

# Supplementary Materials: An Ionic 1,4-Bis(styryl)benzene-Based Fluorescent Probe for Mercury(II) Detection in Water via Deprotection of the Tthioacetal Group

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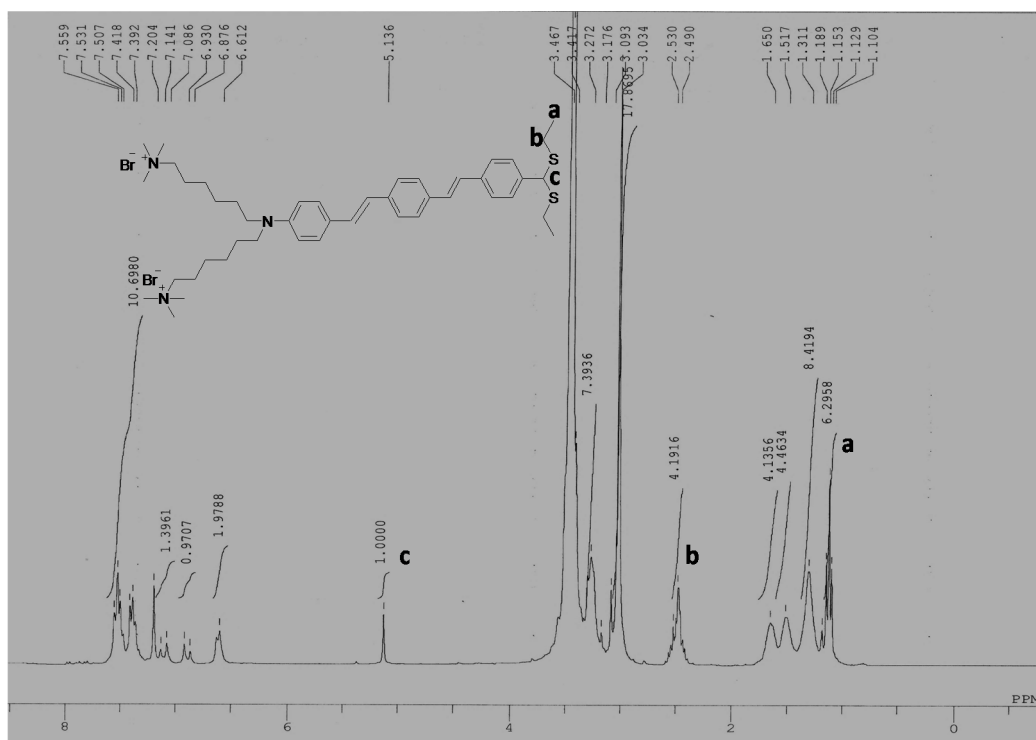


Figure S1. <sup>1</sup>H-NMR spectrum of M1Q in dimethyl sulfoxide-d<sub>6</sub>.

