

Phosphorothioated and Phosphate-Terminal Dumbbell (PP-TD) Probe-Based Rapid Detection of Polynucleotide Kinase Activity

Xianfeng Jiang,^a Xudan Shen,^b Jadera Talap,^b Dan Yang,^b Su Zeng,^b Hui Liu,^{*a} and Sheng Cai^{*b}

^a Sir Run Run Shaw Hospital, School of Medicine, Zhejiang University, Hangzhou, Zhejiang 310020, China.

^b Institute of Drug Metabolism and Pharmaceutical Analysis, Zhejiang Province Key Laboratory of Anti-Cancer Drug Research, College of Pharmaceutical Science, Zhejiang University, Hangzhou, Zhejiang 310058, China.

* Corresponding author, E-mail: caisheng@zju.edu.cn, lhui2010@zju.edu.cn.

Table S1. Sequence information in this study

Name	Sequences
PP-TD Probe	5'- <u>AAGAATTCTTAAGAATTCTT</u> ACATCGTTACGTTAGTGAAAAACCACTAAC-3'P
P-TD Probe	5'-AAGAATTCTTAAGAATTCTTACATCGTTACGTTAGTGAAAAACCACTAAC-3'OH
Loop-6	5'-AAGAATTCTTAAGAATTCTTACATCGTTACGTTAGTGAAAAAACCACTAAC-3'P
Loop-8	5'- <u>AAGAATTCTTAAGAATTCTT</u> ACATCGTTACGTTAGTGAAAAAACCACTAAC-3'P
Bridge-8	5'- <u>AAGAATTCTTAAGAATTCTT</u> ATCGTTACGTTAGTGAAAAACCACTAAC-3'P
Bridge-12	5'-AAGAATTCTTAAGAATTCTTAAACATCGTTACGTTAGTGAAAAACCACTAAC-3'P

* Underlined bases are phosphorothioate modified.

Table S2. Comparison with other reported fluorescence methods

Reference	Time (min)	Linear Range (U/ μ L)	LOD (U/ μ L)
(Li, Xu et al. 2017)	335	1.0×10^{-7} to 1.0×10^{-3}	4.36×10^{-8}
(Zhao, Liu et al. 2017)	280	5.0×10^{-4} to 5.0×10^{-2}	2.0×10^{-4}
(Feng, Wang et al. 2018)	130	1.0×10^{-5} to 3.0×10^{-4}	6.7×10^{-6}
(Chen, Wang et al. 2019)	185	1.0×10^{-6} to 1.0×10^{-5}	7.9×10^{-7}
(Jie, Li et al. 2019)	250	1.0×10^{-6} to 1.0×10^{-2}	1.0×10^{-6}
(Li, Ma et al. 2019)	60	1.0×10^{-5} to 1.25×10^{-2}	1.0×10^{-5}
(Zhang, Liu et al. 2019)	190	7.0×10^{-5} to 1.5×10^{-2}	6.0×10^{-5}
(Zhou, Tong et al. 2019)	200	1.0×10^{-3} to 1.0×10^{-1}	2.0×10^{-4}
(Wang, Wang et al. 2020)	80	1.0×10^{-8} to 2.5×10^{-5}	3.3×10^{-9}
(Wu, He et al. 2020)	30	2.5×10^{-4} to 1.0×10^{-3}	1.1×10^{-4}
(Zhang, Fan et al. 2021)	180	5.0×10^{-10} to 5.0×10^{-6}	1.2×10^{-10}
This work	105	1.0×10^{-6} to 5.0×10^{-5}	7.7×10^{-7}

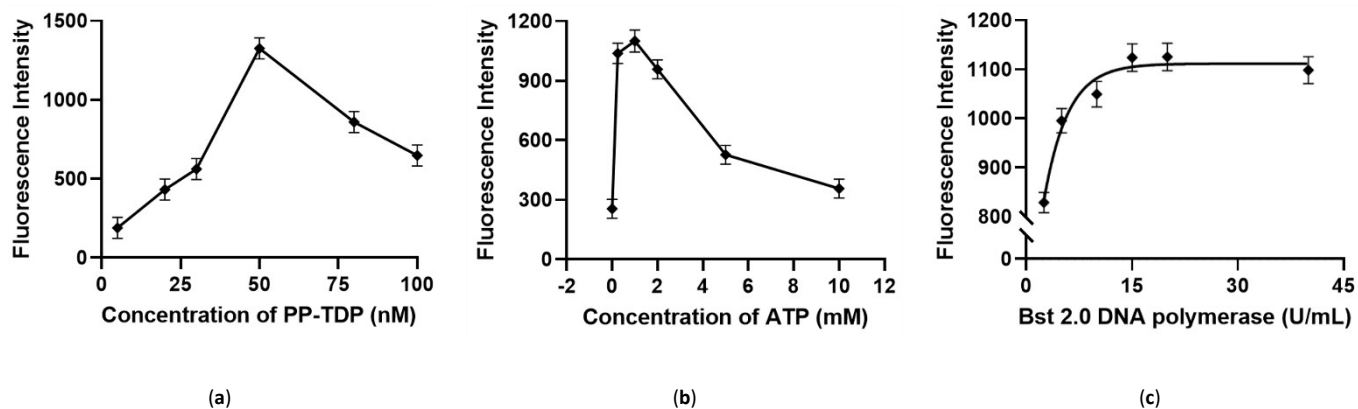


Figure S1. Optimization of the reaction conditions. (a) Fluorescence intensity vs. Concentration of PP-TD probe; (b) Fluorescence intensity vs. Concentration of ATP; (c) Fluorescence intensity vs. Concentration of Bst 2.0 DNA polymerase. The statistical data is presented as mean \pm S.D (n=3).

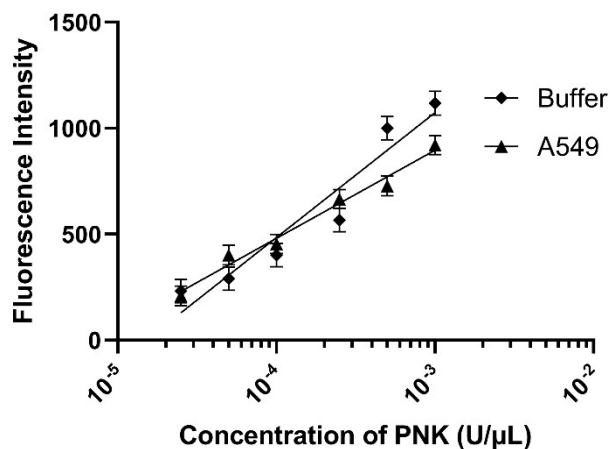


Figure S2. Quantitative detection in diluted cell extracts. The statistical data is presented as mean \pm S.D (n=3).

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