

Supporting Information

Table S1 Results of spiked studies in human serum samples

| | Spiked (nM) | Calculated (nM) | Recovery (%) |
|-----|-------------|-----------------|--------------|
| Cys | 50 | 54.7±3.9 | 109.4 |
| | 100 | 97.2±2.6 | 97.2 |
| Hcy | 50 | 52.1±4.1 | 104.2 |
| | 100 | 98.9±2.8 | 98.9 |

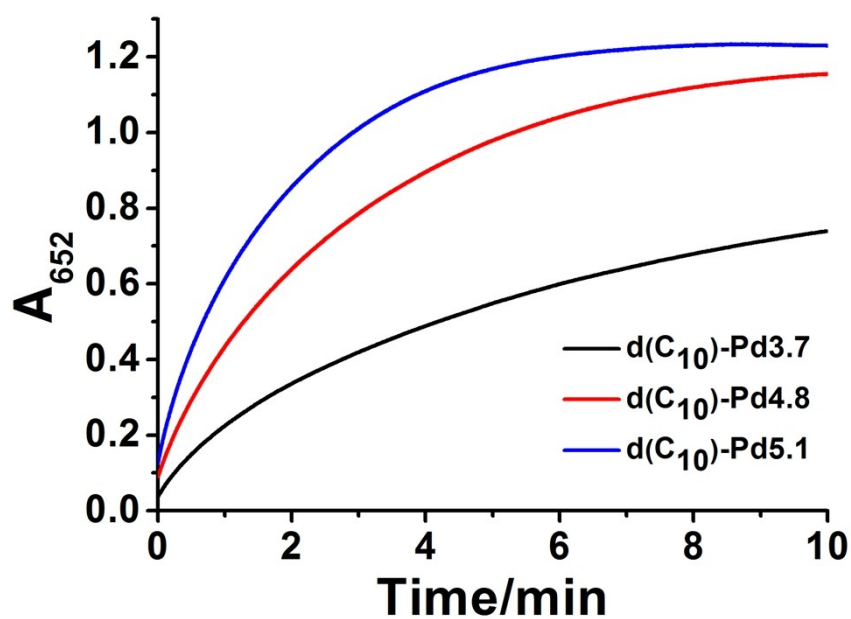


Fig. S1 UV-vis absorption-time course curves of TMB- H_2O_2 reaction system catalyzed by different $d(C_{10})$ -Pd at 20 °C.

