

Supporting Information

Real-time in situ fluorescence imaging of telomerase and miR378 in living cells using two-color DNA tetrahedral nanoprobe combined with molecular beacons

Jiejie Guang ^{a,1}, Shan Wang ^{a,1}, Bingyuan Fan ^a, Ziyao Yu ^b, Yahui Gao ^a, Jinru Pan ^a, Junting Xi ^{c,*}, Wei Meng ^{a,*}, Fang Hu ^{a,*}

Table S1 Nucleic acid sequences.

| Name | Sequences (5'-3') |
|--------|---|
| L1 | ACATTCCTAAGTCTGAAACATTACAGCTTGCTACACGA GAAGAGCCGCCATAGTATTTTGAGGTAAGGACTGC |
| L2-1 | TATCACCAGGCAGTTGACAGTGTAGCAAGCTGTAATTT TT-Cy3 |
| L2-2 | BBQ-650- TTTTAAGAATAACCCTAACCCTAACCCTAACCC |
| L3 | TCAACTGCCTGGTGATAAAACGACACtACgTGGGAATCT ACTATGGCGGCTCTTCTTTTAATCCGTCGAGCAGAGTT |
| L4 | TCGCGATAGCAGGAATGTGCTTCCCACGTAGTGTCGTTT GGGTTAGGGTTATTCTTCATTCAGACTTGCTATCGCGA |
| MB | FAM-CACCGATATACCTTCTGACTCCAAGTCCCGGTG /iBHQ1dT/GCAG TCCTTACCTC |
| miR378 | ACUGGACUUGGAGUCAGAAGG |
| mis1 | ACUGGACUUGGAGUGAGAAGG |
| mis2 | ACUGGACAUGGAGUCACAAGG |
| mis3 | ACUGGACAUGGUGUCACAAGG |
| NC | UUGUACUACAAAAGUACUG |
| miR21 | UAGCUUAUCAGACUGAUGUUGA |
| miR155 | UUA AUGCUAAUCGUGAUAGGGGU |

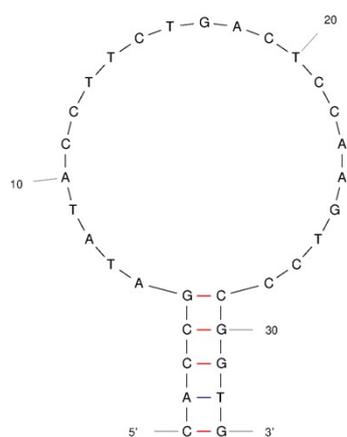


Fig.S1 Secondary structure of the most suitable MB was simulated using mFold online software;
 Sequence: CACCGATATACCTTCTGACTCCAAGTCCCGGTG ($\Delta G = -2.20$)

