

Support Information

Multifunctional electrochemical biosensor with "tetrahedral tripods" assisted multiple tandem hairpins assembly for ultra-sensitive detection of target DNA

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Table S1 DNA sequences used in this work

| Name | Sequences (5' - 3') |
|-----------------------------------|---|
| TTs-S1 | ACATTCCTAAGTCTGAAACATTACAGCTTGCTACACGAGAAGAGCCGC CATAGTATTTAAAGCTCGGCAGCTCCGGCCTGCG |
| TTs-S2 | SH-C6- TATCACCAGGCAGTTGACAGTGTAGCAAGCTGTAATAGATGCGAGGGT CCAATAC |
| TTs-S3 | SH-C6- TCAACTGCCTGGTGATAAAACGACACTACGTGGGAATCTACTATGGCG GCTCTTC |
| TTs-S4 | SH-C6- TTCAGACTTAGGAATGTGCTTCCCACGTAGTGTCTGTATTGGACCC TCGCAT |
| TTS1-miRNA 141 | ACATTCCTAAGTCTGAAACATTACAGCTTGCTACACGAGAAGAGCCGC CATAGTACCATCTTTACCAGACAGTGTACGCATTTTGTGTGTTTGT TGGTA |
| microRNA 141 | UAACACUGUCUGGUAAAGAUGG |
| Target DNA | TCTCAAGGACCACCGCATCTCTACCGCAGGCCGGAGCTGCCGAGCTTT |
| Methylated DNA | TCTCAAGGACCACCGCATCTCTACCGCAGGCCmGGAGCTGCCGAGCTT T |
| Hairpin H1 | GTAGAGATGCGGTGGTCCTTGAGAGAATTCTTAACGTCGCCTCATACT GTCTCAAGGACCACCGCAT |
| Hairpin H2 | TCTCAAGGACCACCGCATCTCTACATGCGGTGGTCCTTGAGACAGTAT GCCTAGCAGAGTT |
| Hairpin H3 | TCGCCTCCTAGCAGAGTTACTTTGAAACTCTGCTAGGAGGCGACGTTA AGAATTC |
| Hairpin H4 | TCAAAGTAACTCTGCTAGGAGGCGAGAATTCTTAACGTCGCCTCCTAG CAGAGTT |
| Hairpin H1-miRNA 141 | TACCAGACAAACACAACAAAATGCGGAATTCTTAACGTCGCCTCATACT TGCGCATTTTGTGTGTTTGT |
| Hairpin H2-miRNA141 | CGCATTTTGTGTGTTTGTCTGGTACAAACACAACAAAATGCGCAGTAT GCCTAGCAGAGTT |
| Single-stranded capture probe | SH-C6-TTTTTAAAGCTCGGCAGCTCCGGCCTGCG |
| Single-base mismatched DNA | TCTCAAGGACCACCGCATCTCTACCGCAGGCCGGAGCTGCCGAGCTTT |
| Two-base mismatched DNA | TCTCAAGGACCACGGCATCTCTACCGCAGGCCGGAGCTGCCGAGCTTT |
| Multi-base Mismatched DNA | TCTCAAGGAGCAGGGCATCTCTACCGCACGCCGGAGCTGCCGAGCTTT |
| Non-complementary base mismatch-A | CGTGGAAGCACTCTTGCTAAACACGCAGGCCGGAGCTGCCGAGCTTT |
| Non-complementary base mismatch-B | TCTCAAGGACCACCGCATCTCTACCGTGGAAAGCACTCTTGCTAAACA |

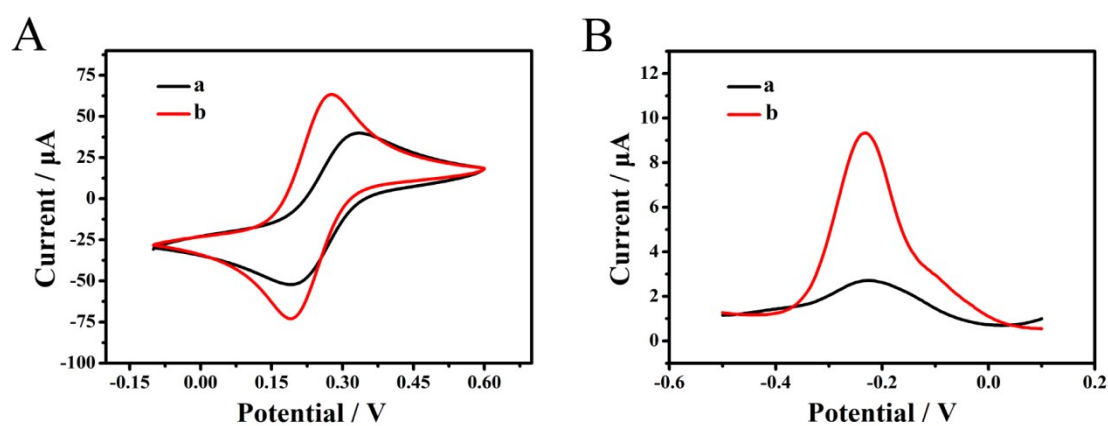


Fig. S1. CV (A) and DPV(B) of the fabricated biosensor. (a) MTHsA/MCH/TTs/Au electrode. (b) MTHsA/miRNA/MCH/TTs/Au electrode. CV current response of the incubation process carried out in 5 mM $\text{Fe}(\text{CN})_6^{3-/4-}$ with 0.1 M KCl solution, pH=7.4. DPV current response of the incubation process carried out in 10mM Tris-HCl with 50 μM RuHex.