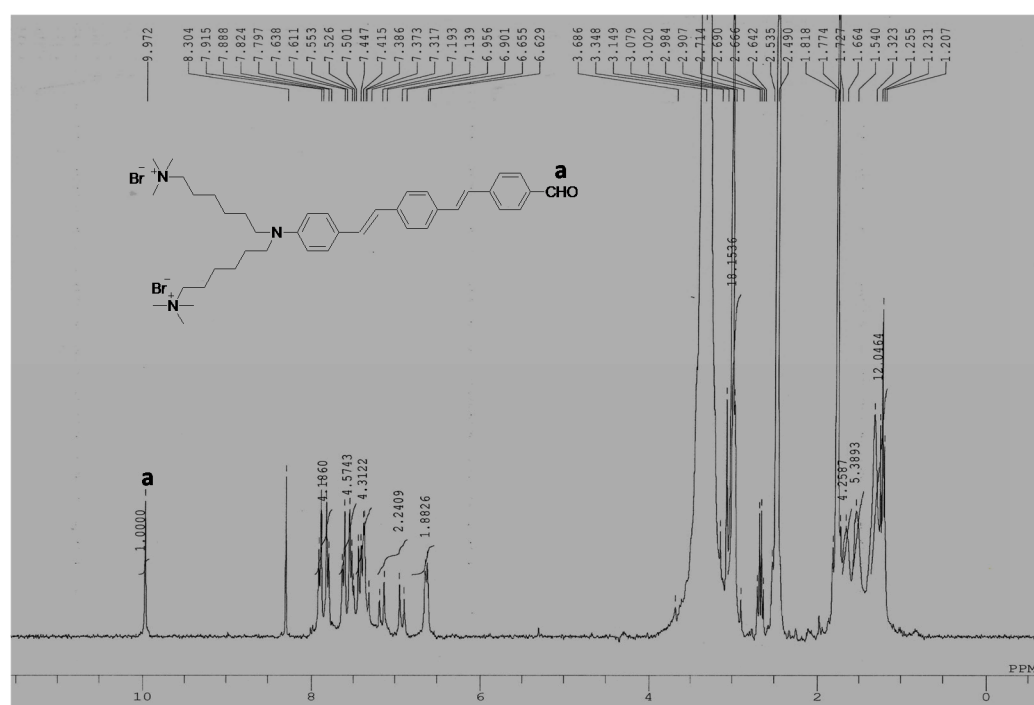
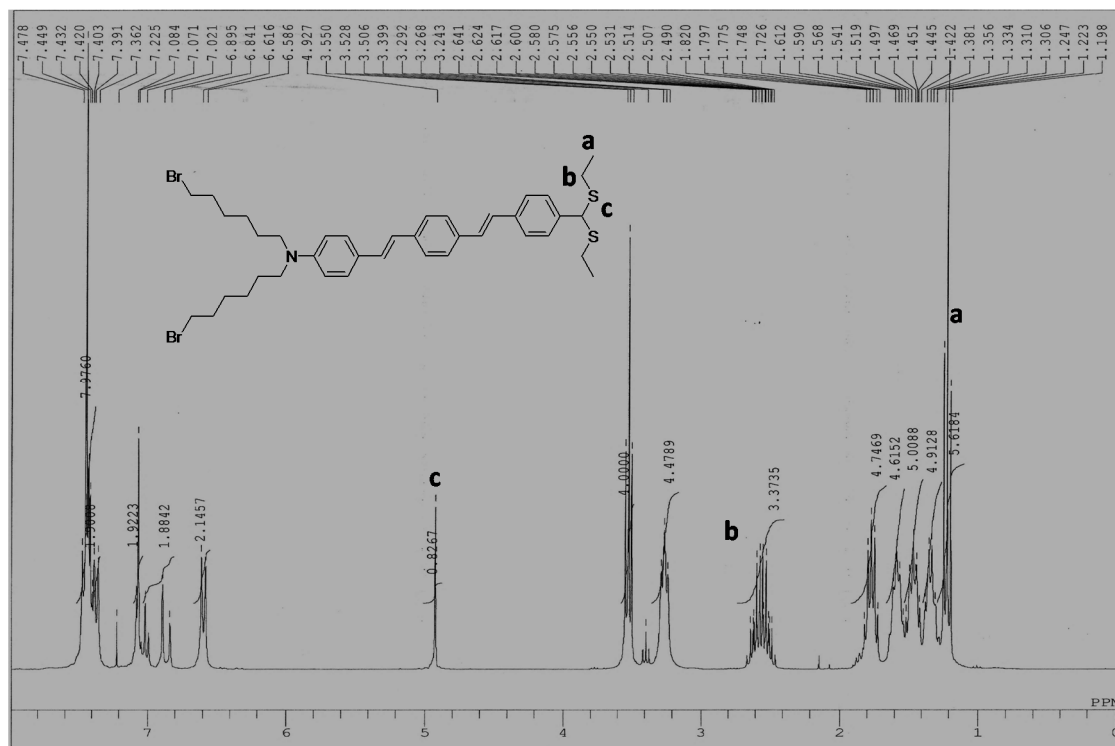
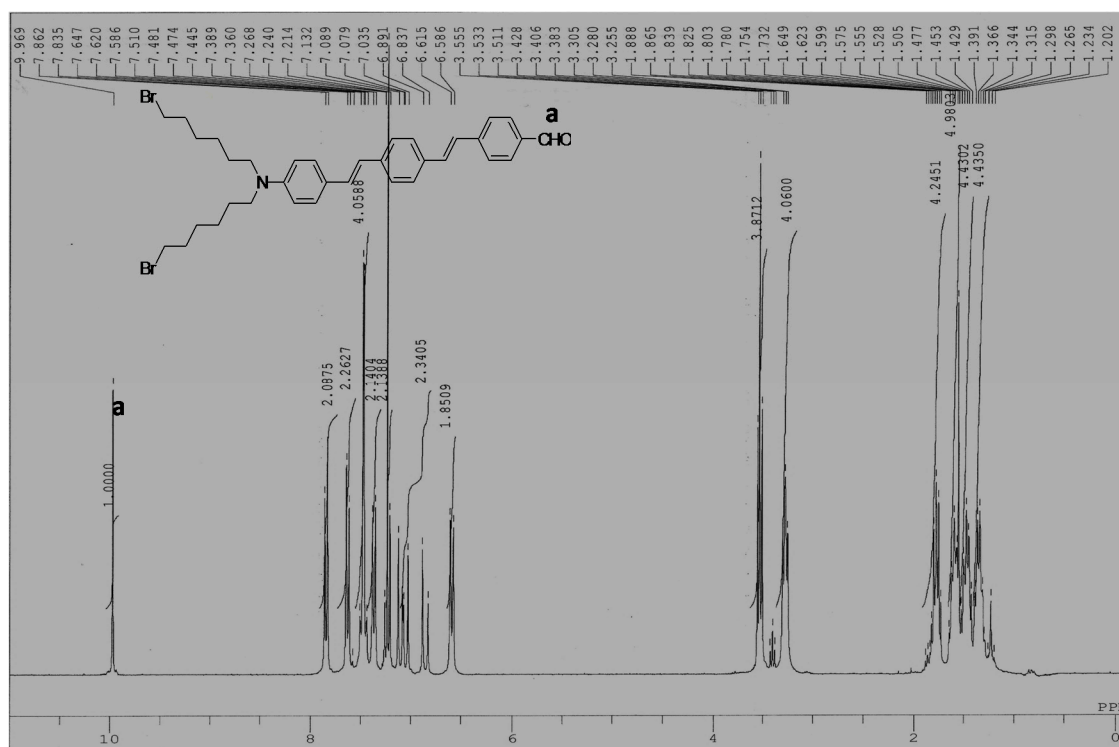
