

Supporting Information

Electrochemical DNA biosensor based on the “Y” junction structure and restriction endonuclease-aided target recycling strategy

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Experimental Section

Materials. DNA oligonucleotides were synthesized by TaKaRa Biotechnology Co., Ltd. (Dalian, China). Restriction endonuclease HaeIII were from New England Biolabs, Inc.. 6-Mercapto-1-hexanol (MCH) and hexaammineruthenium (III) chloride (RuHex) were from sigma. All of the chemical reagents were of analytical grade or higher. Ultrapure water (18.2 M Ω ·cm) was used throughout.

Modification of Au Electrode with different surface coverage. Au electrodes (2 mm in diameter, 99.99%, Shanghai Chenhua Equipment, China) were cleaned before surface modification. The cleaned electrodes were modified as our previously work (*Anal. Chim. Acta*, 2011, **688**, 163–167). The low surface coverage was obtained by incubation of electrodes with 1 μ M of capture probe in the 0.05 M NaCl solution for 30 min. The medium surface coverage was prepared by incubation of electrodes with 2 μ M of capture probe in 10 mM phosphate buffer (pH 7.3) for 120 min. The high surface coverage was prepared by incubation of electrodes with 5 μ M of capture

probe in the 1 M NaCl solution for 120 min. After rinsed with water, capture probe modified Au electrodes were dried under nitrogen. Then, the electrodes were immersed in 1 mM MCH at 25 °C for 60 min. After rinsed with water, the electrodes were dried under nitrogen. The coverage of capture probes on the Au film was measured by chronocoulometry (*Anal. Chem.*, 1998, **70**, 4670–4677).

Feasibility study. The feasibility of enhanced biosensors was investigated. Capture probe was first modified on Au electrode, then the mixture of 1 μM probe 4, 0.1 U/μL and 1nM cDNA was added and incubated for 60 min at 37 °C. After washing thoroughly with 10 mM PBS buffer, the electric currents before and after reaction were recorded respectively.

Electrochemical detection. All electrochemical measurements were performed with a CHI660C electrochemical workstation (Shanghai Chenhua Equipment, China). The conventional three-electrode system was employed, which consisted of Au working electrode, platinum wire auxiliary electrode, and KCl saturated calomel reference electrode. Chronocoulometry was carried out at a pulse period of 250 ms and pulse width of 500 mV. The electrolyte was 10 mM Tris-HCl buffer (pH 7.4) containing 50 μM $[\text{Ru}(\text{NH}_3)_6]^{3+}$. Alternating-current voltammetry (ACV) was performed in 10 mM PBS buffer (pH 7.0) containing 200 mM NaCl. ACV was recorded at a frequency of 100 Hz and potential amplitude of 4 mV. All electrolyte buffers were deoxygenated via purging with nitrogen before experiments.

