

## Supporting Information

### Label-Free Detection of $\beta$ -Amyloid Peptides ( $A\beta$ 40 and $A\beta$ 42): A Colorimetric Sensor Array for Plasma Monitoring of Alzheimer's Disease

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**Fig. S9.** Heat map of the absorbance response patterns for sensor elements against Aβ40, Aβ42 and HSA at concentrations (a) 50, (b) 75, (c) 100, (d) 150, (e) 200, (f) 300, (g) 400, and (h) 500 nmol L<sup>-1</sup> at 3 wavelengths of 420, 530, 620 nm. In the rainbow color scale, a dark red is for the highest positive value and a dark purple is for the highest negative value. 11S

**Table S1.** Leave-one-out analysis by Jackknifing in linear discriminant analysis. Each group contains 8 concentrations of analyte (Aβ40, Aβ42, and HSA) with 6 replicates. 12S

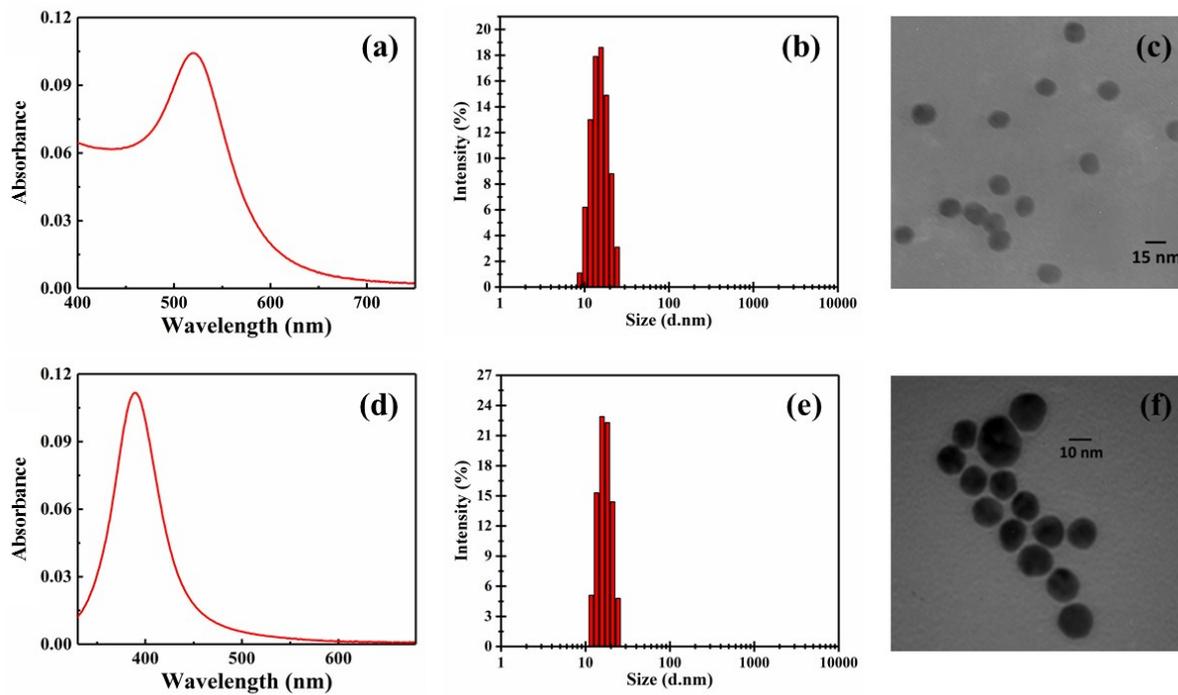
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**Table S2.** Leave-one-out analysis by Jackknifing in linear discriminant analysis. Each group contains one concentration of analyte with 6 replicates. 14S

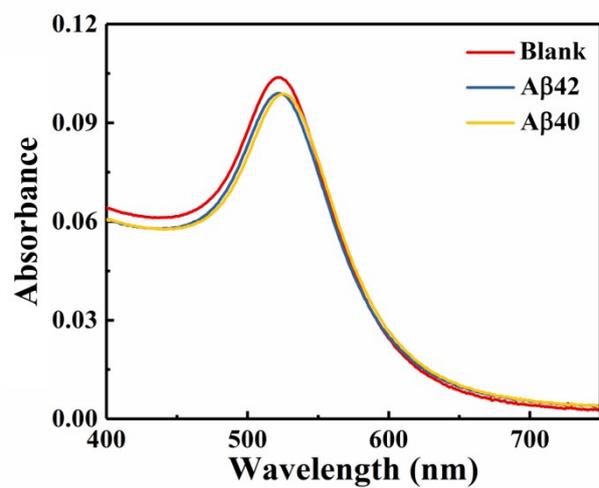
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**Table S3.** Leave-one-out analysis by Jackknifing in linear discriminant analysis. Each group contains analyte or interferences with 6 replicates. 16S