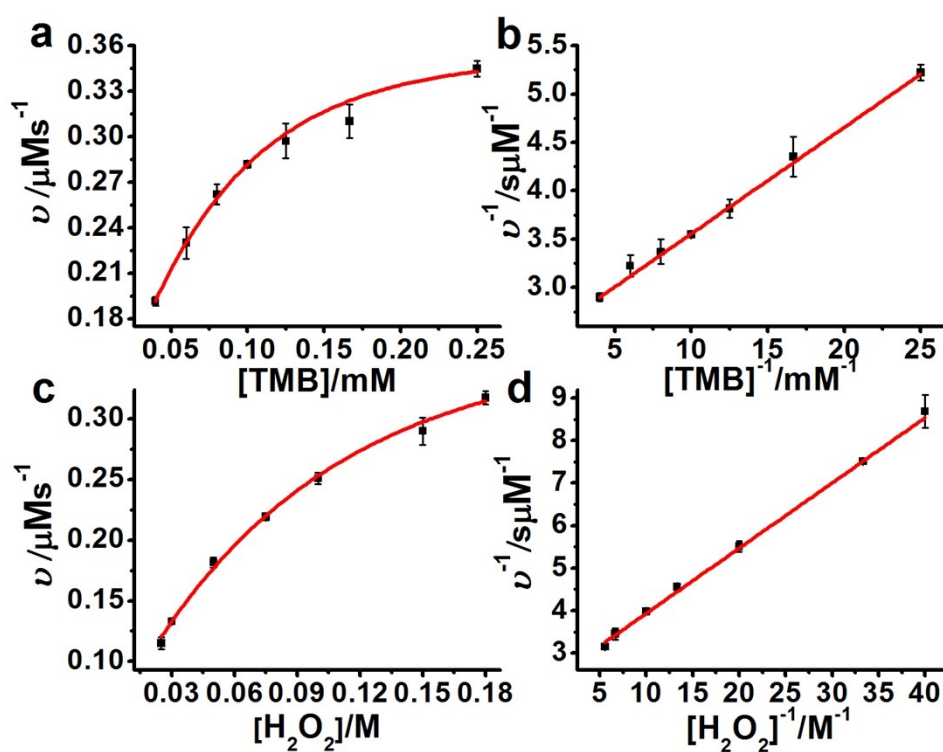


Fig. S2 Steady-state kinetics of d(C₁₀)-Pd5.1 are measured through the oxidization of TMB in the presence of H₂O₂ at 20 °C at pH 4.0 using 900 nM Pd (calculated from precursor): (a) The concentration of H₂O₂ is fixed at 125 mM and the TMB concentration is varied, (c) The concentration of TMB is fixed at 0.125 mM and the H₂O₂ concentration is varied, (b) and (d) are double-reciprocal plots of (a) and (c), respectively.

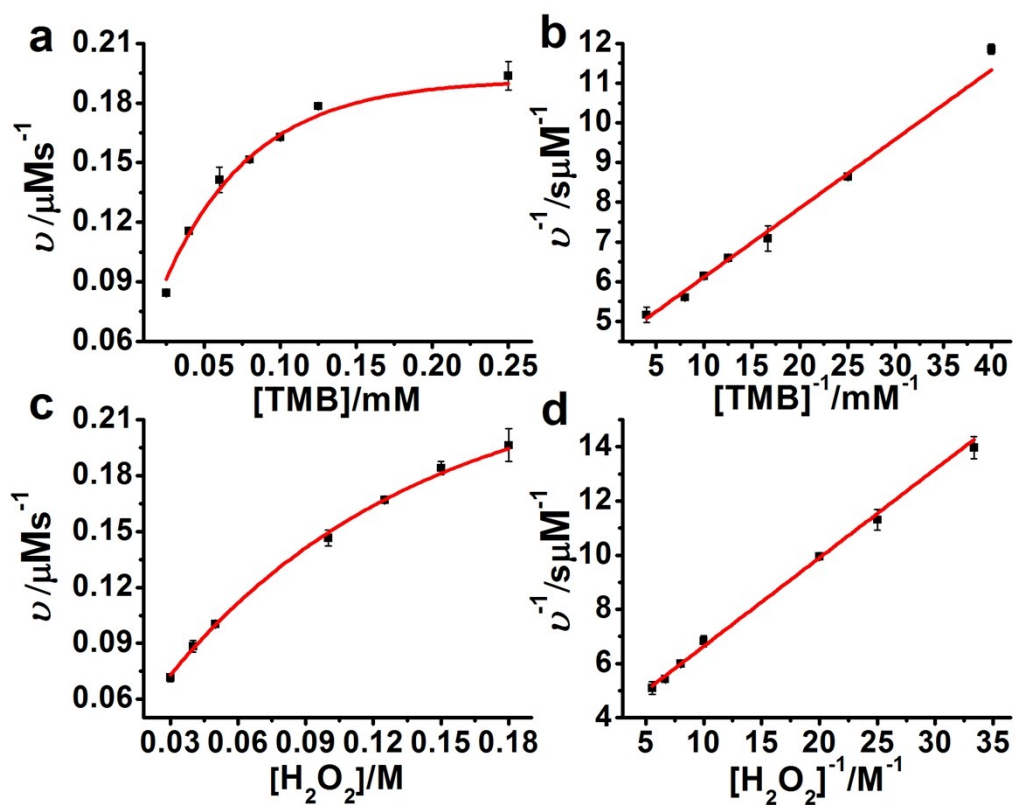


Fig. S3 Steady-state kinetics of d(C₁₀)-Pd_{4.8} are measured using 900 nM Pd (calculated from precursor) in the presence of 150 nM Cys: (a) The concentration of H₂O₂ is fixed at 125 mM and the TMB concentration is varied, (c) The concentration of TMB is fixed at 0.125 mM and the H₂O₂ concentration is varied, (b) and (d) are double-reciprocal plots of (a) and (c), respectively.