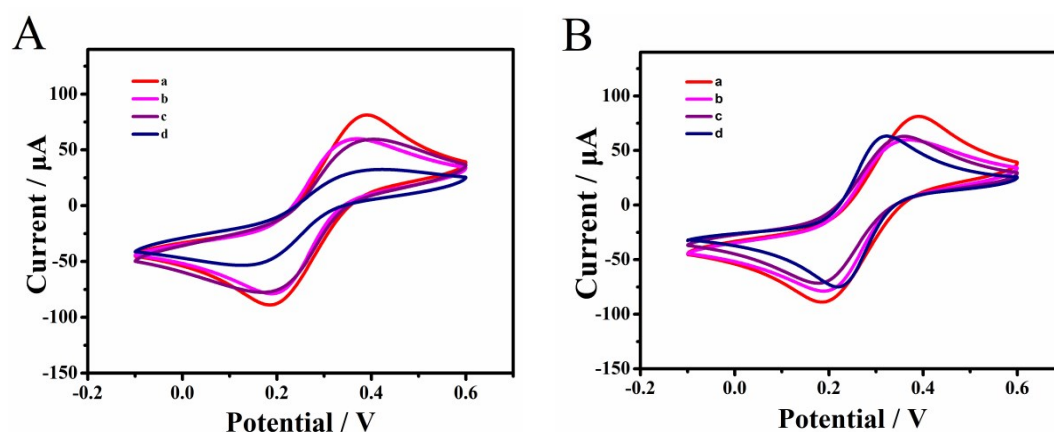


Fig. S2. CV (A) and (B) of the developed biosensor were respectively methylated DNA and unmethylated DNA after 50 U/mL Hpa II endonuclease digestion. (a) TTs/Au electrode, (b) Methylated DNA or Unmethylated DNA /MCH/TTs/Au electrode, (c) Hpa II enzyme /Methylated DNA or Unmethylated DNA/MCH/TTs/Au electrode, (d) MTHsA/Hpa II enzyme /Methylated DNA or Unmethylated DNA/MCH/TTs/Au electrode.

