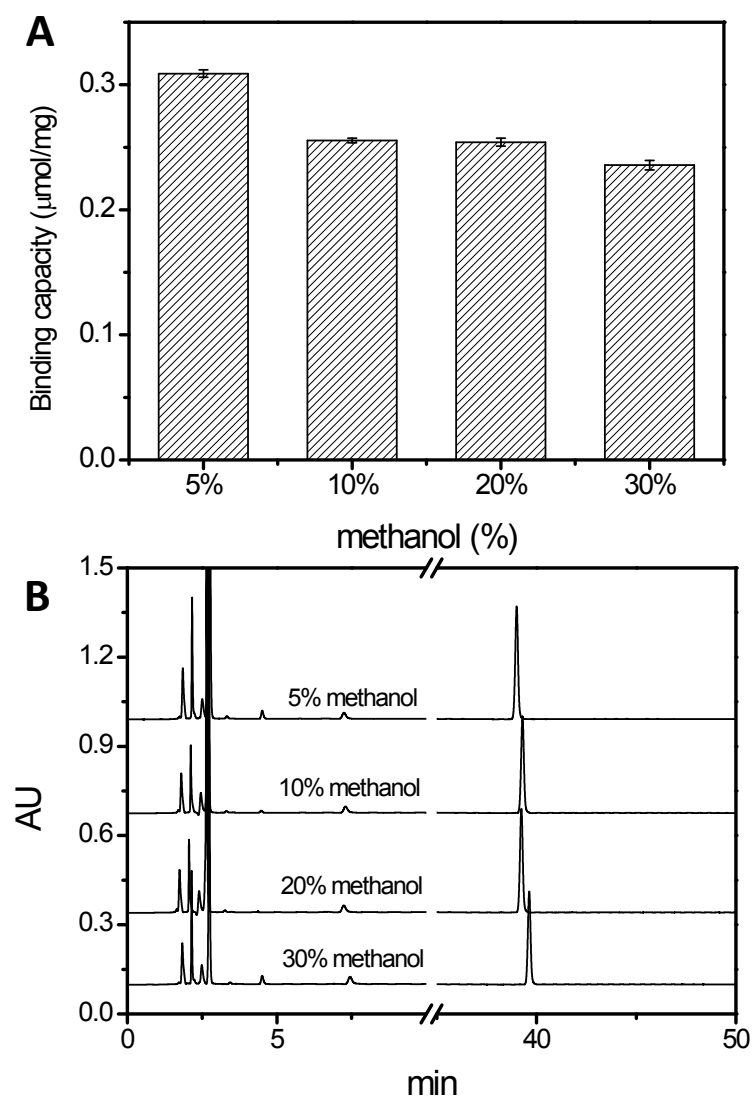


Figure S3. Binding performances of BA-MSNs towards ginsenoside Re at different amount of methanol in loading buffer. (A) binding capacity and (B) corresponding chromatograms. * indicates ginsenoside Re.

