

Supplementary Information

Colorimetric and Optical Hg(II) Ions Sensor Developed with Conjugates of M13-Bacteriophage and Silver Nanoparticles

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M13-bacteriophage growth, purification and amplification process:

M13_{wild} was grown and purified following standard biochemical protocols described in literature.¹⁻⁴ Briefly, one colony of E. coli XL-1 blue was grown in 3 ml of LB media to mid-log phase (E. coli XL-1 blue culture) and infected with 10 μ L of M13_{wild}. The culture was incubated at 37°C with shaking for 12 h, and then centrifuged to remove E. coli. The M13_{wild} was collected by PEG/NaCl (20% PEG and 2.5 mol/L NaCl) precipitation and reconstituted in Tris-buffered saline (10 mM). The typical yield was ~ 20 mg of M13_{wild} per liter. The final concentration was determined spectrophotometrically using an extinction coefficient of 3.84 cm²/mg at 269 nm.^{3,5}

Attachment of cation binding peptides at the major coat protein (Gene VIII) of M13 is reported previously.⁶ Briefly, two primers were designed to insert 4E into the gene VIII protein: 5'-ATATATCTGCAGNKGAAAGAGGAAGAGCCCGCAAAGCGGCCTTTAACTCCC-3' (4E), and 5'-GGAAGCTGCAGCGAAAGACAGCATCGGAACGAGG-3' (linearization primer). To collect M13_{4E} phages, the inverse polymerase chain reaction (PCR) cloning method was performed using the above mentioned primers (the linearization primer with M13_{4E} primer). The PCR product was purified then the amplified plasmid, and phage plaques were verified by DNA sequencing. Furthermore, we have amplified the M13_{4E} phage for our experiments, and the methods were same to the M13_{wild} as mentioned above.^{3,5}

References:

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3. G.P. Smith and V.A. Petrenko, Phage display, *Chem. Rev.*, 1997, **972**, 391-410.

