Supplementary Information

Ascorbic acid-loaded apoferritin assisted carbon dots-MnO₂ nanocomposites for selective and sensitive detection of trypsin

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Experimental Procedure

Preparation of APOAA

In particular, APO was dispersed in 2 mL of 10 mM PBS (pH 7.4) and the solution was adjusted to pH 2.0 with 0.1 M HCl solution under continuous stirring. The solution was further stirred for 30 min to make the APO completely disassociated. Then 100 μL of 1.5 M AA solution was added into APO solution dropwise with stirring to allow AA to diffuse around the subunits of the APO. After that, the pH of solution was adjusted to 8.5 with 0.1 M NaOH. The obtained mixture was stirred for another 3 h at room temperature to allow AA enters the APO core. Subsequently, to remove free AA out of APO, the mixture solution was dialyzed against 10 mM PBS (pH 7.8) for 48 h using MD10 dialysis bag with a molecular weight cutoff (MWCO) of 8000-14000. The obtained APOAA solution was stored in amber glass bottles at 4 °C for further use.

Preparation of MnO₂ nanosheets

MnCl₂ aqueous solution and tetramethylammonium hydroxide (TMAOH) containing 3 wt% of H₂O₂ were mixed quickly and then stirred for 12 h at room temperature. The dried aggregates were washed with methanol and water several times, and following dried at 45 °C. Then the product dispersing in water and following sonicated for 12 h to get the resulting suspension (brown colloid).

Figures and Tables

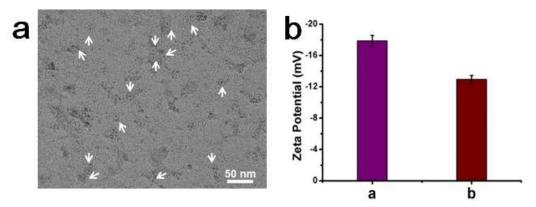


Figure S1. (a) TEM images of APOAA and (b) Zeta Potential of APO (column a) and APOAA (column b)

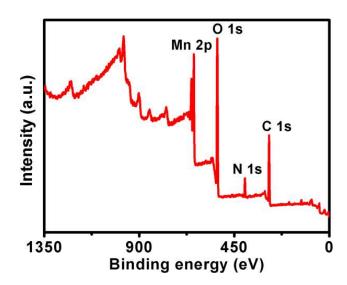


Figure S2. XPS survey spectrum of the CDs-MnO₂

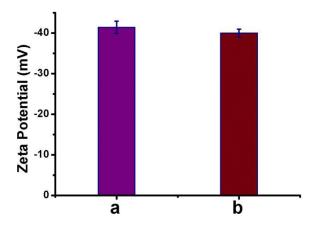


Figure S3. Zeta Potential of MnO₂ synthesized newly (column a) and stored for 6 months (column b)

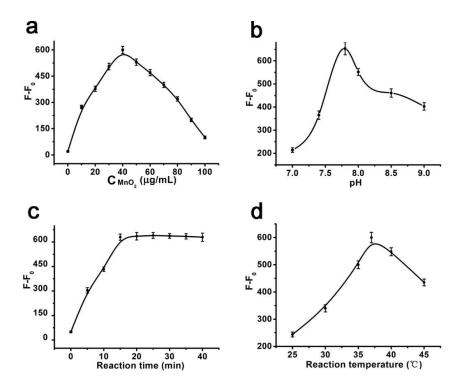


Figure S4. Dependence of the fluorescence recovery value of CMA on (a) MnO_2 nanosheets concentration (b) reaction pH (c) reaction time (d) reaction temperature (the concentration of trypsin is 300 ng/mL). The error bars were derived from the standard deviation of three measurements. Error bar = SD (n = 3). F_0 and F are the fluorescence intensities of CMA in the absence and presence of trypsin, respectively.

Table S1 Comparison of this method with the newly reported approaches for trypsin determination.

Materials	Methods	Linear range (ng/mL)	LOD (ng/mL)	Ref.
		(IIg/IIIL)	(ng/mz)	
Cytochrome c	Chemiluminescence	0-100	0.7	[1]
CdTe QDs	Photoelectrochemial	30-450	2.7	[2]
Cytochrome c-Heme-TMB	Colorimetric	5-2000	4.5	[3]
Au nanoclusters	Colorimetric	$9 \times 10^2 - 1 \times 10^6$	600	[4]
Au nanoclusters	Fluorescence	$1 \times 10^4 - 1 \times 10^8$	2	[5]
GFP	Fluorescence	0-400	0.0282	[6]
CDs-MnO ₂	Fluorescence	1-500	0.3411	This work

GFP: green fluorescent protein

Table S2 Recoveries results of the determination of trypsin in serum samples.

Commle	Added	Found	Recovery
Sample	(ng/mL)	(ng/mL)	(%)
Sample 1	0	0	-
Sample 2	50	48 ± 0.5	96
Sample 3	100	103±0.4	103
Sample 4	200	196±0.2	98
Sample 5	350	355±0.3	101.4
Sample 6	450	446±0.4	99.1

^a Mean value of three independent measurements.

References

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