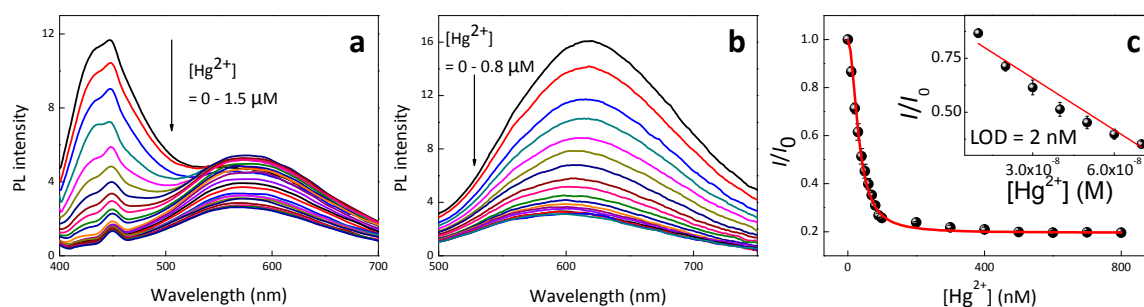
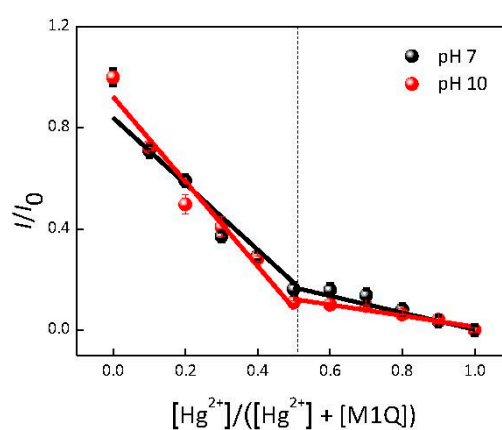