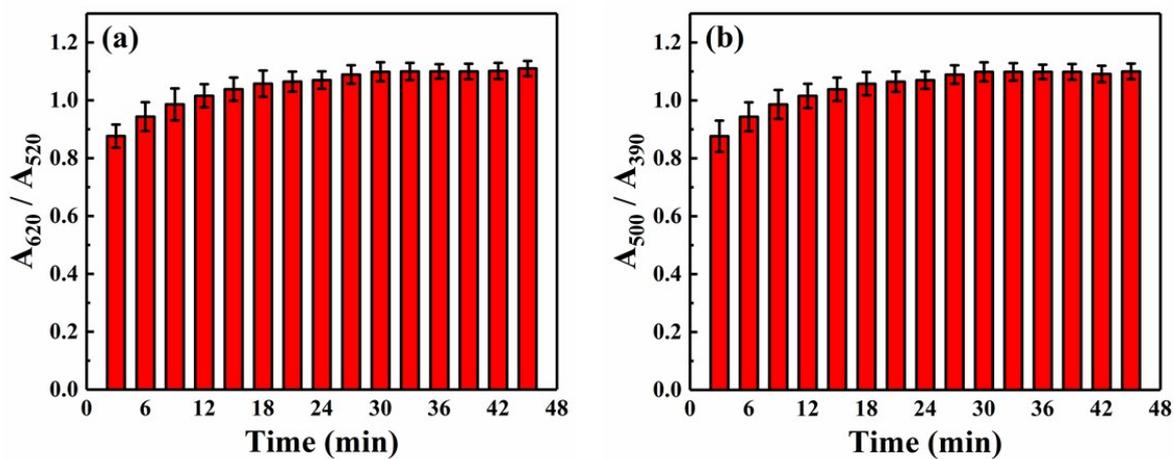
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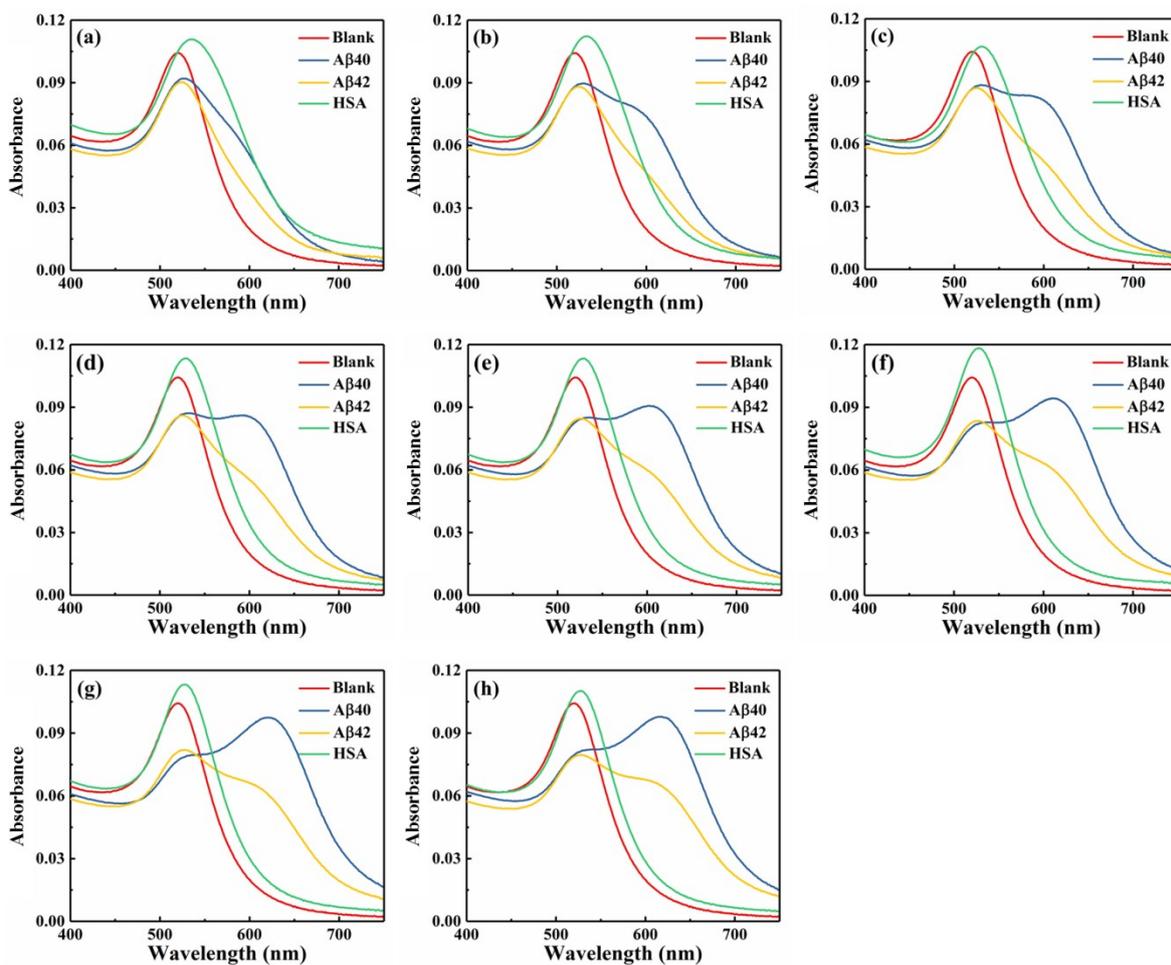
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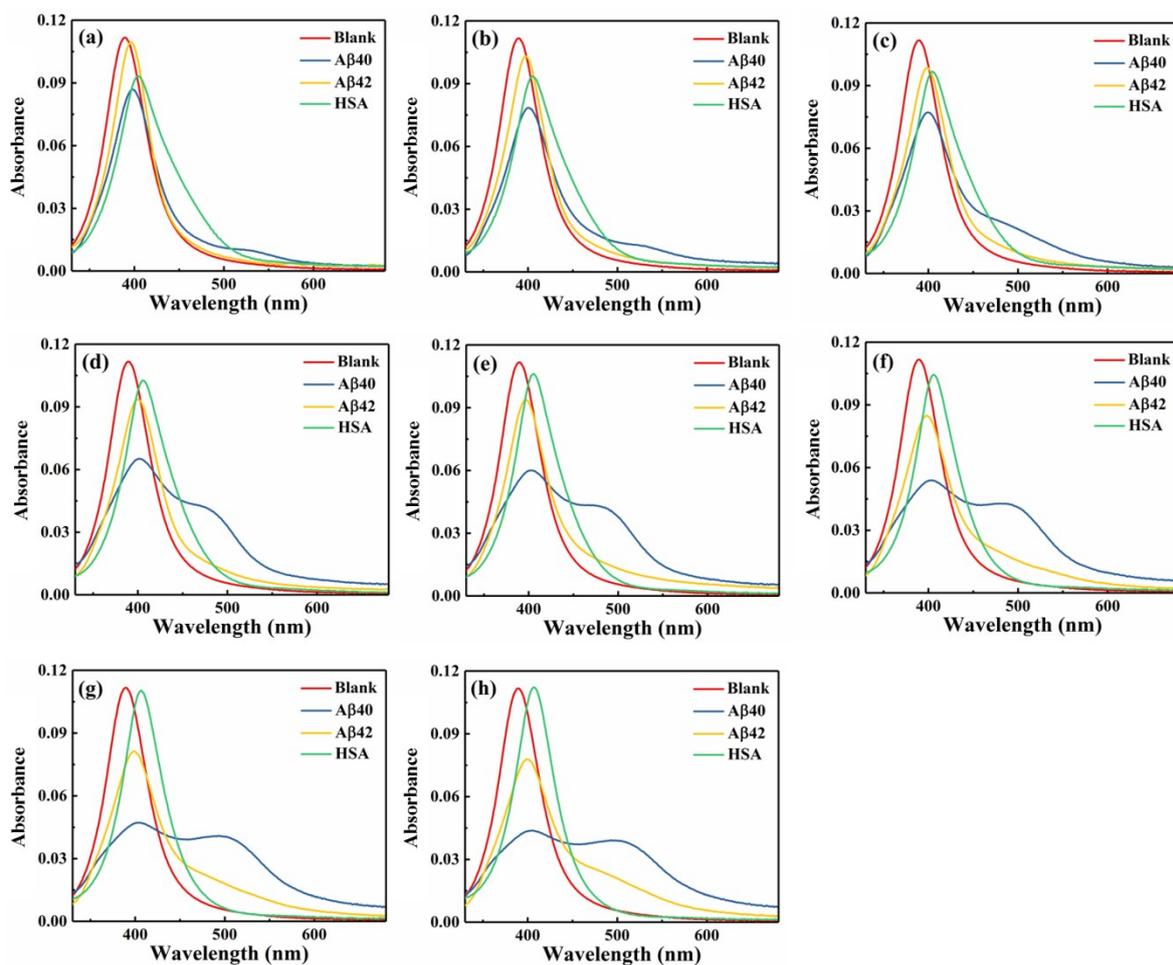
