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# **Supplementary Materials**

## Ultrasensitive detection of flap endonuclease 1 using a

# chemiluminescence optical fiber biosensor with

hybridization chain reaction

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### Pretreatment of optical fiber

4 cm bare fiber (SFS400/440/700 T, numerical aperture 0.22. Wyoptics Technology Co., Ltd., Shanghai, China) was used to construct sensor in the assay. First, the distal end of optical fiber (about 1.5 cm) was immersed into chloroform solution for 10 min and then in 24% HF for 20 min to remove the acrylic sheath and silica cladding. After rinsing with 0.1 M HCl to get rid of impurities, the fiber was inserted in 0.1 M NaOH for 1 h to activate the silanol group on the surface. The obtained bare core of the optical fiber was washed with ultrapure water and dried by nitrogen gas.

### Table S1 Sequences of DNAs used in the present study.

| Name             | Sequence (from 5 'to 3')                     |
|------------------|--|
| СР               | COOH-TCACACCATGTCACTCCAGCAATTTGGAGTGACATG    |
| $H_1$            | TGGAGTGACATGGTGTGACATGTCACTCCAAATTGC-biotin  |
| $\mathrm{H}_{2}$ | biotin-TCACACCATGTCACTCCAGCAATTTGGAGTGACATG  |
| 5' flap          | TGGAGTGACATGGTGTGA                           |
| Т                | CCCCACTCGCTGCAATCACTGAACGTTACTAGAT           |
| D                | TGGAGTGACATGGTGTGAGTGATTGCAGCGAGTGGGG-biotin |
| U                | ATCTAGTAACGTTCAT                             |

| Table S2 Comparison of diffe | rent methods for FEN | 1 determination. |
|------------------------------|----------------------|------------------|
|------------------------------|----------------------|------------------|

| Method                       | Detection system                        | LOD                                | Linear range   | Ref       |
|------------------------------|---|------------------------------------|--|-----------|
| Reverse transcription        |   | semiquantitative                   |  | 1         |
| polymerase chain reaction    |   |                                    |  |           |
| Western blot                 |   | qualitative                        |  | 2         |
| Immunohistochemistry         |   | qualitative                        |  | 3         |
| Fluorescence method          | DNA-Ag nanoclusters sensor              | 10 pM                              | 20–1000 pM   | 4         |
| Enzyme-assistant rolling     | Dumbbell DNA-SYBR green I               | 15 fM                              | 20-8000 fM   | 5         |
| circle amplification         | fluorescence sensor                     |                                    |  |           |
| Nt.BstNBI-induced tandem     | G-quadruplex-thioflavin T               | 0.002-3.6 fmol                     | 1.68 amol  | 6         |
| signal amplification*        | fluorescence sensor                     | (0.001–1.5 U)                      | (0.75 mU)  |           |
| Enzyme-assistant exponential | DNA-SYBR green I fluorescence           | 0.5 fM                             | 1 fM-10 pM   | 7         |
| amplification reaction       | sensor                                  |                                    |  |           |
| Enzyme-free branched*        | DNA-Ru(phen) <sub>3</sub> <sup>2+</sup> | 0.052 fM                           | 0.16–1.4×10 <sup>4</sup> fM                            | 8         |
| hybridization chain reaction | electrochemiluminescence sensor         | $(2.2 \times 10^{-2} \text{ U/L})$ | $(6.5 \times 10^{-2} - 6.5 \times 10^{3}  \text{U/L})$ |           |
| Enzyme-free Hybridization    | Chemiluminescent optical fiber          | 3.4 fM                             | 10 fM-74 pM  | This work |
| chain reaction               | sensors                                 |                                    |  |           |

\*The unit of FEN1 was unified molar concentration according the conversion formula of 1 U/L = 2.4 fM which is obtained by New England biolabs Inc.



Fig. S1 The picture of the biosensor system (A) and optical fiber sensing unit (B).



Fig. S2. Correlation between the proposed method and ELISA (n=5). The data shows 95% confidence interval of the mean.

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