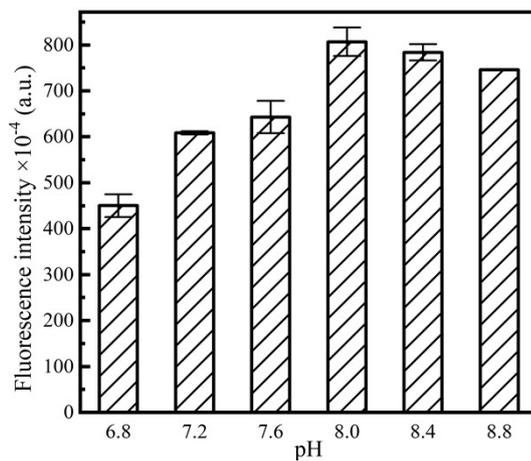


Fig.S2 Comparing fluorescence differences at different pH levels

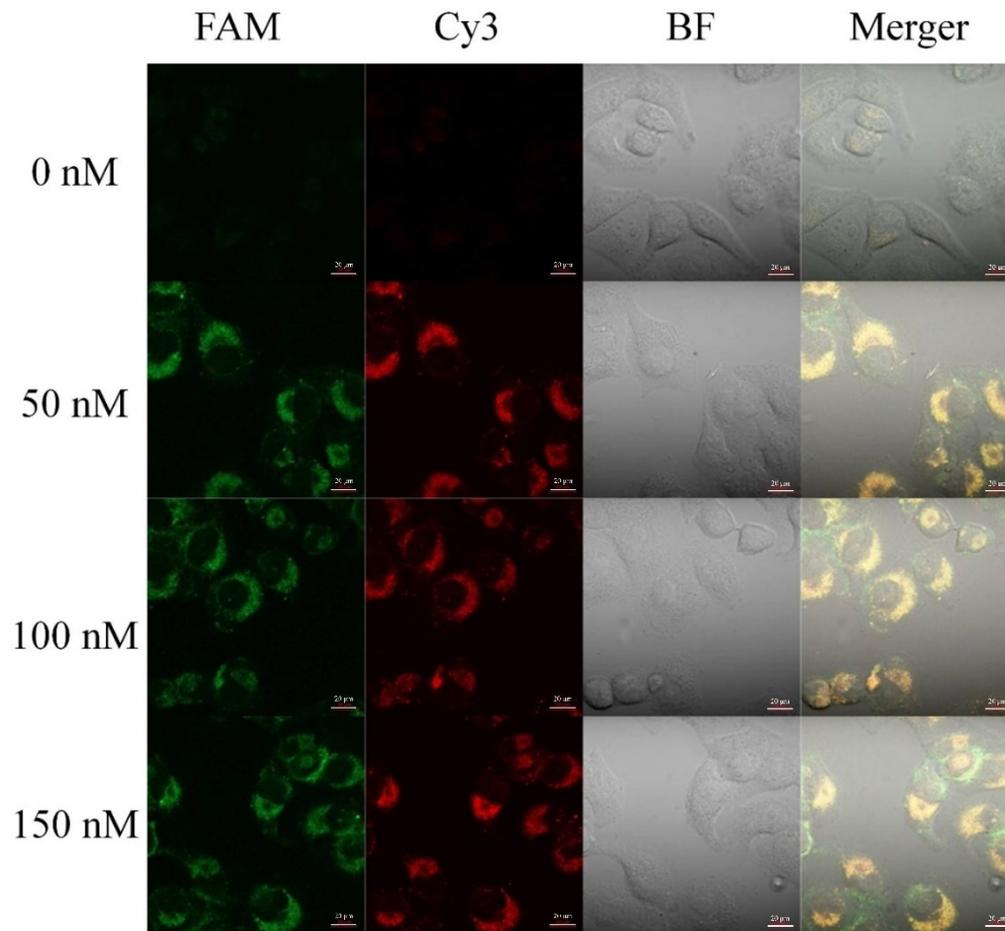


Fig.S3 Confocal fluorescence images of different DTP concentrations incubated with HeLa cells for 9 h; scale: 20 μ m

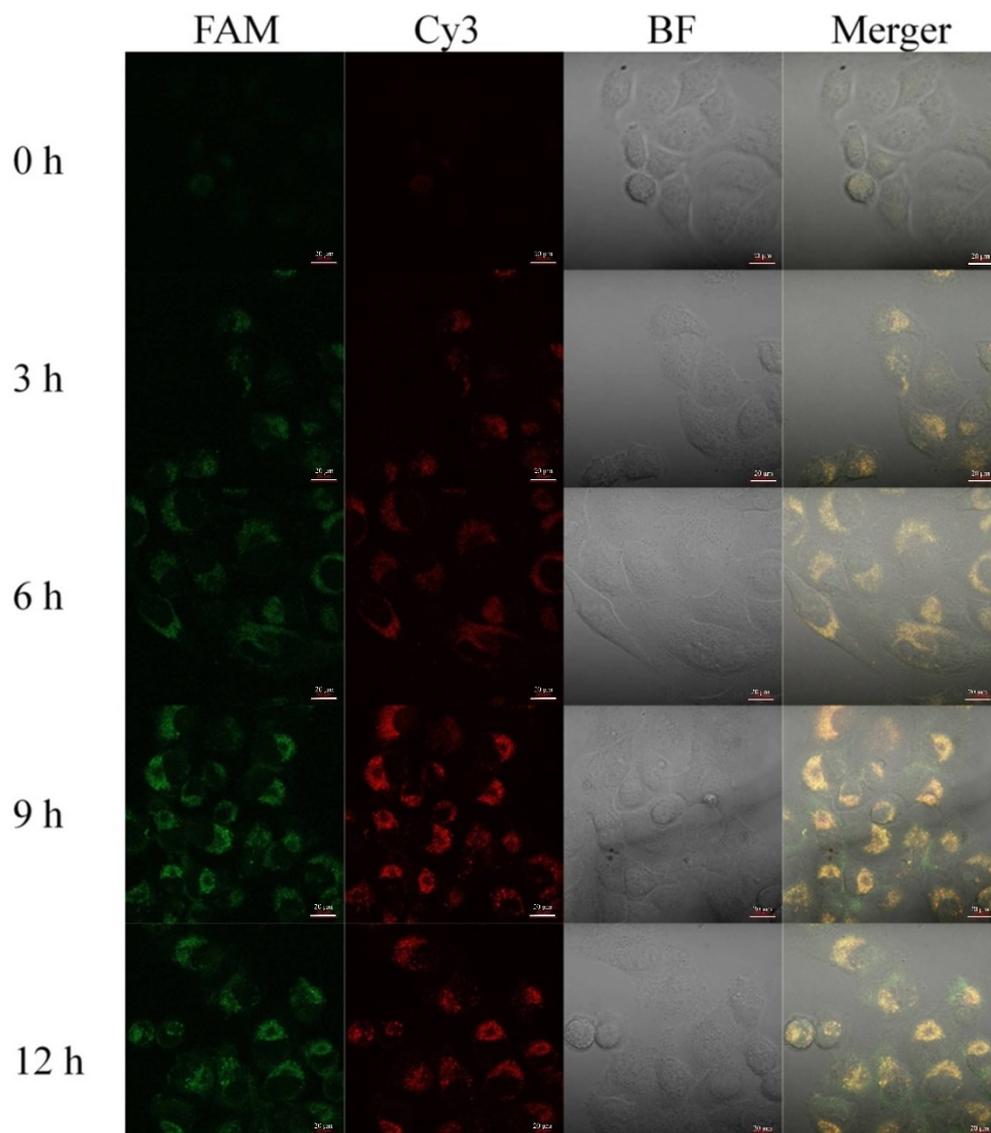


Fig.S4 Confocal fluorescence images of MB-DNA tetrahedral probe incubated with HeLa cells for different durations; scale: 20 μM