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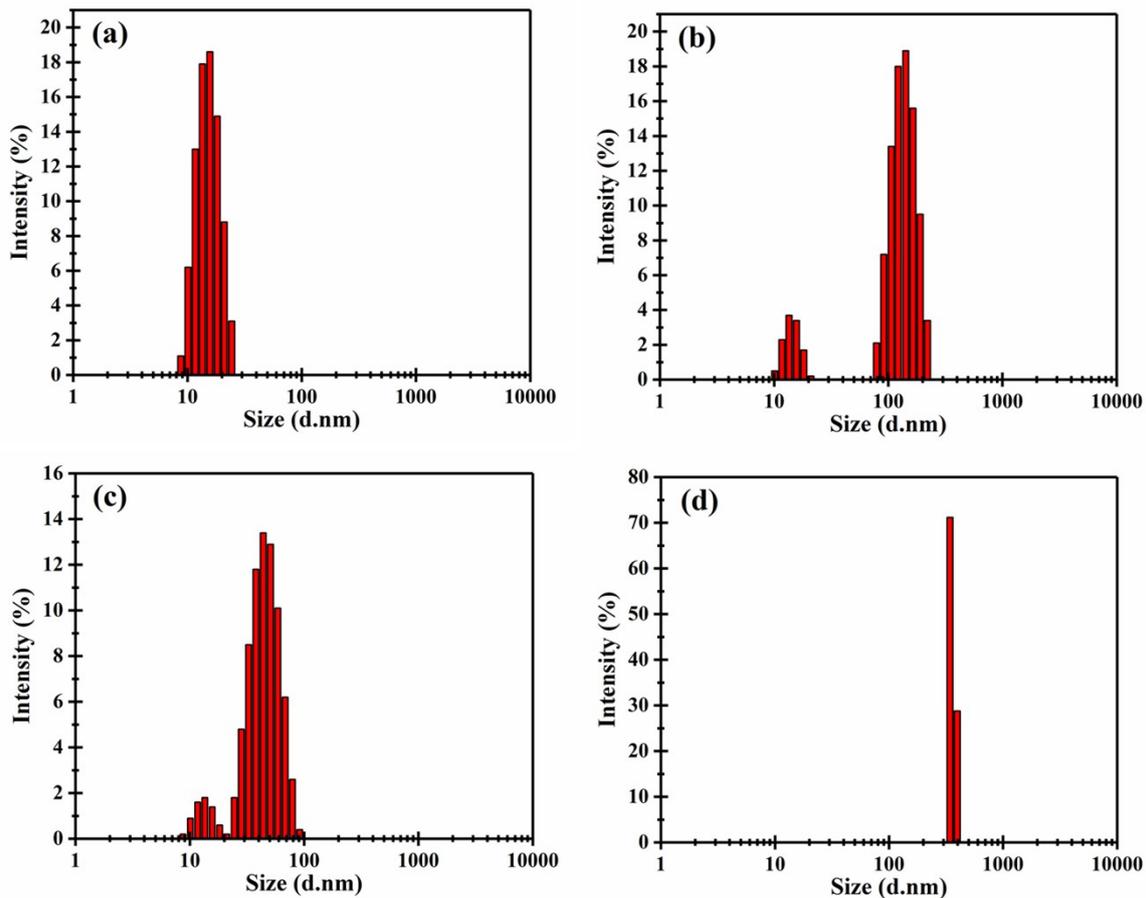
**Fig. S3.** Effect of time on the aggregation process of Aβ40 (300 nmol L<sup>-1</sup>) in the presence of 130 μmol L<sup>-1</sup> of Cu(II), 0.27 nmol L<sup>-1</sup> of (a) AuNPs or (b) AgNPs and pH 5.0.



