

Supporting Information

Template-free assembly of Cu,N-codoped hollow carbon nanospheres as low-cost and highly efficient peroxidase nanozymes

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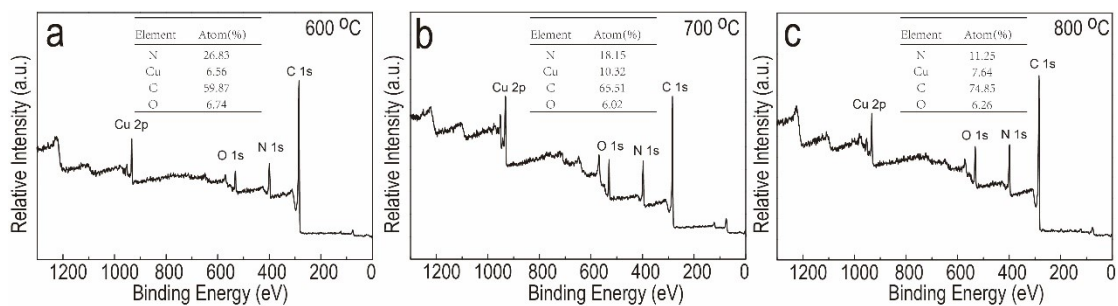


Fig. S1. XPS survey scans of CuNHfCN calcinated at 600 °C, 700 °C and 800 °C. Insets: contents of C, N, O and Cu.

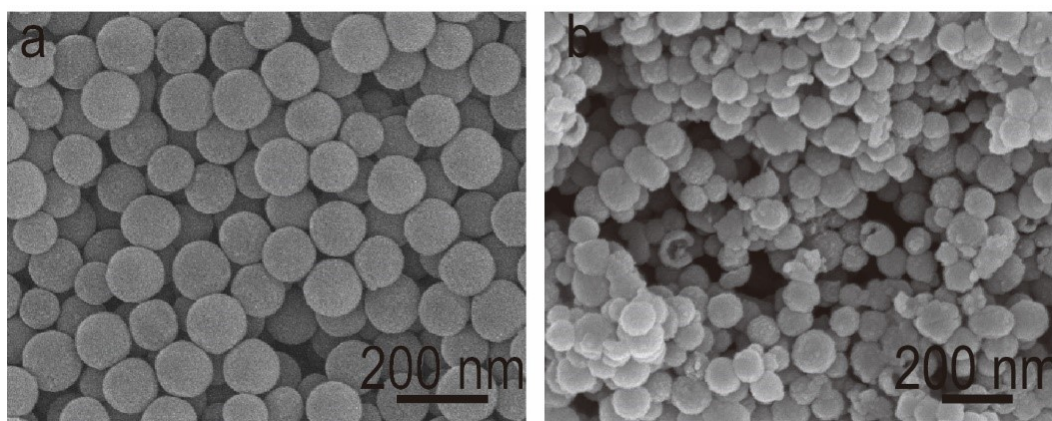


Fig. S2. SEM images of the (a) hollow Cu-PmPD nanospheres and (b) CuNHfCN.

