

Supporting Information

Tetrahedral DNA Nanostructure-decorated Electrochemical Platform for Simple and Ultrasensitive EGFR Genotyping of Plasma ctDNA

Xuyao Wang,^{†,*} Jianping Wu,[‡] Weilin Mao,[‡] Xia He,[§] Liming Ruan,[†] Junlan Zhu,[†] Peng Shu,[†] Zhenqi Zhang,[†] Bitao Jiang[†] and Xingguo Zhang^{†,*}

[†] Precision Medicine Center, Beilun People's Hospital, Zhejiang University School of Medicine First Affiliated Hospital Beilun Branch, Ningbo, Zhejiang, 315806, P. R. China

[‡] Department of Clinical Laboratory, Zhejiang University School of Medicine First Affiliated Hospital, Hangzhou, Zhejiang, 310003, P. R. China

[§] Department of Clinical Laboratory, Shengzhou People's Hospital, Zhejiang University School of Medicine First Affiliated Hospital Shengzhou Branch, Shengzhou, Zhejiang, 312400, P. R. China

Corresponding Author

*E-mail: zhxgblhospital@163.com. E-mail: xuyaowang22@qq.com

Contents

Table of the sequences used. Photograph of SPE used. Additional data and figures of the assay.

Table S1. Sequences of the DNA oligonucleotides employed in this study.

Name	Sequence (from 5' to 3')	Remarks
Tetra-B	SH-C ₆ - TATCACCAGGCAGTTGACAGTGTAGCAAG CTGTAATAGATGCGAGGGTCCAATAC	The sequence to form the scaffold of the tetrahedron
Tetra-C	SH-C ₆ - TCAACTGCCTGGTGATAAAACGACACTACG TGGGAATCTACTATGGCGGCTCTTC	The sequence to form the scaffold of the tetrahedron
Tetra-D	SH-C ₆ – TTCAGACTTAGGAATGTGCTTCCCACGTAG TGTCGTTTGTATTGGACCCTCGCAT	The sequence to form the scaffold of the tetrahedron
Tetra-2A19	ACATTCCTAAGTCTGAAACATTACAGCTTG CTACACGAGAAGAGCCGCCATAGTATTTTT TTTTTGGATCCCAGAAGGTGAGAAAGTTAA AATTCCCGT	The sequence to form the scaffold of the tetrahedron with a pendant probe complementary to exon 19 target
Tetra-2A21	ACATTCCTAAGTCTGAAACATTACAGCTTG CTACACGAGAAGAGCCGCCATAGTATTTTT TTTTTAAACACCGCAGCATGTCAAGATCAC AGATTTTGG	The sequence to form the scaffold of the tetrahedron with a pendant probe complementary to exon 21 target
Tetra-3A	TTCCCCACGAGGGCTTCAACTCTATTTTTT TTTACATTCCTAAGTCTGAAACATTACAGCT TGCTACACGAGAAGAGCCGCCATAGTA	The sequence to form the scaffold of the tetrahedron with a pendant probe complementary to the assistant probe
19ass	TAGAGTTGAAGCCCTCGTGGGGAATTGGA TCCCAGAAGGTGAGAAAGTTAAATTCCC GT	Complementary to exon 19 and Tetra-3A
21ass	TAGAGTTGAAGCCCTCGTGGGGAATTAAA CACCGCAGCATGTCAAGATCACAGATTTTG G	Complementary to exon 21 and Tetra-3A
19AntiT	TTAACGTCTTCCTTCTCTCTGTCATAGGG ACTCTGGATCCCAGAAGGTGAGAAAGTTA AAATTCCCGTCGCTATCAAAACATCTCCGA AAGCCAACA	Anti-target sequence of exon 19
21AntiT	GTCGCTTGGTGCACCGCGACCTGGCAGCC AGGAACGTACTGGTGAAAACACCGCAGCA TGTCAGATCACAGATTTTGGGCGGACCA AACTGCTGGGTG	Anti-target sequence of exon 21, the red nucleotide is adjusted to fit 21MR
19F	TGCCAGTTAACGTCTTCCTTC	Forward primer
19F2	TGCCAGTTAACGTCTTCCTTC	An extra nucleotide was added to fix the T _m . Labeled in red.