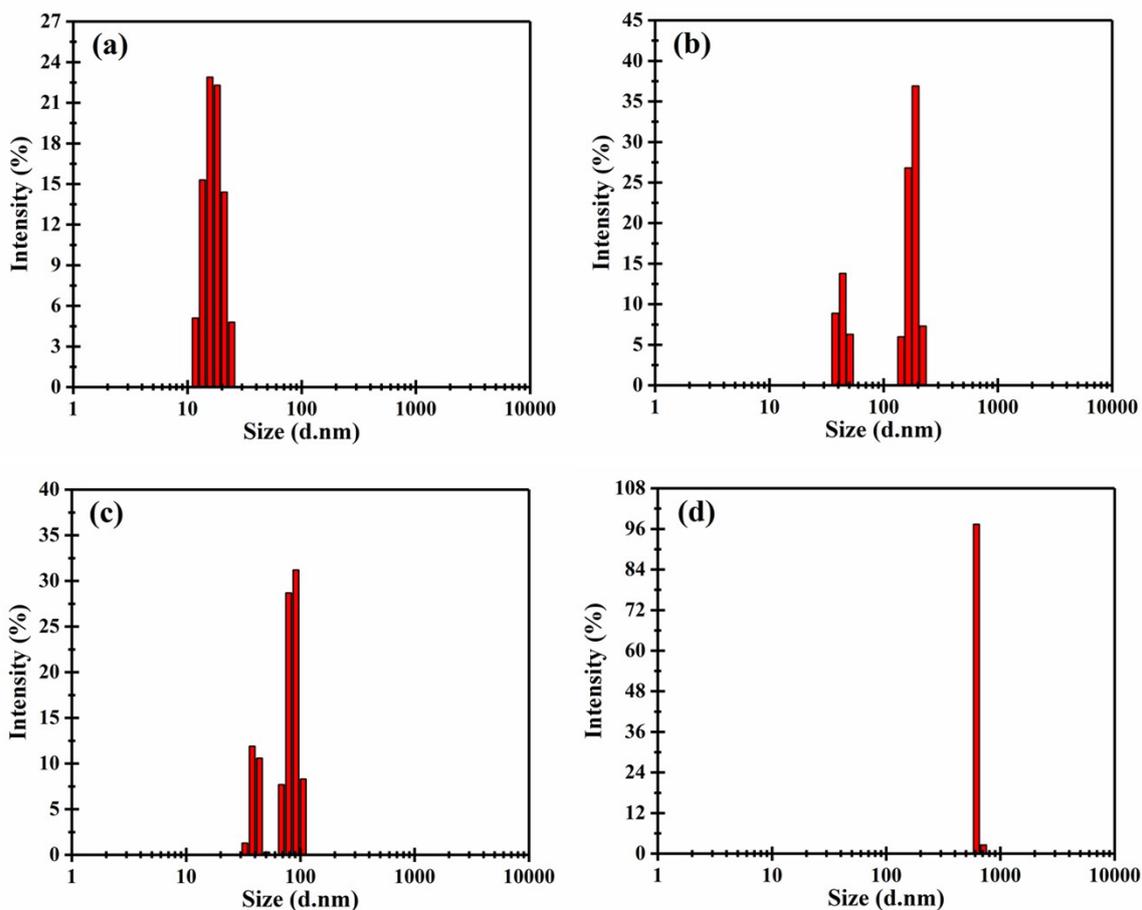
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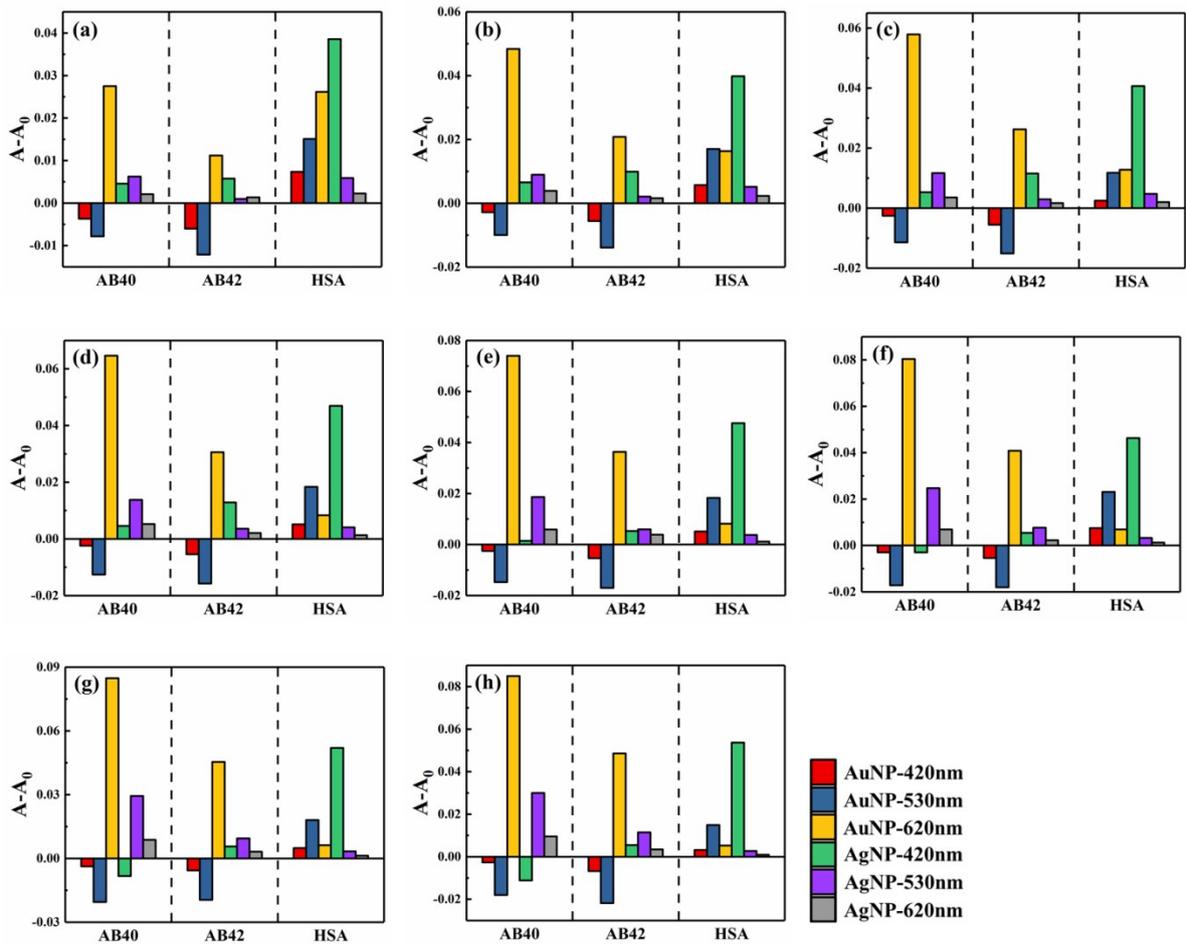
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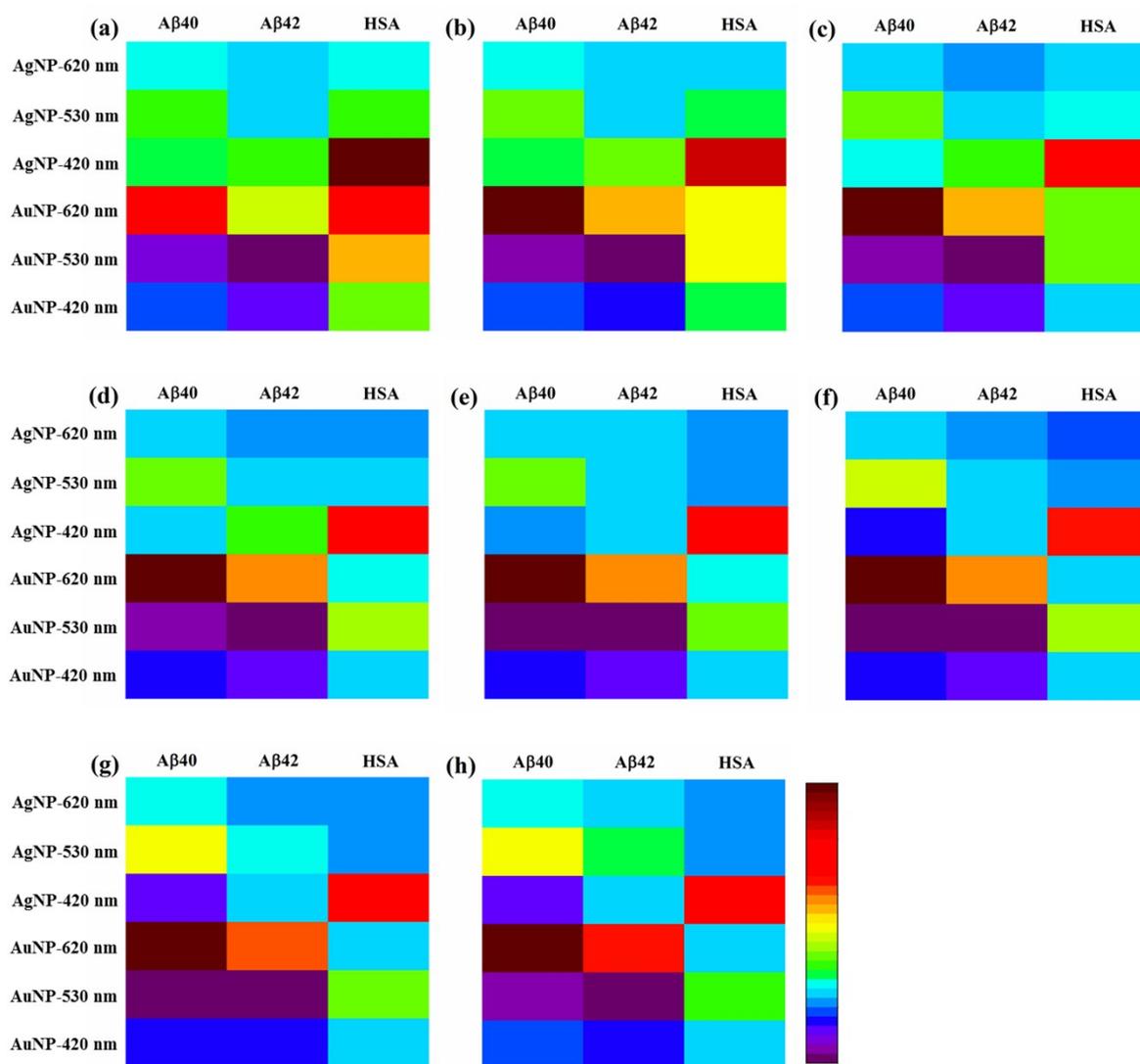
**Fig. S6.** Intensity size distribution of the AuNPs (a), AuNPs (0.27 nmol L<sup>-1</sup>) in the presence of 130 μmol L<sup>-1</sup> Cu(II), pH 5.0 at 20 min and 300 nmol L<sup>-1</sup> of Aβ40 (b), AuNPs (0.27 nmol L<sup>-1</sup>) in the presence of 130 μmol L<sup>-1</sup> Cu(II), pH 5.0 at 20 min and 300 nmol L<sup>-1</sup> of Aβ42 (c) and AuNPs (0.27 nmol L<sup>-1</sup>) in the presence of 130 μmol L<sup>-1</sup> Cu(II), pH 5.0 at 20 min and 300 nmol L<sup>-1</sup> of HAS (d).