Figure S2. <sup>1</sup>H-NMR spectrum of M1Q-CHO in dimethyl sulfoxide-d<sub>6</sub>.

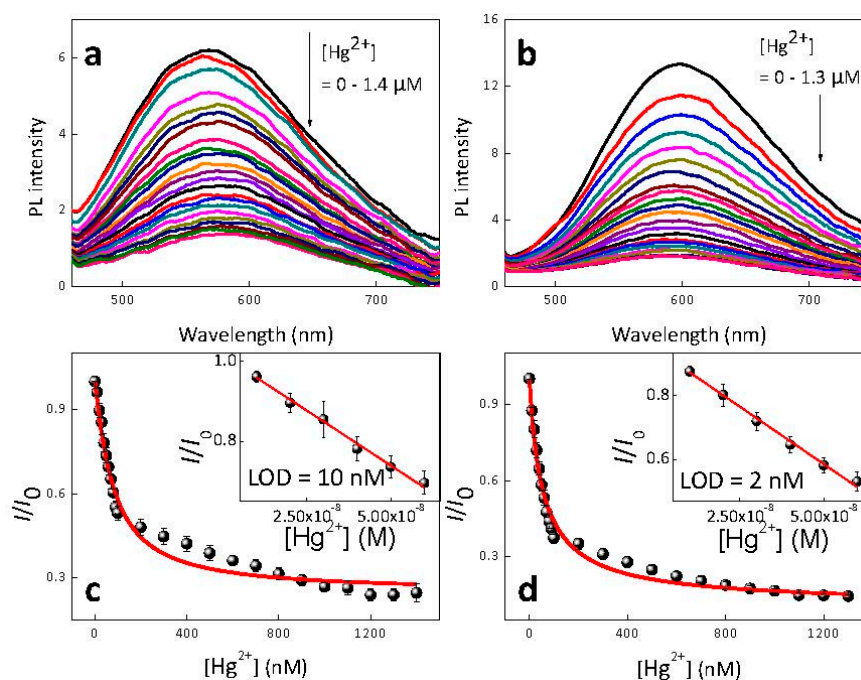
Figure S3.  $^1\text{H-NMR}$  spectrum of M1 in  $\text{CDCl}_3$ .Figure S4.  $^1\text{H-NMR}$  spectrum of M1-CHO in  $\text{CDCl}_3$ .



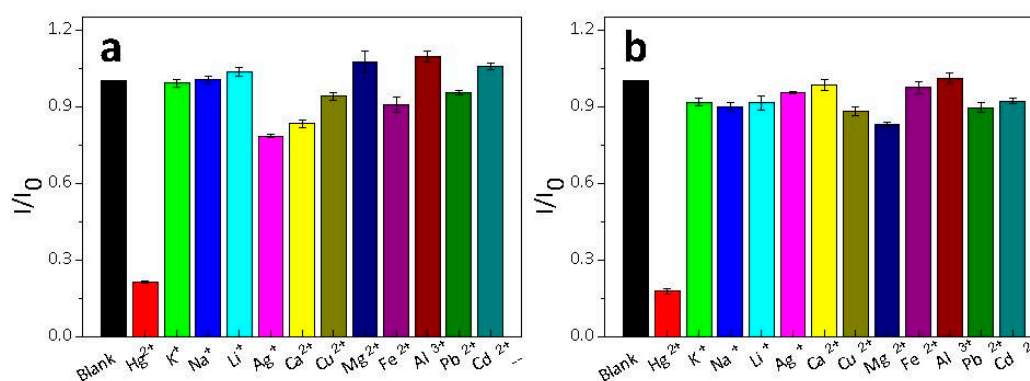
**Figure S5.** (a,b) PL emission spectra of M1Q in pH 4 and pH 10 with increasing  $[Hg^{2+}]$ . Excitation at 390 nm. (c) Intensity ratio versus  $[Hg^{2+}]$  at pH 10. Inset: linear range of titration curve varying  $[Hg^{2+}]$ . The error bars represent the standard deviation of three independent measurements.  $[M1Q] = 1.0 \times 10^{-6}$  M in water.