19WR	Biotin-GGCTTTCGGAGATGTTGCTTCTC	Complementary to wild type
19MR1	Biotin-TGTTGGCTTTCGGAGATGTT TTG	Complementary to mutated type. Three nucleotides were appended at the split. Labeled in red.
19MR2	Biotin-GGCTTTCGGAGATGTT TTGATAG	Complementary to mutated type. Seven nucleotides were appended at the split. Labeled in red.
21F	GCATGAACTACTTGGAGGAC	Forward primer
21F2	G GCATGAACTACTTGGAGGAC	An extra nucleotide was added to fix the Tm. Labeled in red
21WR	Biotin-CACCCAGCAGTTTGGTCA	Complementary to wild type
21MR1	Biotin-CACCCAGCAGTTTGG TCC	Complementary to mutated type, with an extra mismatch at -2 position. Labeled in red.
21MR2	Biotin-CACCCAGCAGTTTGGCCC	Complementary to mutated type



Figure S1. Photograph of SPE used in this platform.

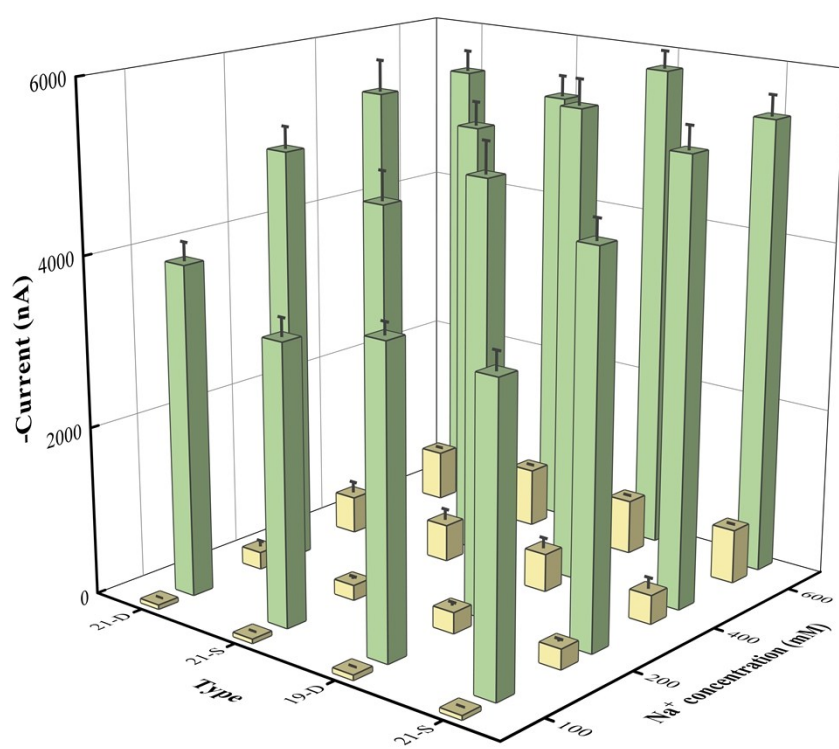


Figure S2. Optimal conditions for Na⁺ concentration.

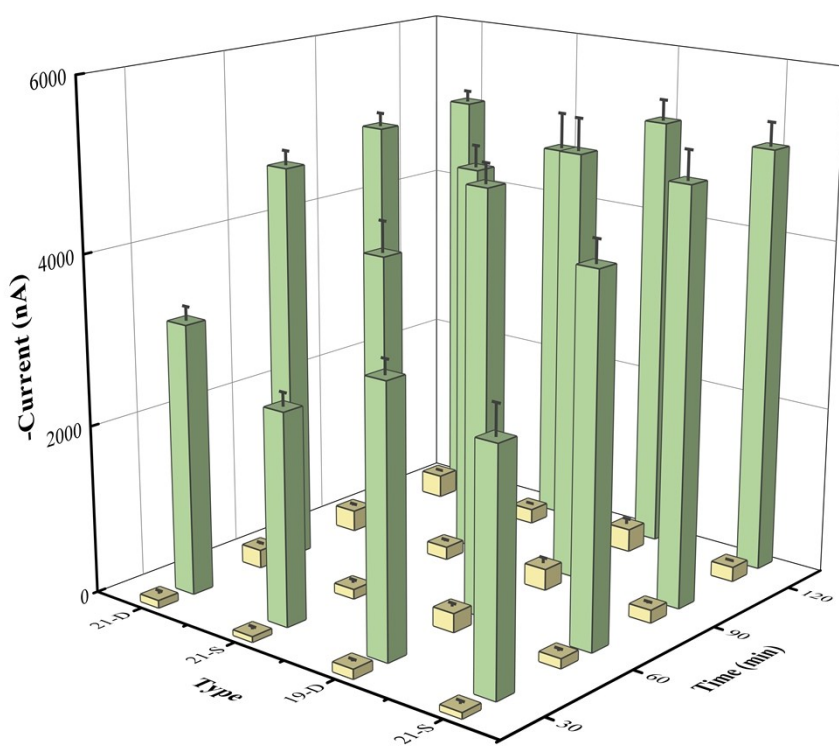


Figure S3. Optimal conditions for hybridization time.