**Fig. S7.** Intensity size distribution of the AgNPs (a), AgNPs ( $0.27 \text{ nmol L}^{-1}$ ) in the presence of  $130 \text{ } \mu\text{mol L}^{-1}$  Cu(II), pH 5.0 at 20 min and  $300 \text{ nmol L}^{-1}$  of A $\beta$ 40 (b), AgNPs ( $0.27 \text{ nmol L}^{-1}$ ) in the presence of  $130 \text{ } \mu\text{mol L}^{-1}$  Cu(II), pH 5.0 at 20 min and  $300 \text{ nmol L}^{-1}$  of A $\beta$ 42 (c) and AgNPs ( $0.27 \text{ nmol L}^{-1}$ ) in the presence of  $130 \text{ } \mu\text{mol L}^{-1}$  Cu(II), pH 5.0 at 20 min and  $300 \text{ nmol L}^{-1}$  of HAS (d).



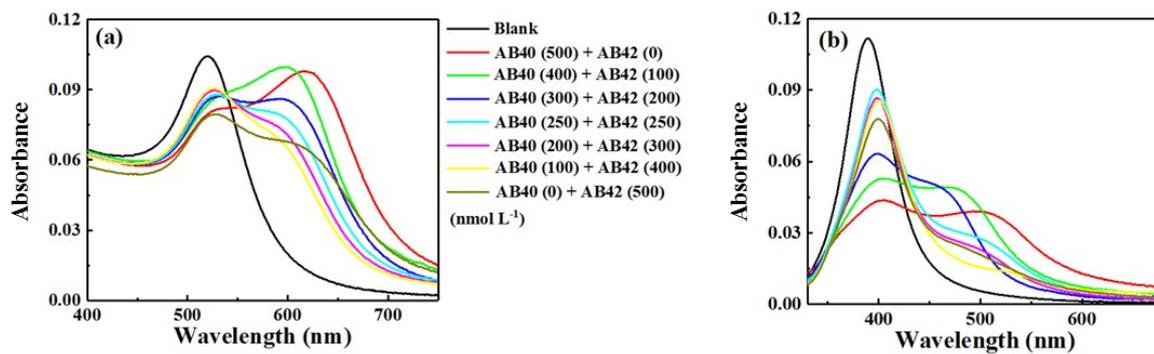
**Fig. S8.** Absorbance response patterns of sensor elements against Aβ40, Aβ42 and HSA at concentrations (a) 50, (b) 75, (c) 100, (d) 150, (e) 200, (f) 300, (g) 400, and (h) 500 nmol L<sup>-1</sup> at 3 wavelengths of 420, 530, 620 nm.