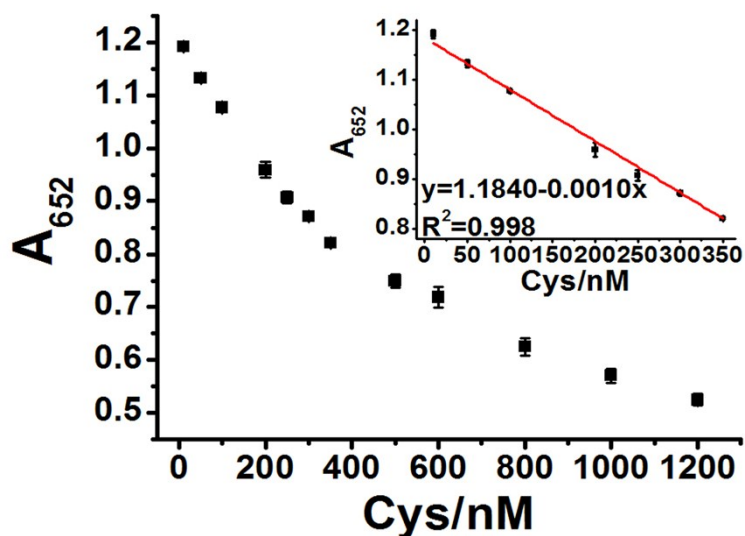


Fig. S4 Plots of the A_{652} with the concentration of Cys using d(C₁₀)-Pd5.1, and the inset is corresponding calibration curve.

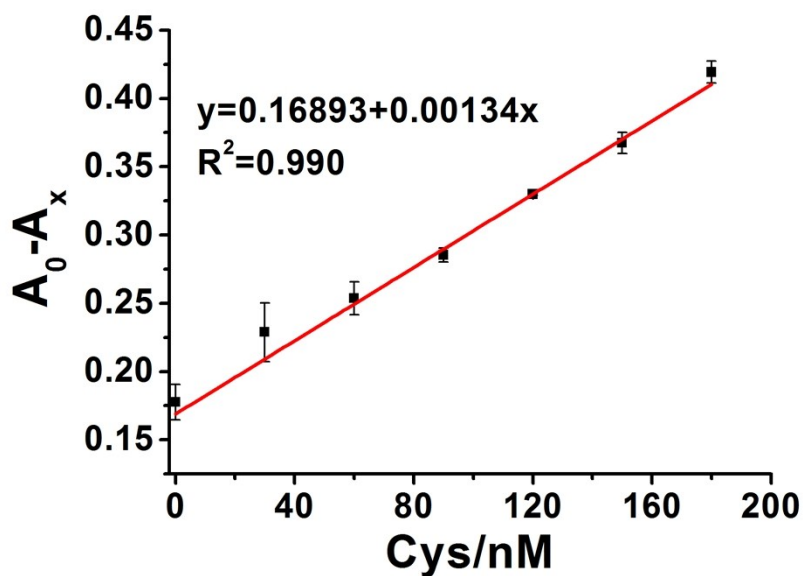


Fig. S5 Calibration curve for Cys in human serum samples, A_0 represents the A_{652} collected from the TMB- H_2O_2 reaction in NaH_2PO_4 - Na_2HPO_4 buffer (pH 4.0) catalyzed by d(C₁₀)-Pd4.8, and A_x represents the A_{652} collected from the TMB- H_2O_2 reaction in diluted serum (pH 4.0) catalyzed by d(C₁₀)-Pd4.8 in the absence and presence of Cys. The A_{652} signal was collected at 10 min after initiation. The concentration of Cys in diluted human serum was determined as the epitaxial linear fellowship at horizontal axis.