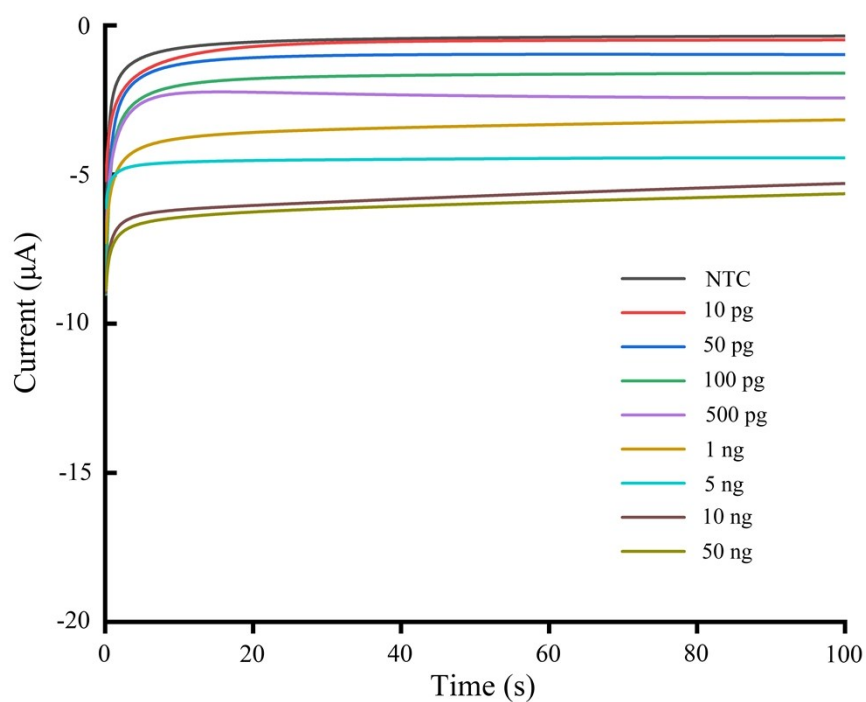


Figure S4. Current values of different amount of 19del DNA input. The signal gradually rises as more DNA is added.

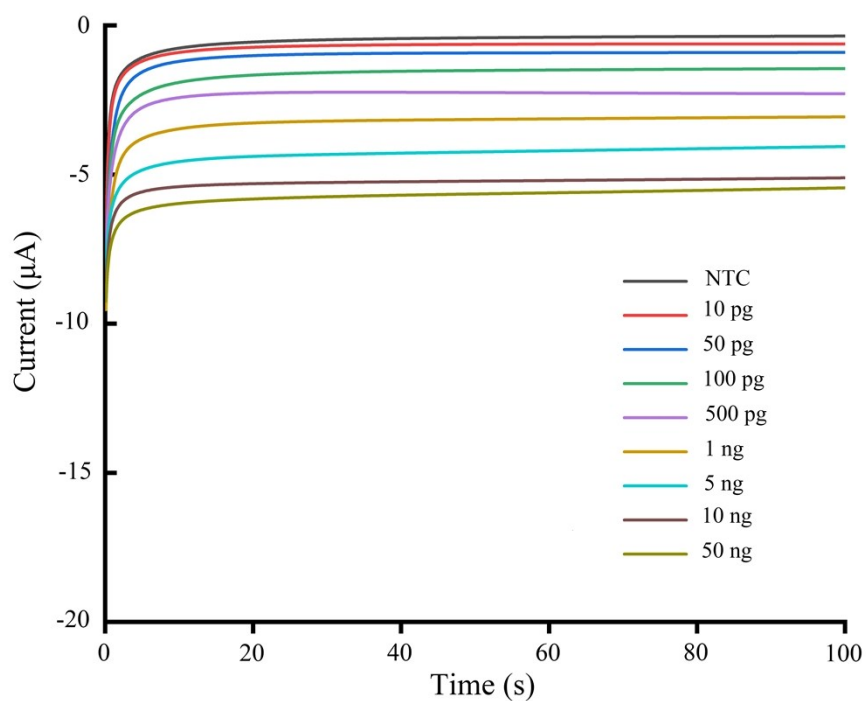


Figure S5. Current values of different amount of L858R DNA input. The signal gradually rises as more DNA is added.