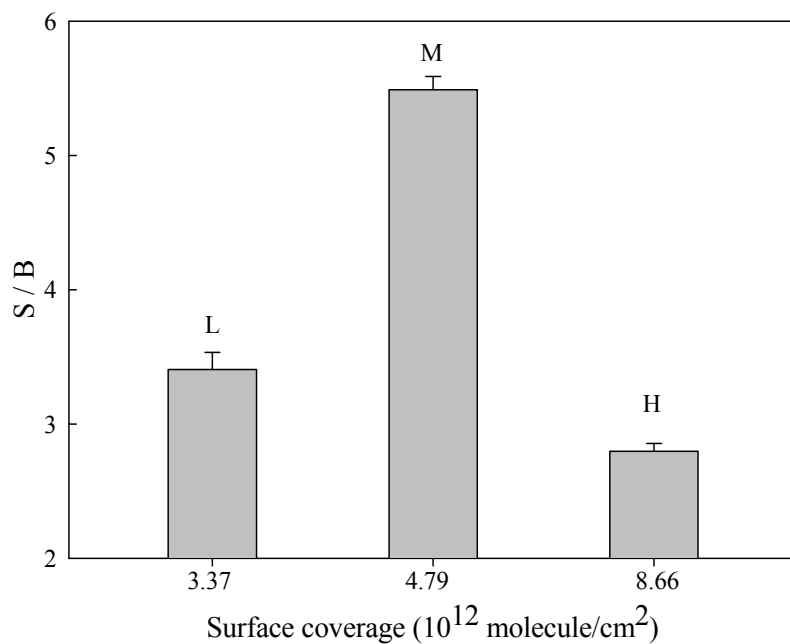


Fig. S1 Signal / background ratio (S/B) corresponding to (L) low surface coverage (3.37×10^{12} molecule/cm 2), (M) medium surface coverage (4.79×10^{12} molecule/cm 2) and (H) high surface coverage (8.66×10^{12} molecule/cm 2). The error bars represent the standard deviation of six measurements. Concentration of HaeIII: 0.1 U/ μ L; Assistant probe: assistant probe4; Concentration of assistant probe: 1 μ M; Temperature: 37 $^{\circ}$ C; Reaction time: 60 min.

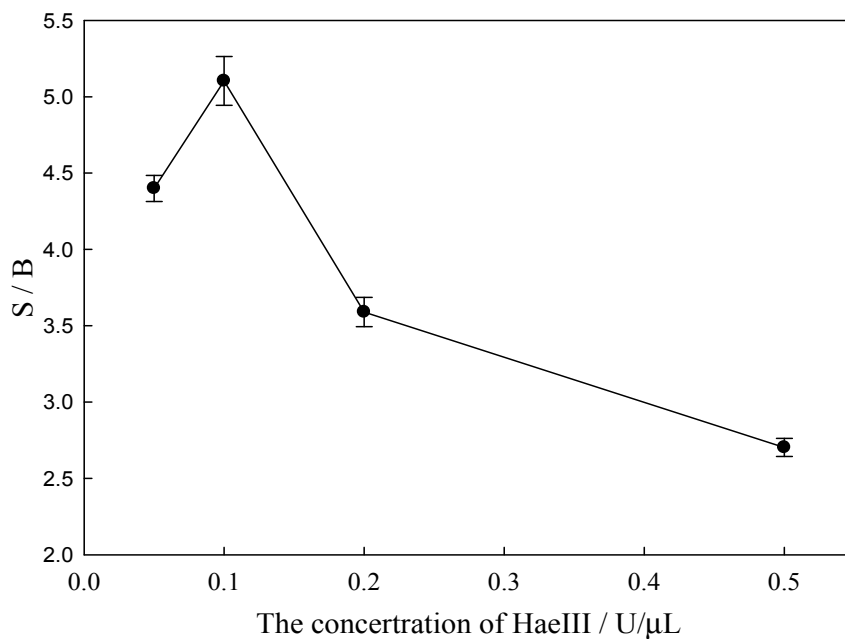


Fig. S2 Signal / background ratio (S/B) corresponding to different concentration of HaeIII. The error bars represent the standard deviation of six measurements. Concentration of cDNA: 10 nM; Surface coverage: 4.79×10^{12} molecule/cm²; Assistant probe: assistant probe4; Concentration of assistant probe: 1 μM; Temperature: 37 °C; Reaction time: 60 min.