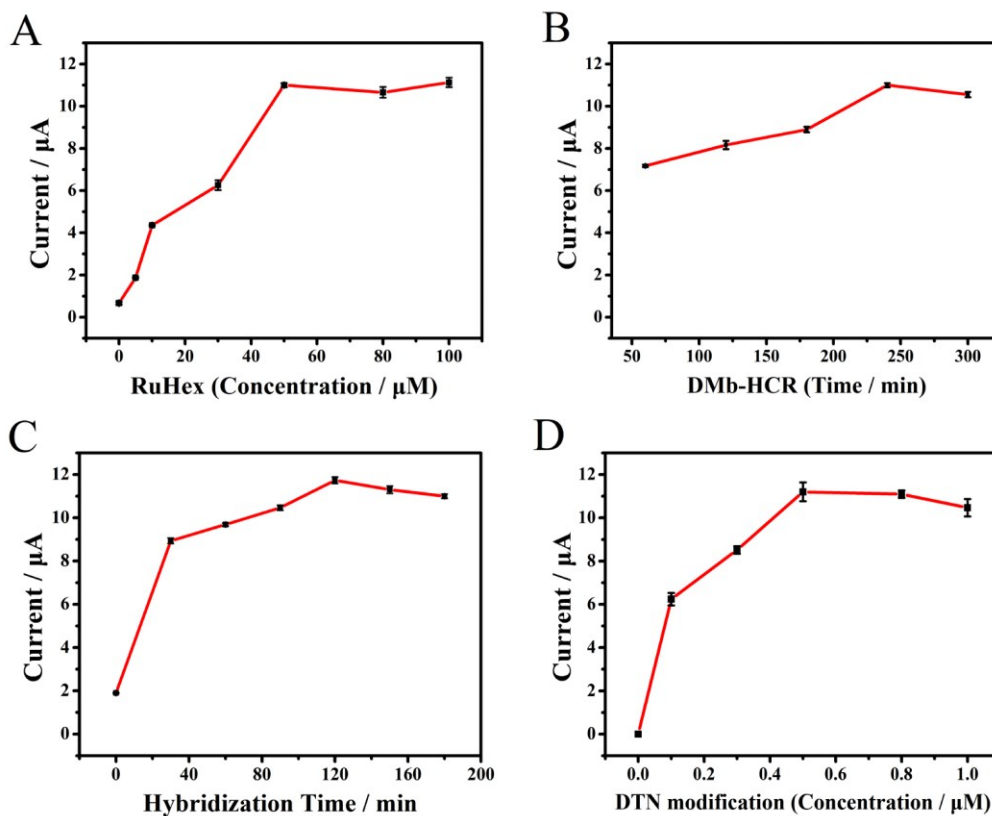


Fig. S3. (A) The optimization of RuHex concentration. (B) The optimization of MTHsA reaction time. (C) The optimization of DNA hybridization time with the target DNA. (D) The optimization of TTs concentration. The error bars show the standard deviations of electrochemical measurements taken from the three tests.

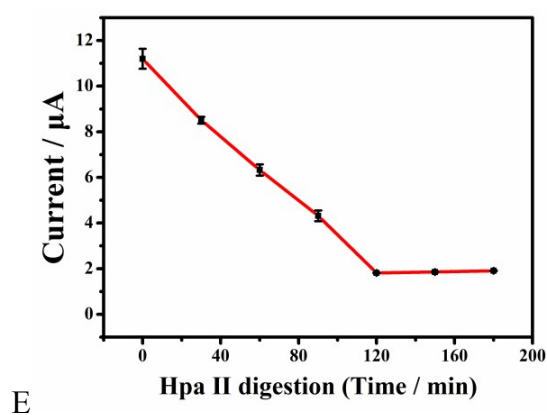


Fig. S4. The optimization of digestion time of Hpa II endonuclease. Error bars show the standard deviations of electrochemical measurements taken from the three tests.

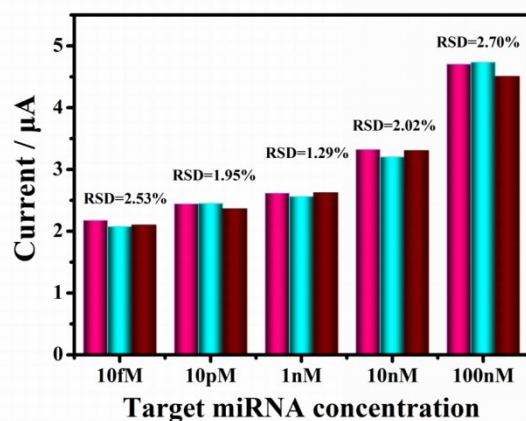


Fig. S5. Comparison of the reproducibility for different target miRNA concentration in 10% human serums (n=3) on the fabricated electrochemical biosensor.

Table S2 Electrochemical analysis results of spiking DNA into human serum.

| Samples | Added DNA | Detected DNA | Recovery (%) (n = 3) | RSD (%) |
|---------|-----------|--------------|----------------------|---------|
| 1 | 10 fM | 10.06 fM | 100.6 | 0.98 |
| 2 | 100 fM | 105.54 fM | 105.54 | 0.72 |
| 3 | 1 pM | 1.017 pM | 101.71 | 0.52 |
| 4 | 10 pM | 9.857 pM | 98.57 | 1.01 |
| 5 | 100 pM | 96.03 pM | 96.03 | 0.72 |

Table S3 The comparison of the constructed electrochemical biosensor towards MTHsA with other reported methods for target DNA detection.

| Amplification strategy | Analytical technique | Linear range | LOD | Reference |
|------------------------|----------------------|-------------------|---------|-----------|
| ACTRAa/Exo III | DPV | 10 fM to 500 fM | 10 fM | 1 |
| NHCRb/HRP | Amperometric | 1 fM to 1000 fM | 0.4 fM | 2 |
| CHAc/AuNP | Colorimetric | 50 pM to 700 pM | 9 pM | 3 |
| TMSDd/MB | DPV | 0.01 pM to 100 pM | 8.2 fM | 4 |
| ZrHCF MNPs e/MBs | DPV | 1.0 fM to 1.0 nM | 0.43 fM | 5 |
| CSDPRf/HCR | DPV | 10 fM to 1 nM | 8fM | 6 |
| HCR/HRP | Chronoamperometric | 1 aM to 1 pM | 0.93 aM | 7 |

| | | | | |
|-------------|-----|----------------|---------|-------------|
| DNCg/HCR | DPV | 5 aM to 1 pM | 5 aM | 8 |
| MTHsA/RuHex | DPV | 1 aM to 100 pM | 0.59 aM | This method |

^a ACTRA, autocatalytic target recycling amplification; ^b NHCR, nonlinear hybridization chain reaction; ^c CHA, catalyzed hairpin assembly; ^d TMSD, toehold-mediated strand displacement; ^e ZrHCF MNPs, magnetic zirconium hexacyanoferrate nanoparticles; ^f CSDPR, circular strand-displacement polymerase reaction; ^g DNC, dendritic DNA concatamer.

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