**Fig. S9.** Heat map of the absorbance response patterns for sensor elements against A $\beta$ 40, A $\beta$ 42 and HSA at concentrations (a) 50, (b) 75, (c) 100, (d) 150, (e) 200, (f) 300, (g) 400, and (h) 500 nmol L<sup>-1</sup> at 3 wavelengths of 420, 530, 620 nm. In the rainbow color scale, a dark red is for the highest positive value and a dark purple is for the highest negative value.

**Table S1.** Leave-one-out analysis by Jackknifing in linear discriminant analysis. Each group contains 8 concentrations of analyte (A $\beta$ 40, A $\beta$ 42, and HSA) with 6 replicates.

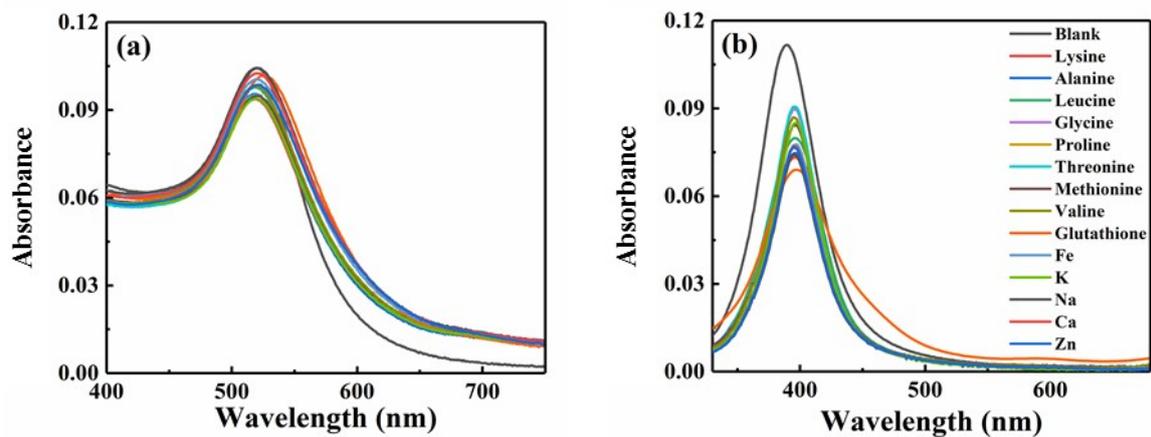
	Predicted Group			
	A $\beta$ 40	A $\beta$ 42	HSA	Total
A $\beta$ 40	48	0	0	48
	100.00%	0.00%	0.00%	100.00%
A $\beta$ 42	0	48	0	48
	0.00%	100.00%	0.00%	100.00%
HSA	0	0	48	48
	0.00%	0.00%	100.00%	100.00%
Total	48	48	48	144
	33.33%	33.33%	33.33%	100.00%