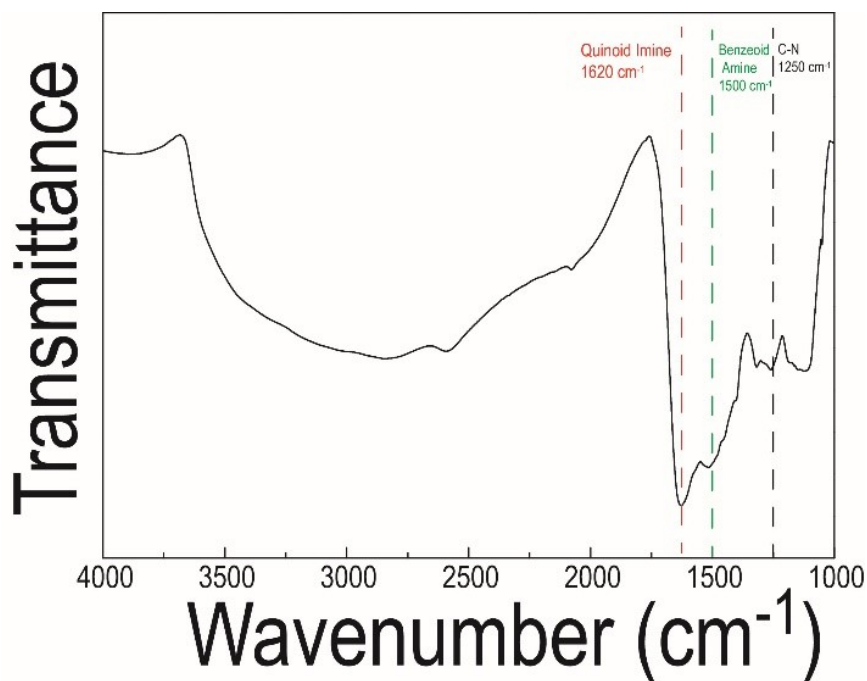


Fig. S3. FT-IR spectrum of the hollow Cu-PmPD nanospheres

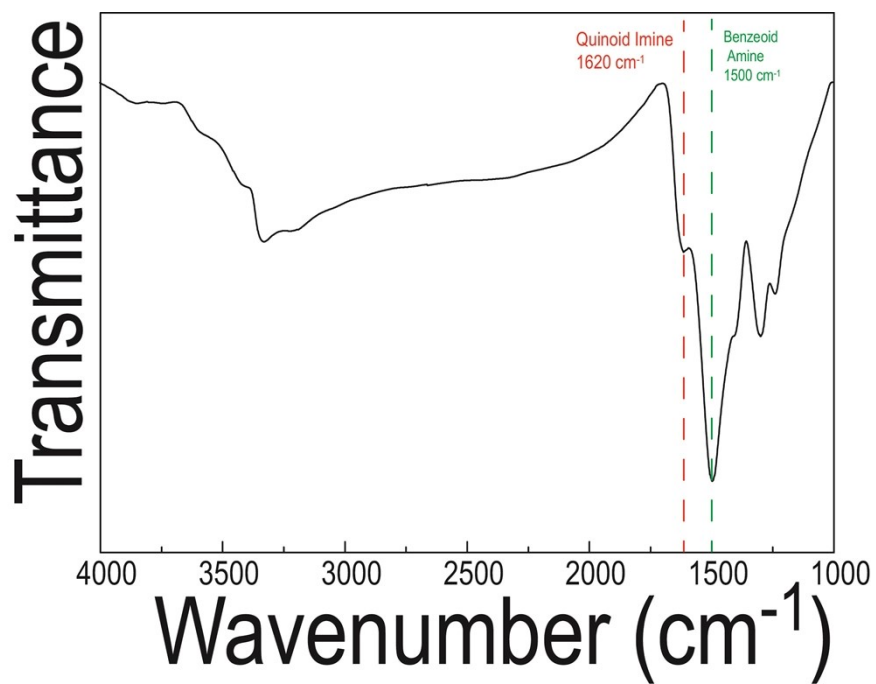


Fig. S4. FT-IR spectrum of the interior parts of the Cu-PmPD nanospheres (the interior parts was obtained according to the literature^{1,2}. Briefly, the interior parts was dissolved in the reaction solvent, which could be obtained through centrifugation and freeze drying)