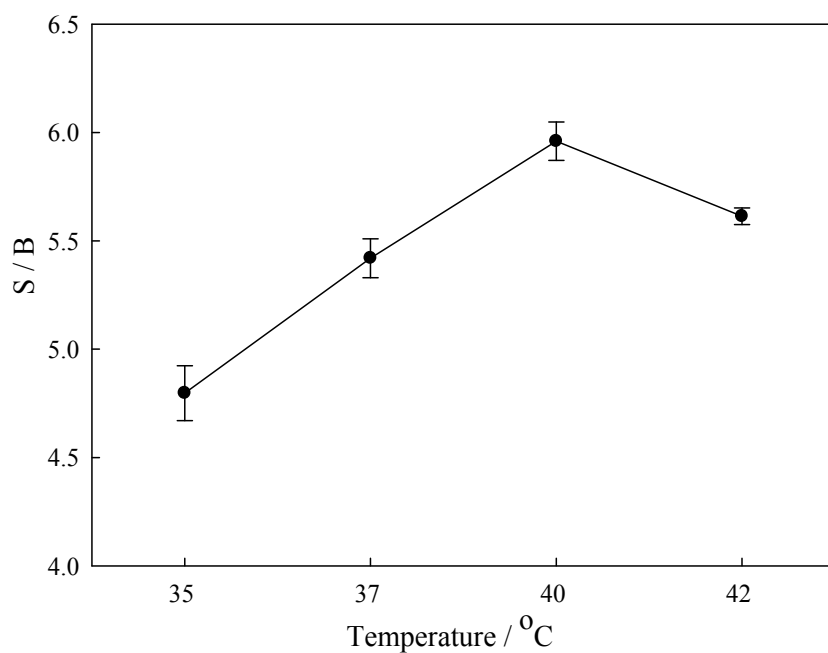


Fig. S3 Signal / background ratio (S/B) corresponding to different reaction temperature. The error bars represent the standard deviation of six measurements. Concentration of cDNA: 10 nM; Concentration of HaeIII: 0.1 U/ μ L; Surface coverage: 4.79×10^{12} molecule/ cm^2 ; Assistant probe: assistant probe3; Concentration of assistant probe: 1 μ M; Reaction time: 60 min.

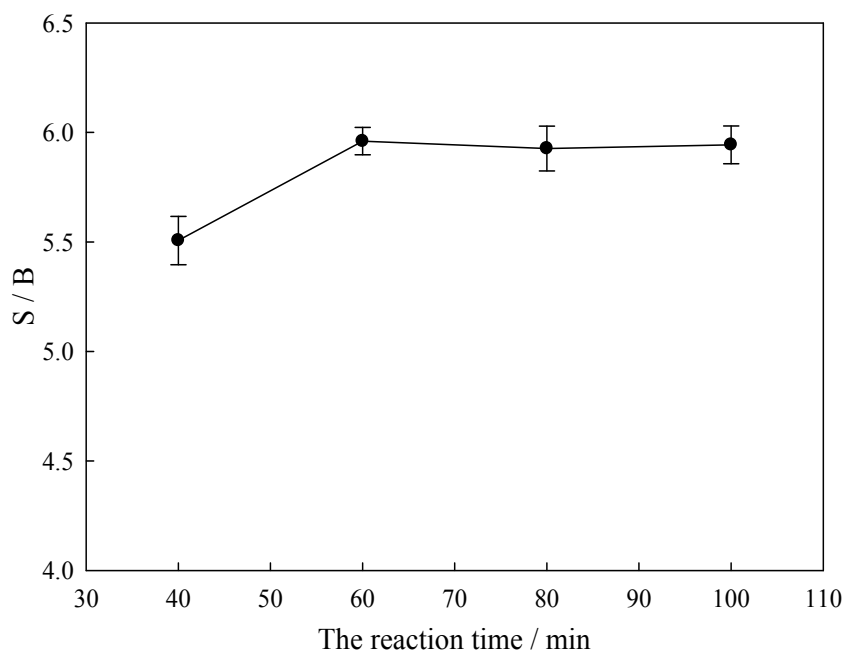


Fig. S4 Signal / background ratio (S/B) corresponding to different reaction time.

The error bars represent the standard deviation of six measurements. Concentration of cDNA: 10 nM; Concentration of HaeIII: 0.1 U/ μ L; Surface coverage: 4.79×10^{12} molecule/ cm^2 ; Assistant probe: assistant probe3; Concentration of assistant probe: 1 μ M; Temperature: 40 $^{\circ}$ C.