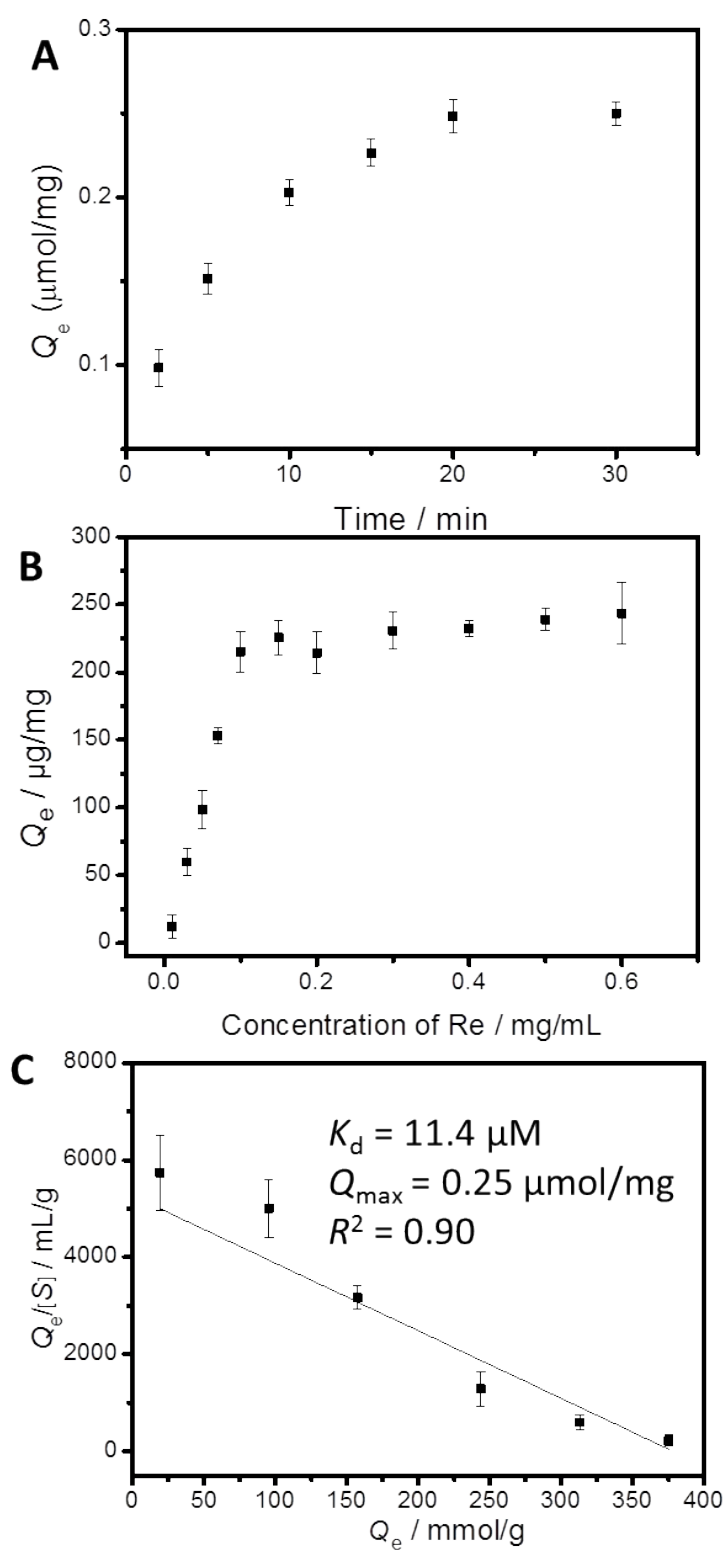


Figure S4. Binding performances of BA-MSNs towards ginesnoside Re (A) binding equilibrium, (B) binding isotherm and (c) corresponding scatchard plots of binding isotherm of BA-MSNs.