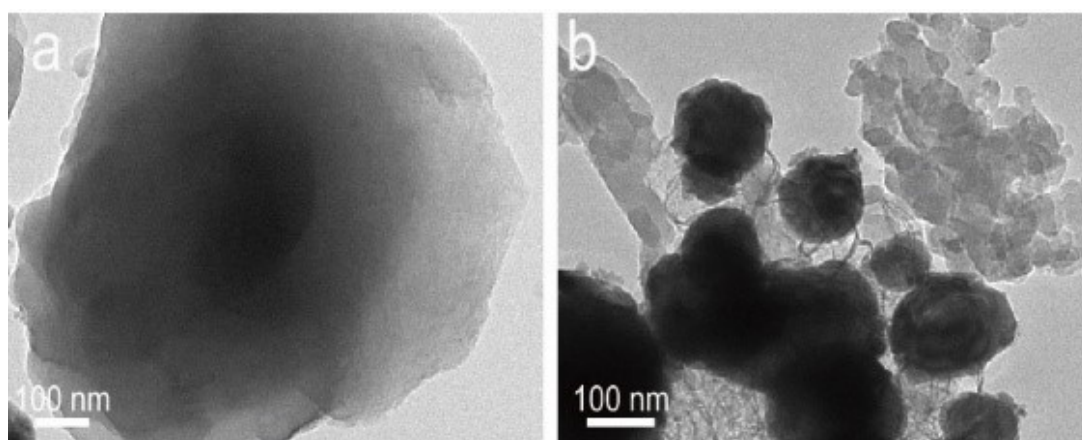


Fig. S5. TEM images of the (a) NC and (b)CuN

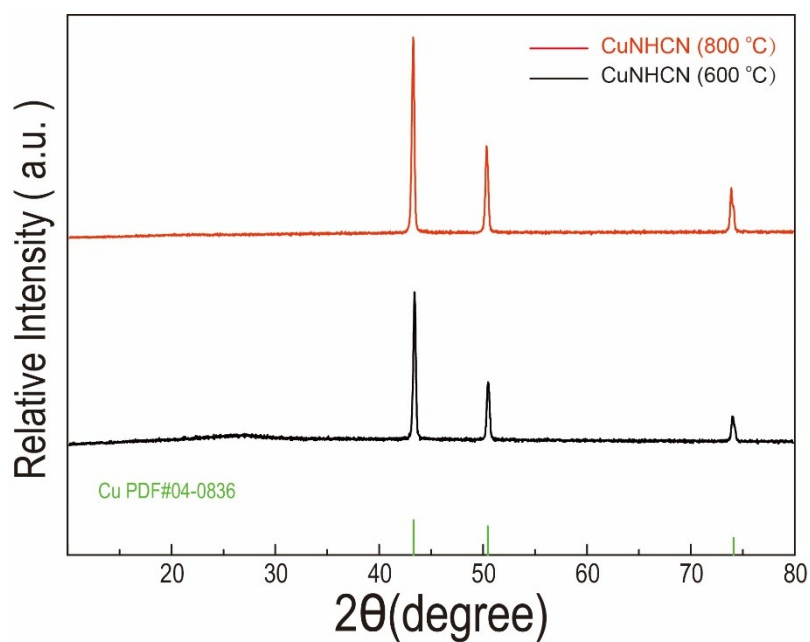


Fig. S6. N₂ adsorption-desorption isotherm curves and pore size distribution curves of (a) CuC, (b) NC, (c) CuNHCN.