4. Y. Seo, S. Manivannan, I. Kang, S. W. Lee and K. Kim, *Biosens. Bioelectron.*, 2017, **94**, 87-93.

5. J. Sambrook and D.W. Russell, Cold Spring Harbor Laboratory Press 2001.

6. J. P. Park, M. Do, H. E. Jin, S. W. Lee and H. Lee, *ACS Appl. Mater. Interfaces*, 2014, **6**, 18653-18660.

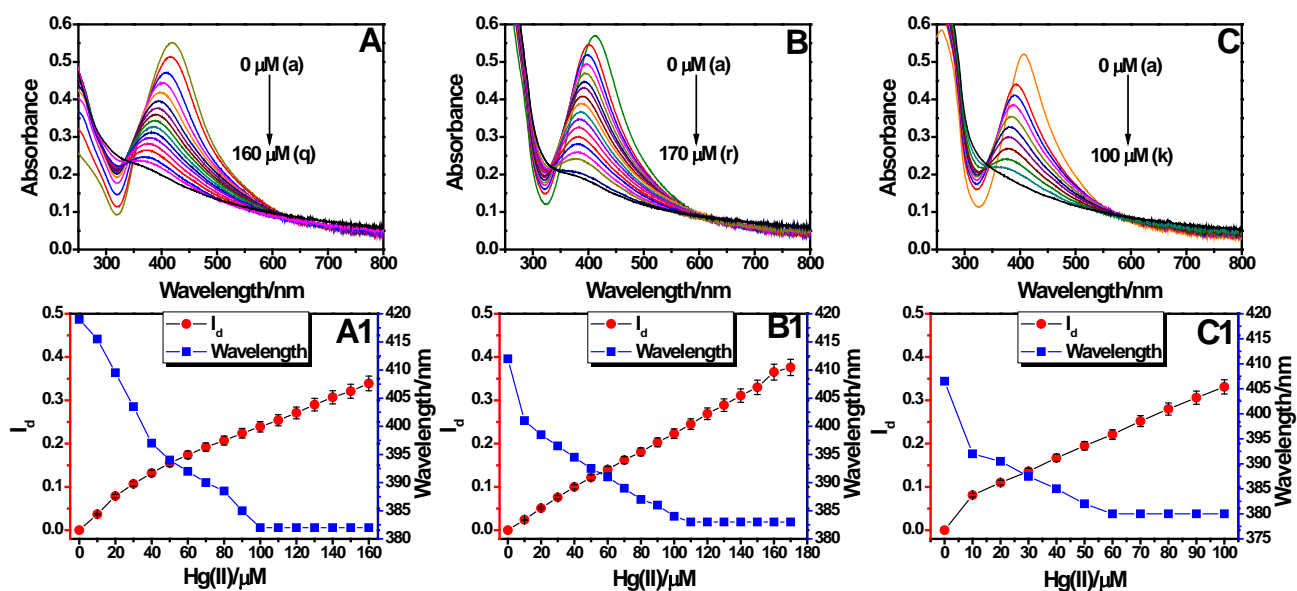


Figure. S1 (A-C) SPR absorption spectral changes and (A1-C1) plots of changes in the absorption intensity and wavelength observed for (A and A1) SSG-Ag NPs, (B and B1) SSG-Ag_{wild} NCBs, and (C and C1) SSG-Ag_{4E} NCBs upon each addition of 10 μM of Hg(II) ions.

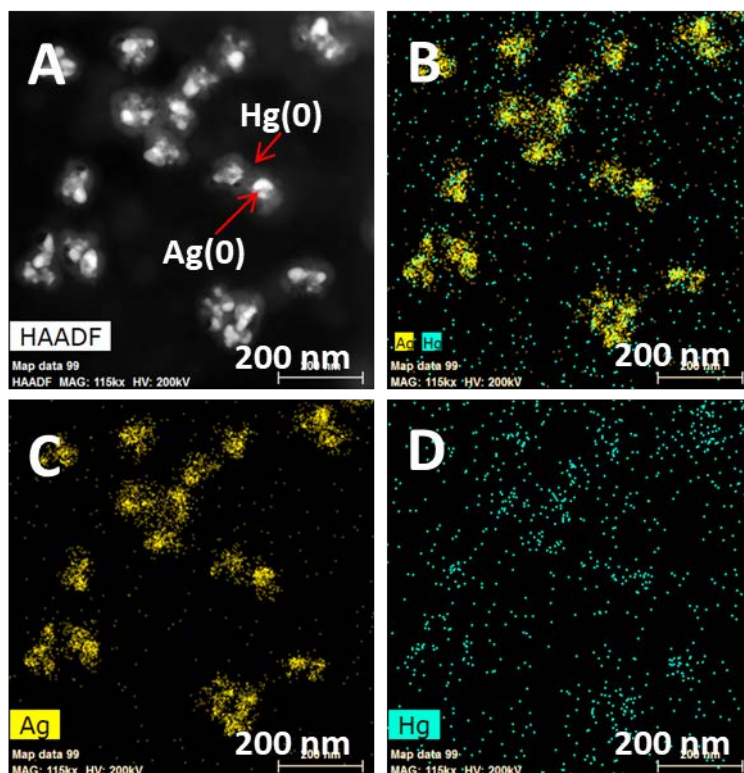


Figure. S2 (A) STEM-HAADF images and (B-D) STEM-EDX mapping analyses of AgHg-amalgam crystals.