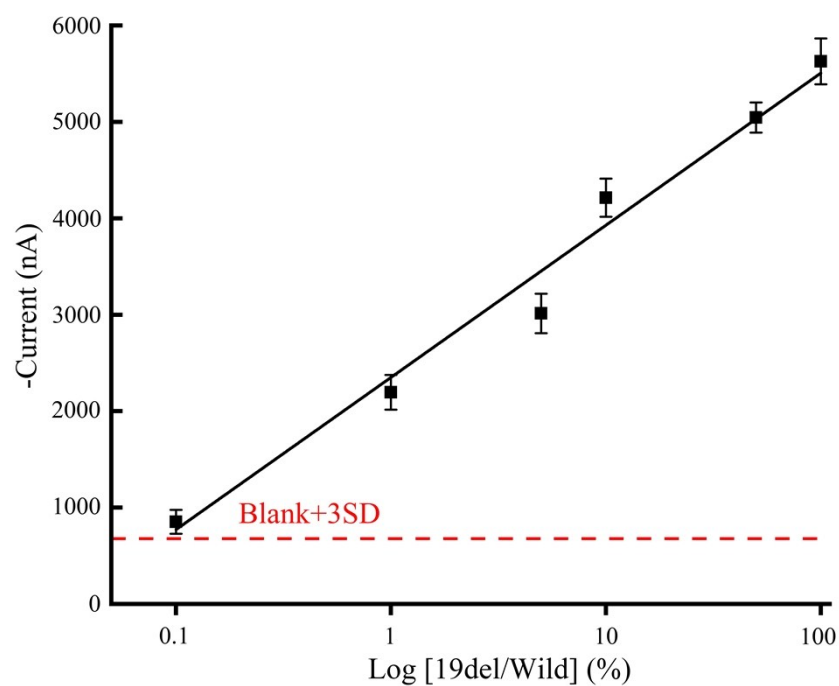


Figure S6. Chronoamperometric currents of the different percentages of 19del in the mixture.

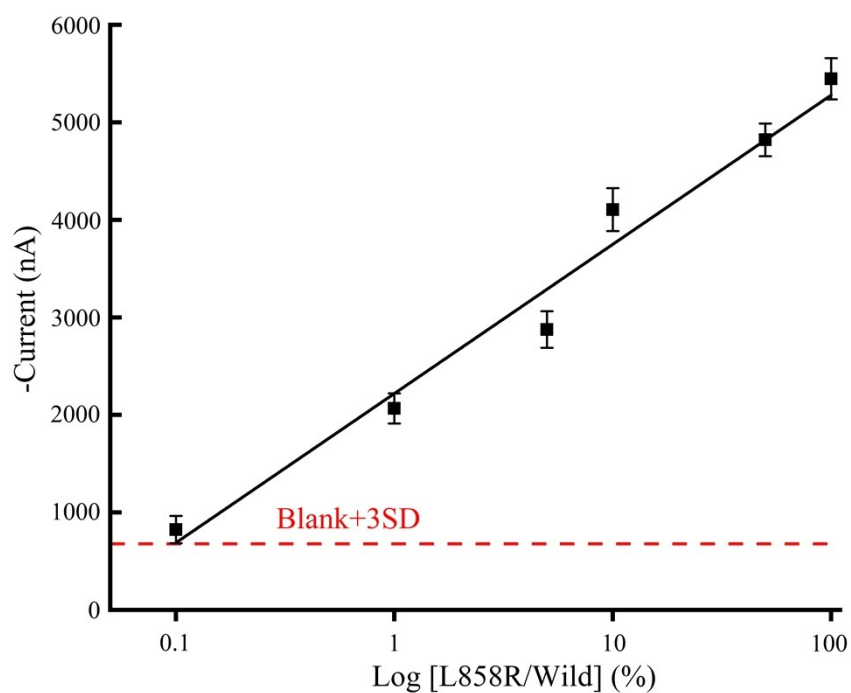


Figure S7. Chronoamperometric currents of the different percentages of L858R in the mixture.

Table S2. The reproducibility of the platform.

No.	1	2	3	SD	Average	RSD
Current (nA)	7523	7674	7390	142.1	7529	1.89%

Table S3. The list of abbreviations in the main text.

Full phrase	Abbreviations
Non-small cell lung cancer	NSCLC
Epidermal growth factor receptor tyrosine kinase inhibitors	EGFR-TKIs
Epidermal growth factor receptor	EGFR
Circulating tumor DNA	ctDNA
Next-generation sequencing	NGS
Amplification refractory mutation system	ARMS
Droplet digital PCR	dd-PCR
affordable, sensitive, specific, user-friendly, rapid and robust, equipment-free, and deliverable to end-users	ASSURED
Point-of-care testing	POCT
Hybridization chain reaction	HCR
Rolling circle amplification	RCA
Linear-After-The-Exponential	LATE
Single-stranded DNA	ssDNA
Screen-printed electrodes	SPE
3,3',5,5'-tetramethylbenzidine	TMB
Avidin-horseradish peroxidase	HRP
Cyclic voltammetry	CV
Fetal bovine serum	FBS
Three standard deviations	3SD
Non-template control	NTC