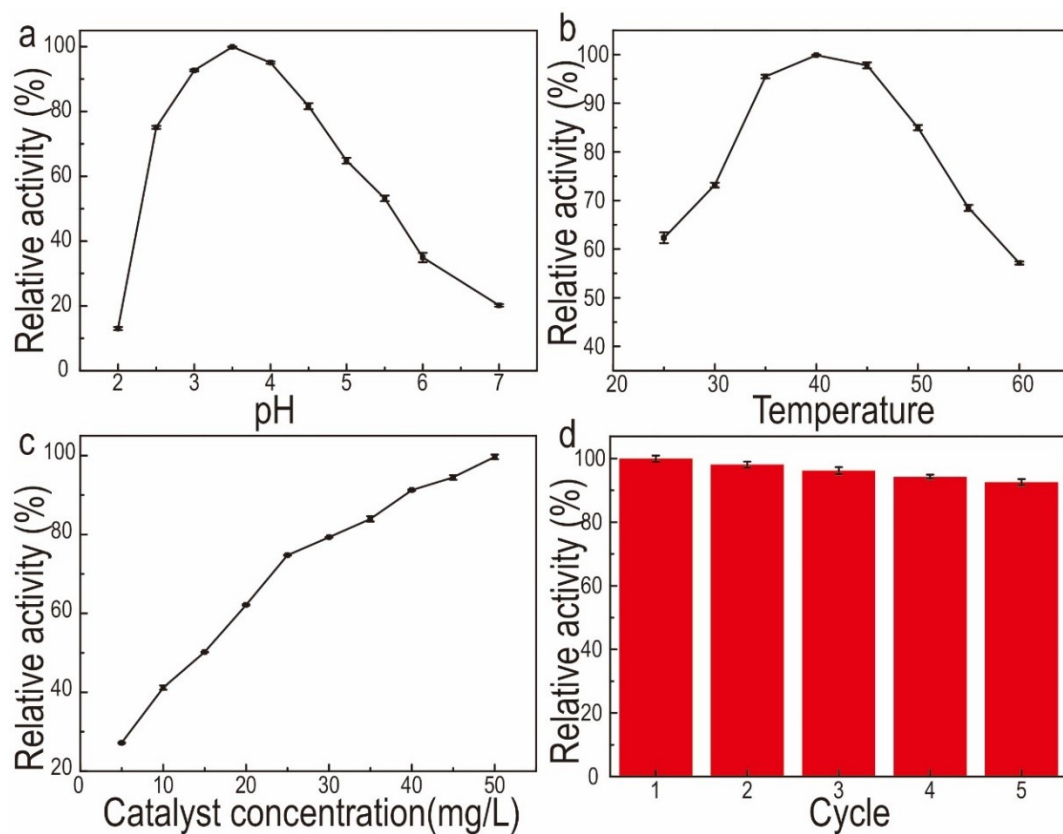


Fig. S7. Effects of (a) pH, (b) temperature, (c) concentration and (d) stability of CuNHCN under optimized reaction conditions. Error bar shows the standard deviation of three independent measurements.

