

# Electronic Supplementary Material

## Detection of Locus-Specific N<sup>6</sup>-Methyladenosine Modification Based on Ag<sup>+</sup>-Assisted Ligation and Supersandwich Signal Amplification

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**Table S1 DNA oligonucleotides used in the work.**

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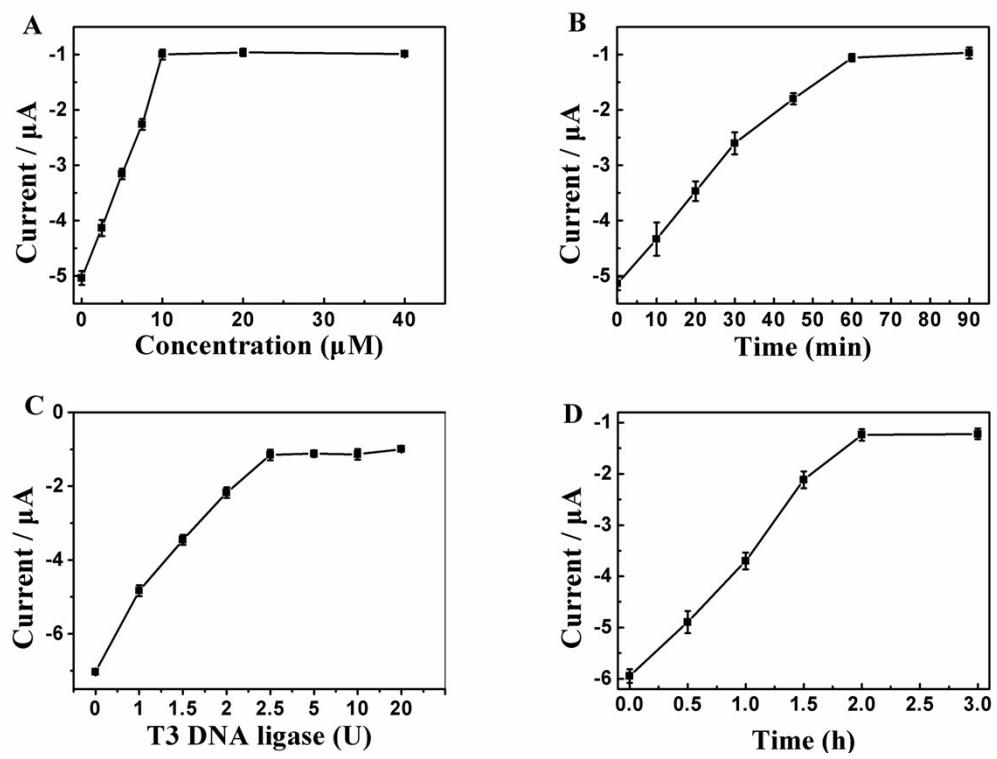
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Name	Sequences (5'-3')
I	SH-(CH <sub>2</sub> ) <sub>6</sub> -TCC GAG CCC GAC GCA TGA TCT GTA CTT GAC-OH
T	ATC AAG TAC AGA TCA TGC GTT GCA CGG TCG /6-MedA/TC AAG TAC AGA TCA TGC GTC GGG CTC GGA
TA	ATC AAG TAC AGA TCA TGC GTT GCA CGG TCG ATC AAG TAC AGA TCA TGC GTC GGG CTC GGA
P1	CGA CCG TGC AAC GCA TGA
P2	CGA CCG TGC AAC GCA TGA TCT GTA CTT GAT CCC TAA CCC TAA CCC TAA
H1	CCC TAA CCC TAA GTA GTG TTA GGG TTA GGG TTA GGG-MB
H2	MB-CAC TAC TTA GGG TTA GGG CCC TAA CCC TAA CCC TAA

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**Fig. S1.** The optimization of (A)  $\text{Ag}^+$  concentration. (B) The reaction time of  $\text{Ag}^+$  mediated base pairing of A and C. (C) T3 DNA ligase concentration. (D) The reaction time of T3 DNA ligase.

**Table S2. Comparison of the proposed method with other reported methods**

Detection strategy	Target	Sample source	Detection limit	Ref.
	analyte			
Optical	DNA	Human lung cancer	0.8 fM	33
	methylation			
Photoelectrochemical	DNA	Synthetic oligonucleotides	35 fM	34
	methylation			
Electrochemical	DNA	Synthetic CpG oligonucleotides	Not mentioned	35
	methylation	(20-mer and 60-mer)		
Liquid chromatography	Global DNA	Normal and tumor tissues of	0.05 pg	36
tandem mass spectrometry	methylation	colorectal cancer		
High-performance capillary electrophoresis	Global DNA	Methotrexate-resistant A549 cells	1 μM	37
	methylation			
Fluorimetric nanobiosensor	DNA	Sequence of APC gene	0.31 fM	38
	methylation			
	Synthetic			
Electrochemical	DNA	Synthetic oligonucleotides	44.427 pM	This work
	methylation			