**Figure S6.** Job plots of M1Q in water at pH 7 and pH 10.



**Figure S7.** (a,b) PL spectra of M1Q in Han River and in tap water with increasing  $[Hg^{2+}]$ ; (c,d) Normalized PL intensity ( $I/I_0$ ) of M1Q with changing  $[Hg^{2+}]$ . Excitation at 390 nm. Inset: LOD determination. The error bars represent the standard deviation of three independent measurements.  $[M1Q] = 1.0 \times 10^{-6}$  M.



**Figure S8.** Selectivity test for M1Q in the Han River (a) and in tap water (b) in the presence of metal ions by excited at 390 nm. The error bars represent the standard deviation of three independent measurements. [M1Q] =  $1.0 \times 10^{-6}$  M; [Hg<sup>2+</sup>] =  $2.0 \times 10^{-6}$  M; [other metal ion] =  $2.0 \times 10^{-6}$  M.

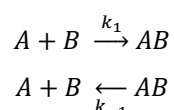
#### \* Determination of limit of detection, Hill equation and dissociation constant.

The terms of limit of detection (LOD), Hill equation, dissociation constant, etc. are closely related to sensor characteristics. The LOD is the lowest analyte concentration that can be distinguished from the absence of that analyte [1]. The value of LOD can be calculated by the following equation:  $\text{LOD} = 3.3 \times \sigma / \text{slope}$ . The  $\sigma$  is the standard deviation in the PL intensity measurements of probe itself without analytes. The slope of  $I/I_0$  vs. [Hg<sup>2+</sup>] was determined via linear fitting [2].

The Hill equation, which was originally formulated by Archibald Hill in 1910, describes the sigmoidal binding curve of analyte as a function of the analyte concentration [3]. By determining the degree of analyte binding to probe, the detection range was determined by fitting the titration curve using the Hill equation, where  $I/I_0$  transits from 10% to 90% of its signal output [4,5].

A dissociation constant,  $K_d$  is equal to the concentration of free A when half of the total molecules of B are associated with A ( $A_x B_y \rightleftharpoons xA + yB$ ) [6].

The binding/dissociation equilibrium of M1/M1Q (denoted as A) and [Hg<sup>2+</sup>] (denoted as B) can be described by



$k_1$ : association rate constant,  $k_{-1}$ : dissociation rate constant.

For the above bimolecular reaction, the equilibrium dissociation constant ( $K_d$ ) can be defined as

$$K_d = \frac{k_{-1}}{k_1} = \frac{[A][B]}{[AB]}$$

or

$$\frac{[AB]}{[A]} = \frac{[B]}{K_d}$$

$$\text{Fraction A bound} = \frac{[AB]}{[A] + [AB]} = \frac{1}{\frac{[A]}{[AB]} + 1} = \frac{1}{\frac{K_d}{[B]} + 1} = \frac{[B]}{K_d + [B]}$$

From this expression, if Fraction A bound =  $[B]/(K_d + [B]) = 0.5$ , then  $[B] = K_d$  [7]. Thus,  $K_d$  is simply determined by measuring the Hg<sup>2+</sup> concentration when the  $I/I_0$  ratio decreases to half of the original value without Hg<sup>2+</sup> (Fraction A bound = 0.5 from a titration curve).

## References

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