**Fig. S10.** Absorbance spectra of (a) AuNPs and (b) AgNPs ( $0.27 \text{ nmol L}^{-1}$  in the presence of  $130 \text{ } \mu\text{mol L}^{-1}$  Cu(II), pH 5.0 before and after exposure to total  $500 \text{ nmol L}^{-1}$  of A $\beta$ 40, A $\beta$ 42 and their mixtures at 20 min.

**Table S2.** Leave-one-out analysis by Jackknifing in linear discriminant analysis. Each group contains one concentration of analyte with 6 replicates.

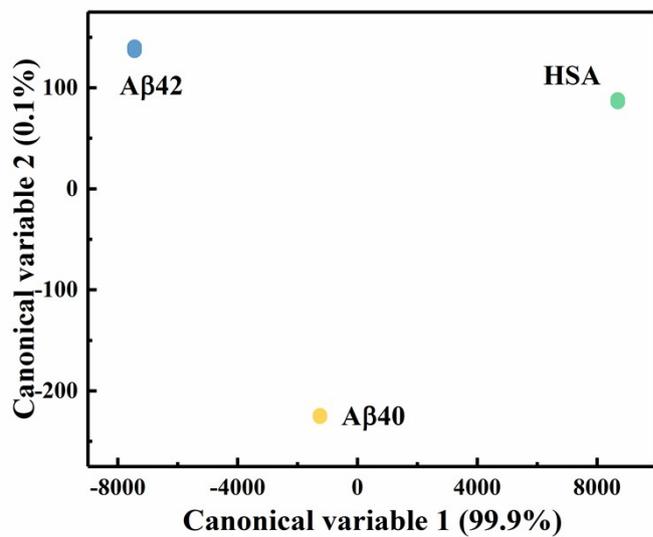
	Predicted Group							Total
	1	2	3	4	5	6	7	
1	6	0	0	0	0	0	0	6
	100.00%	0.00%	0.00%	0.00%	0.00%	0.00%	0.00%	100.00%
2	0	6	0	0	0	0	0	6
	0.00%	100.00%	0.00%	0.00%	0.00%	0.00%	0.00%	100.00%
3	0	0	6	0	0	0	0	6
	0.00%	0.00%	100.00%	0.00%	0.00%	0.00%	0.00%	100.00%
4	0	0	0	6	0	0	0	6
	0.00%	0.00%	0.00%	100.00%	0.00%	0.00%	0.00%	100.00%
5	0	0	0	0	6	0	0	6
	0.00%	0.00%	0.00%	0.00%	100.00%	0.00%	0.00%	100.00%
6	0	0	0	0	0	6	0	6
	0.00%	0.00%	0.00%	0.00%	0.00%	100.00%	0.00%	100.00%
7	0	0	0	0	0	0	6	6
	0.00%	0.00%	0.00%	0.00%	0.00%	0.00%	100.00%	100.00%
Total	6	6	6	6	6	6	6	42
	14.29%	14.29%	14.29%	14.29%	14.29%	14.29%	14.29%	100.00%



**Fig. S11.** Absorbance spectra of (a) AuNPs and (b) AgNPs ( $0.27 \text{ nmol L}^{-1}$  in the presence of  $130 \text{ } \mu\text{mol L}^{-1}$  Cu(II), pH 5.0 before and after exposure to the lysine, alanine, leucine, glycine, proline, threonine, methionine, valine, glutathione, FeCl<sub>3</sub>, KCl, NaCl, CaCl<sub>2</sub>, and ZnCl<sub>2</sub>, as interferences.

**Table S3.** Leave-one-out analysis by Jackknifing in linear discriminant analysis. Each group contains analyte or interferences with 6 replicates.

	Predicted Group				
	A $\beta$ 40	A $\beta$ 42	HSA	Interferences	Total
A $\beta$ 40	6	0	0	0	6
	100.00%	0.00%	0.00%	0.00%	100.00%
A $\beta$ 42	0	6	0	0	6
	0.00%	100.00%	0.00%	0.00%	100.00%
HSA	0	0	6	0	6
	0.00%	0.00%	100.00%	0.00%	100.00%
Interferences	0	0	0	84	84
	0.00%	0.00%	0.00%	100.00%	100.00%
Total	6	6	6	84	102
	5.88%	5.88%	5.88%	82.35%	100.00%



**Fig. S12.** Two-dimensional LDA plot clustering Aβ40, Aβ42, and HSA. Plasma was spiked with Aβ40, Aβ42 and HAS at concentrations of 500 nmol L<sup>-1</sup>.