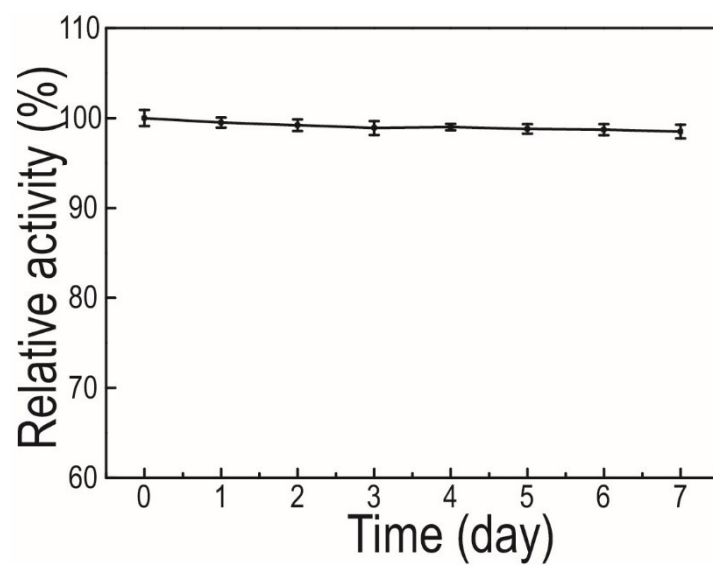


Fig. S8 The storage performance of CuNHCN under optimized reaction conditions

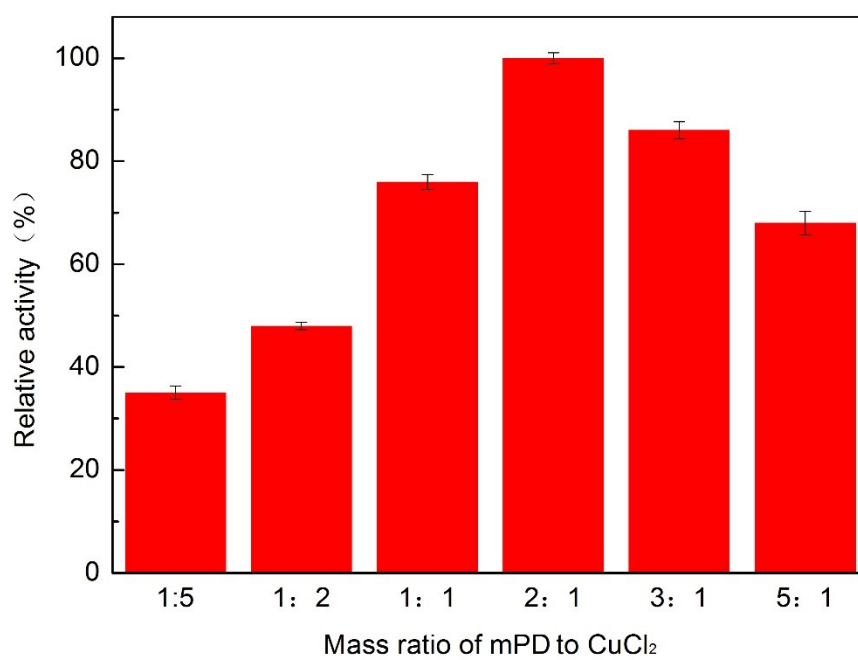


Fig.S9. Influence of the mass ratio of mPD to CuCl₂ on the peroxidase-like catalytic performance

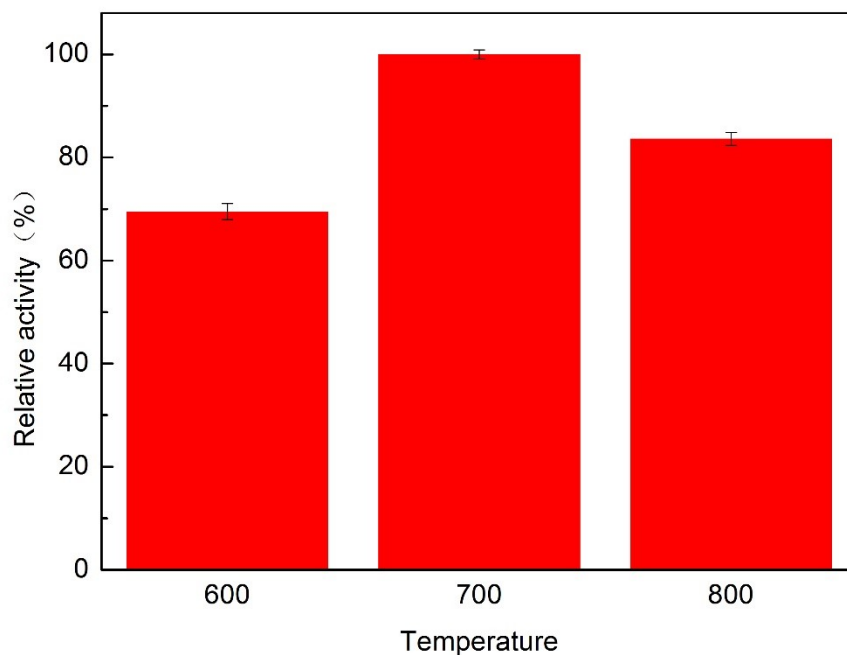


Fig.S10. Influence of the temperature on the peroxidase-like catalytic performance of CuNH₂CN