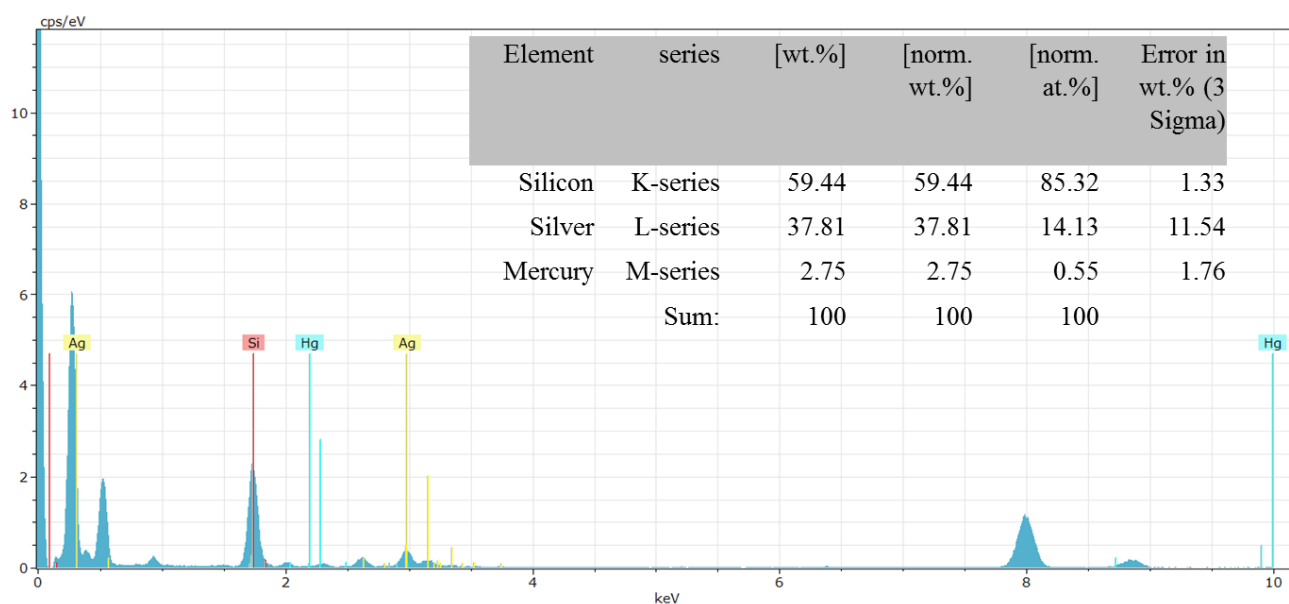


Figure. S3 STEM-EDX point ID analysis of AgHg-amalgam crystals.

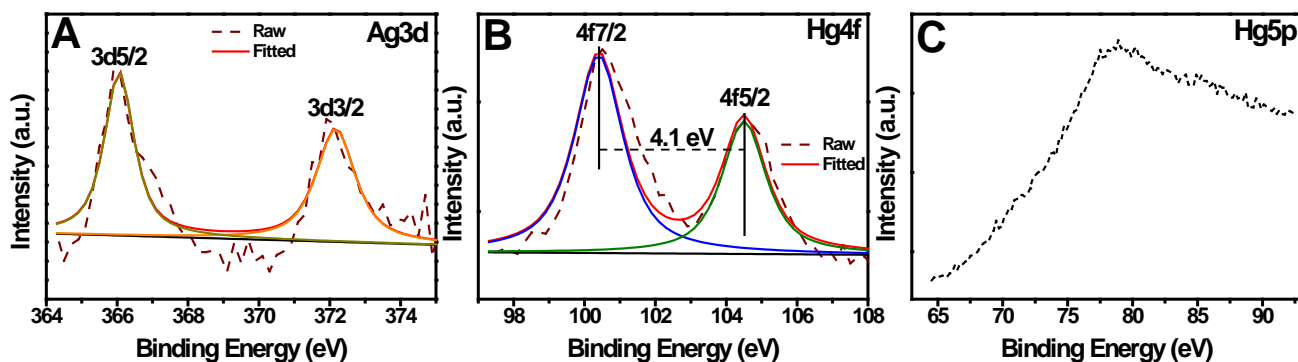


Figure. S4 XPS spectra obtained for the AgHg-amalgam crystals and their corresponding (A) Ag3d, (B) Hg4f, and (C) Hg5p regions of core-level spectra.

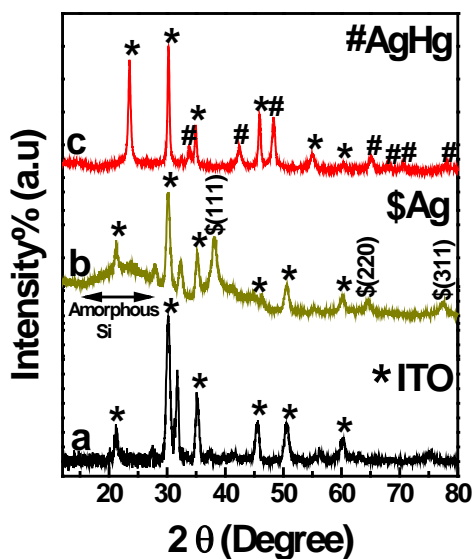


Figure S5. XRD analysis of (a) Bare ITO substrate, (b) SSG-Ag_{4E} CGs, (c) AgHg-amalgam crystals.

Table. S1 Quantification of Hg(II) ions in different water samples by using SSG–Ag_{4E} CGs and ICP-MS.

Hg(II) Ions (μM)							
Samples	SSG–Ag_{4E} NCBs				ICP-MS		
	Added (μM)	Found (μM)	Recovery (%)	RSD (%)	Found (μM)	Recovery (%)	RSD (%)
Tab water	0	0	–	–	0	–	–
	25	24.93	99.73	2.02	24.57	98.28	0.74
	50	50.36	100.73	2.06	49.22	98.44	0.51
Pond water	0	0	–	–	0	–	–
	25	26.34	105.36	0.66	28.10	112.40	1.76
	50	52.85	105.71	1.79	54.20	108.4	0.99