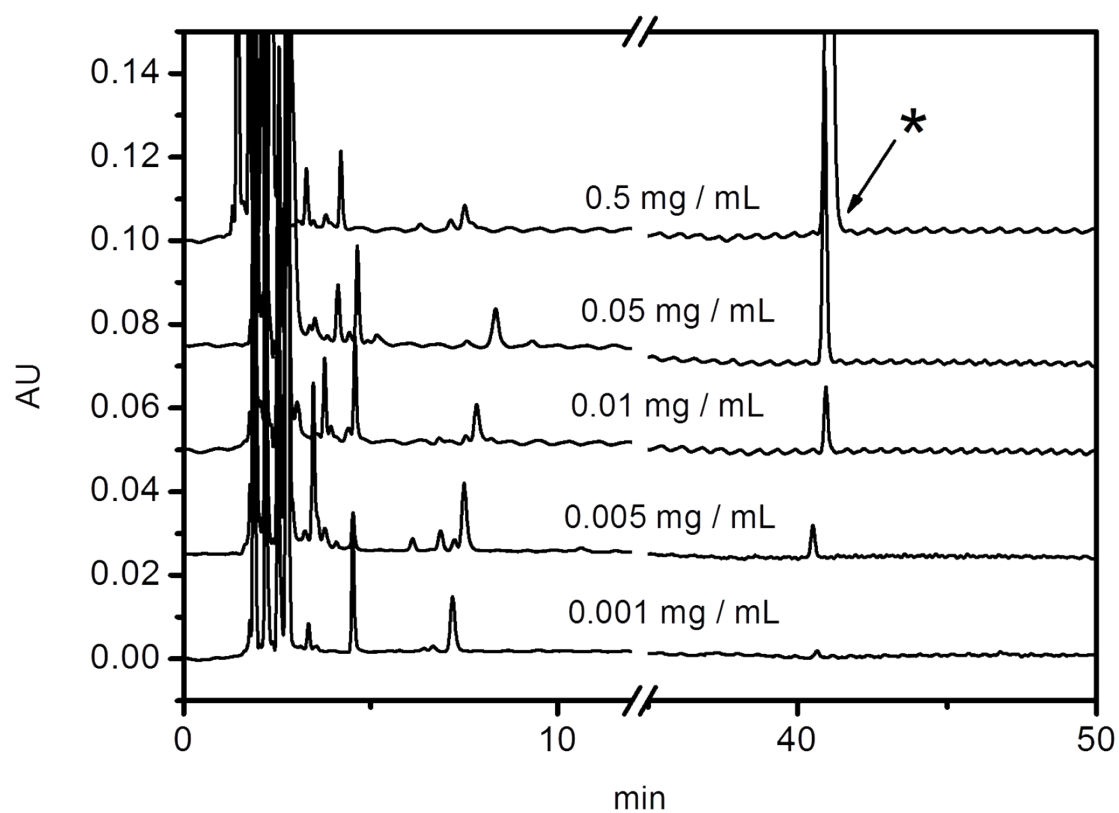


Figure S5. Chromatographic analysis of ginsenoside Re of different concentrations after extracted with BA-MSNs. * indicates ginsenoside Re.

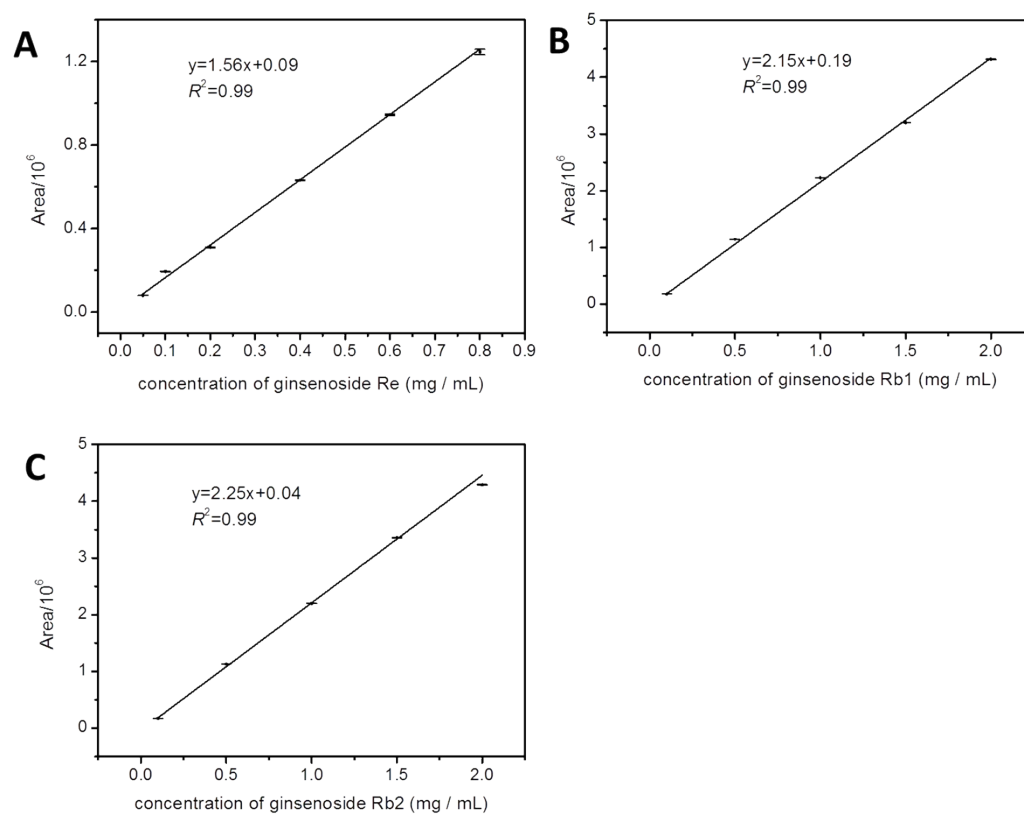


Figure S6. Calibration curve of different ginsenosides (A) Re, (B) Rb1 and (C) Rb2.