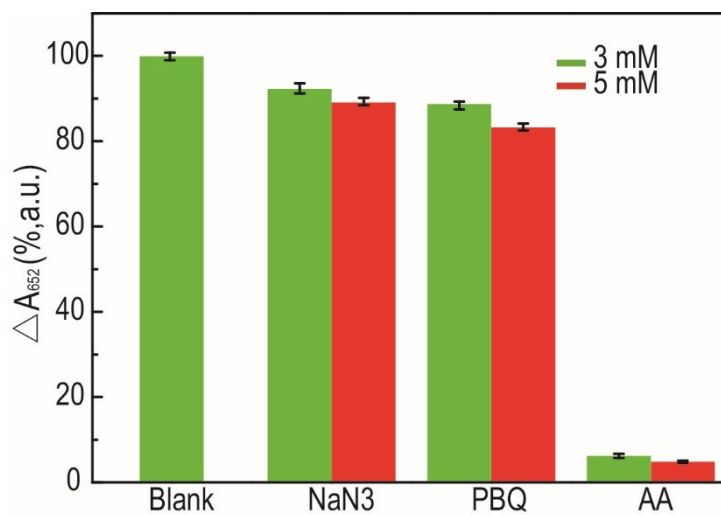


Fig. S11. The influence of different concentrations of ROS scavengers (NaN₃, PBQ and AA) on the A₆₅₂ of the CuNH₂CN/TMB/H₂O₂ system. Reaction condition: 0.15 mM TMB, 0.1 mM H₂O₂, 25 mg L⁻¹ catalyst, pH 3.5 acetate buffer(0.2 M), 4 min reaction at 40 °C.

