Supporting Information

## Self-Standing Aptamers by Artificial Defect-Rich Matrix

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**Figure S1.** Cyclic voltammograms of 1 mM  $[Ru(NH_3)_6]^{3+}$  on different surfaces. (A) Bare Au (curve a) and aptamer-modified Au (curve b). (B) CLL-treated Au (curve a) and aptamer-modified CLL-treated Au (curve b). These measurements were performed in 25 mM TRIS buffer at pH 7.4 with a scan rate of 100 mV/s. (C) The 1 mM  $[Ru(NH_3)_6]^{3+}$  CV response change upon aptamer-modification of bare Au (A) and CLL-treated Au (B) surfaces. (N = 3) The error bars indicate the standard deviation.



**Figure S2.** Surface aptamer density histograms from substrates fabricated by different processes. Each data point is obtained from the average fluorescence intensity of three samples (N = 3). The error bars indicate the standard deviation.



**Figure S3.** Linear correlation curve derived from  $(F_0-F)/F$  against the logarithm of cocaine concentration and (inset) the cocaine concentration. Each data point is obtained from the average fluorescence signal of individual squares (N = 4). The error bars indicate the standard deviation.

## Calculations of dissociation constants (K<sub>d</sub>)

According to literature reports,<sup>1-4</sup> aptamer probe 3 can hybridize with cocaine with a molar ratio of 1:1. Therefore, the  $K_d$  can be defined as the formula given in equation (1):

$$K_{d} = \frac{[aptamer] \times [cocaine]}{[aptamer/cocaine]}$$
(1)

where [aptamer], [cocaine], and [aptamer/cocaine] are the concentrations of unhybridized aptamer, unhybridized cocaine, and hybridized cocaine/aptamer on the gold surface, respectively. Based on 1:1 of aptamer/cocaine binding stoichiometry, when [aptamer] = [aptamer/cocaine], equation (1) is simplified to  $K_d$  = [cocaine]. In this case, when cocaine leads half the total of FAM-labeled aptamer to fluorescence quenching or half-maximal fluorescence image contrast change,  $K_d$  is equal to cocaine concentration.

As shown in Figure S1, the linear fitted equation of correlation is:

$$Y = 0.26 X - 0.076 \tag{2}$$

in which Y and X respectively represent the value of  $(F_0-F)/F$  and the logarithm of cocaine concentration ( $\mu$ M). (Note:  $F_0$  and F represent corresponding relative fluorescence intensities without and with targets, respectively.) The maximum value of  $(F_0-F)/F$  (Y<sub>max</sub>) is 0.62 ± 0.02, while the  $(F_0-F)/F$  value of blank can be seen as zero due to  $F = F_0$ . Thus, Y<sub>max/2</sub> is estimated to be 0.31 ± 0.01, which indicates the half-maximal fluorescence image contrast change. According to equation (2), the calculated X is 0.90 ± 0.03, and thus the cocaine concentration is  $10^{0.90} = 7.97 \pm$ 0.57  $\mu$ M. Due to K<sub>d</sub> = [cocaine] is at which half-maximal fluorescence image contrast change occurs, the K<sub>d</sub> is determined to be 7.97 ± 0.57  $\mu$ M.



**Figure S4.** Selectivity examination of the CLL-fabricated aptamer-anchored platform toward cocaine. Each substrate was treated by 1 mM solutes. Each data point is obtained from the average fluorescence signal of individual squares (N = 4). The error bars indicate the standard deviation. The scale bars are 20  $\mu$ m.



Figure S5. Reusability test of the CLL-fabricated aptamer-anchored platform for several measurement cycles of 100  $\mu$ M cocaine. Each data point is obtained from the average fluorescence signal of individual squares (N = 4). The error bars indicate the standard deviation.



Figure S6. Comparison of the CLL-fabricated aptamer-anchored platform before and after 6-month of storage under 100% relative humidity at 4°C in dark. The selected pattern defect site (red arrows) indicates that the data were collected from the same sample. The scale bars are 20  $\mu$ m.



**Figure S7.** (A) Representing photo and (B-E) corresponding fluorescence images of different sampling spots obtained from the same CLL-fabricated aptamer-anchored substrate, and their responses toward the target. The scale bars are 20 µm.

## REFERENCES

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