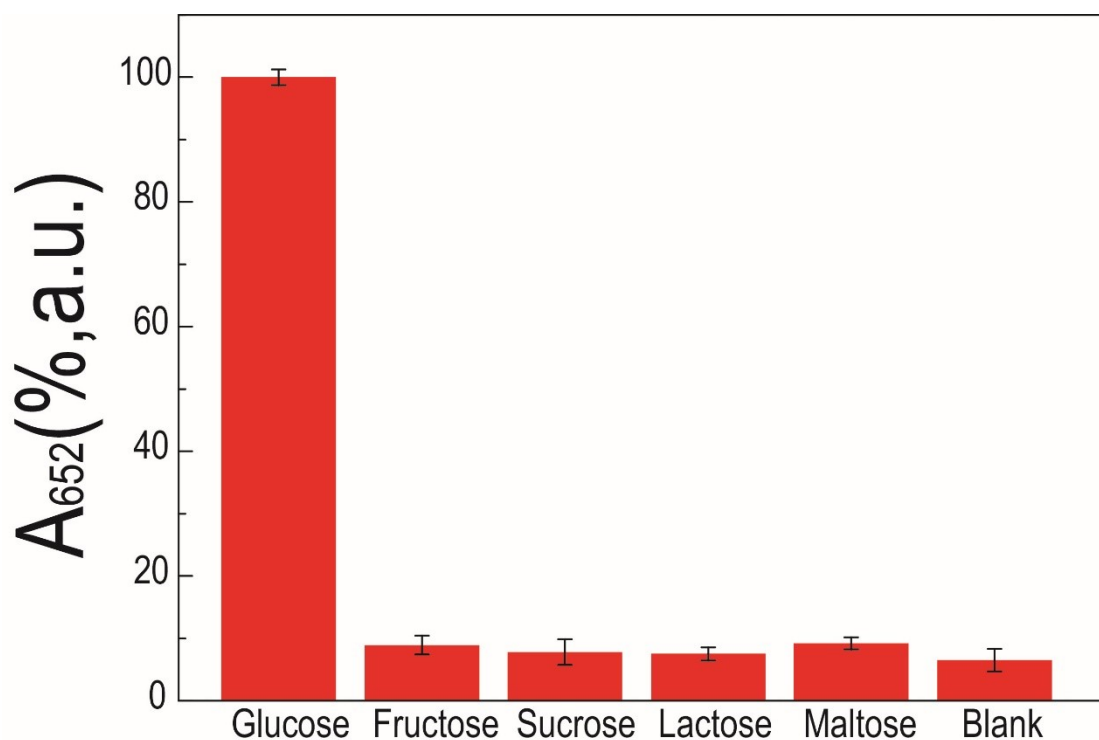


Fig. S12. The influence of different substrates (5 mM fructose, 5 mM sucrose, 5 mM lactose, 5 mM Maltose or 0.5 mM glucose) on the A_{652} of the glucose oxidase/TMB/CuNH₂CN system. Error bar shows the standard deviation of three independent measurements.

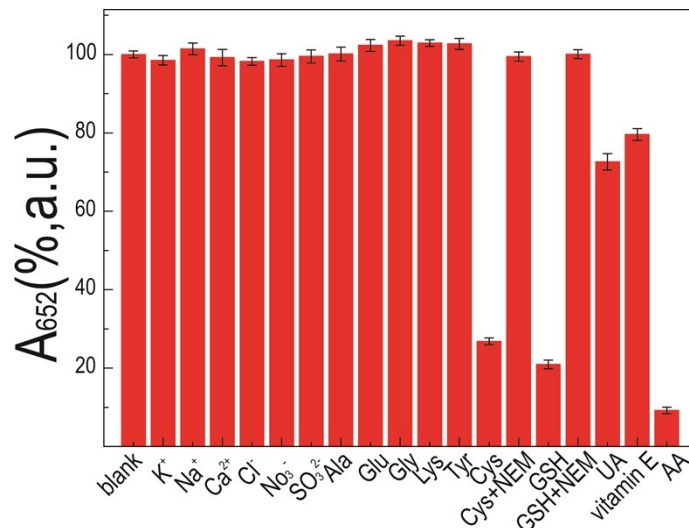


Fig. S13. The influence of different substrates (metal ion(K⁺, Na⁺,Ca²⁺) (5 mM), sulfite (5 mM) , nitrite (0.5 mM), chloride(0.5 mM), alanine (Ala) (5 mM), glutamic acid (Glu) (5 mM), glycine (Gly) (5 mM), lysine (Lys) (5 mM), tyrosine (Tyr) (5 mM),Cysteine (Cys) (0.5 mM), glutathione (GSH) (0.5 mM), Cys+NEM, GSH+NEM, uric acid (UA) (0.5 mM), vitamin E (0.5 mM) and AA (0.5 mM)) on the A_{652} of the H₂O₂/TMB/CuNH₂CN system. Error bar shows the standard deviation of three independent measurements.

Table S1 Comparison of the CuNHCN based on colorimetry method for the determination of glucose and AA.

Material	K_m (mM)		V_{max} (10^{-8} M/s)		Refs
	TMB	H ₂ O ₂	TMB	H ₂ O ₂	
Au NPs /Graphene-	0.38	26.4	18.3	15.41	[3]
Au NPs/PVP-GNs	2.63	104	13.04	11.98	[3]
Cu/Au/Pt TNPs	0.15	2.34	7.33	136.5	[4]
β -CD-Cu-NCs	0.543	32.87	43.4	45.2	[4]
Hollow CuS nanocubes	1.62	0.94	16.64	2.55	[6]
Au loaded nanoporous Fe ₂ O ₃ nanozymes	0.049	138.5	5.882	4.770	[7]
Pt hollow nanodendrites	0.81	6.9	1.2	9.9	[8]
Fe ₃ O with aptamer	0.088	30.24	0.406	0.342	[9]
Au-Ni/g-C ₃ N ₄	0.16	4.47	2.34	6.16	[10]
Co ₃ O ₄ nanocrystals	0.49	1.90	16	12.7	[11]
HRP	0.434	3.70	10.0	8.71	[12]
CuNHCN	0.0655	0.918	5.85	42.84	This work

Reference

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