Fluorescent Conjugated Polyelectrolyte As An Indicator for

Convenient Detection of DNA Methylation

Fude Feng¹, Hongzhong Wang², Lingli Han², Shu Wang¹*

¹Beijing National Laboratory for Molecular Sciences, Key Laboratory of Organic Solids,

Institute of Chemistry, Chinese Academy of Sciences, Beijing 100080, P. R. China.

Email: wangshu@iccas.ac.cn

² Department of Biological Sciences and Biotechnology, Tsinghua University, Beijing 100084, P. R. China.

Supporting Information

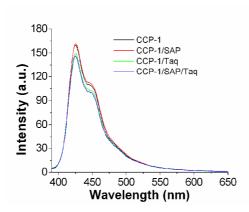


Figure S1. Fluorescence spectra from Hepes buffer solutions (25 mM, pH 8.0) containing (a) CCP-1 (0.25 μ M), (b) CCP-1 (0.25 μ M), SAP (0.17 U/mL), (c) CCP-1 (0.25 μ M), Taq (0.21 U/mL), (d) CCP-1 (0.25 μ M), SAP (0.17 U/mL) and Taq (0.21 U/mL). The excitation wavelength was 380 nm.

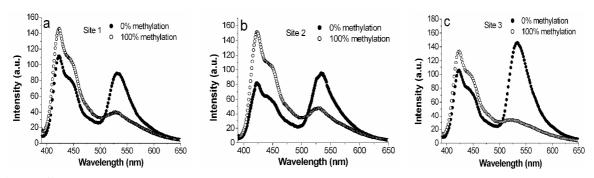


Figure S2. Fluorescence spectra (a-c) from solutions containing CCP-1 and single nucleotide base extension products of methylated plasmid and unmethylated plasmid using unmethylation-specific probes in 1 mL of HEPES buffer solution (25 mM, pH 8). The probes

used in extension reactions were p16-1n for site 1, p16-2n for site 2 and p16-3n for site 3. The amount: 0.67 pmol probe, 1.67 pmol dGTP-Fl, [CCP-1] = 0.25 μ M in RUs. The excitation wavelength is 380 nm.

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1 attcaagage taacaggtat tagettagga tgtgtggcae tgttcttaag gettatatgt
  61 attaatacat catttaaact cacaacaacc cctataaagc agggggcact catattccct
 121 tececetta taattaegaa aaatgeaagg tatttteagt aggaaagaga aatgtgagaa
 181 gtgtgaagga gacaggacag tatttgaagc tggtctttgg atcactgtgc aactctgctt
 241 ctagaacact gagcactttt tctggtctag gaattatgac tttgagaatg gagtccgtcc
 301 ttccaatgac tccctcccca ttttcctatc tgcctacagg cagaattctc ccccgtccgt
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 481 gcagtccgac tctccaaaag gaatcctttg aactagggtt tctgacttag tgaaccccgc
 541 geteetgaaa ateaagggtt gagggggtag ggggacaett tettagtegt acaggtgatt
601 tcgatttctc ggtggggctc tcacaactag gaaagaatag ttttgctttt tcttatgatt
 661 aaaagaagaa gccatacttt ccctatgaca ccaaacaccc cgattcaatt tggcagttag
 721 gaaggttgta tcgcggagga aggaaacggg gcgggggcgg atttcttttt aacagagtga
 781 acgcactcaa acacgccttt getggcagge gggggagege ggctgggage agggaggccg
 841 gagggcggtg tggggggcag gtggggagga gcccagtcct ccttccttgc caacgctggc
                                                                         Site 2
 901 tetggcgagg getgetteeg getggtgeee eegggggaga eccaacetgg ggcgacttea
 961 ggggtgccac att CG ctaag tgctcggagt taatagcacc tcctccgage actcgctcac
1021 agcgtcccct tgcctggaaa gatacCGcgg tccctccaga ggatttgagg gacagggtCG
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1141 ggtgggcgg accgagtgcg ctcggcggct gcggagaggg gtagagcagg cagcgggcgg
1201 cggggagcag catggagccg gcggcggga gcagcatgga gccttcggct gactggctgg
1261 ccacggccgc gg
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Scheme-S1. The sequence of the *p16* promoter region of human colon cancer cell line HT29 containing the three methylated CpG sites and the positions of first and second rounds of PCR.

Completed Reference 14 in the text

Bibikova, M.; Lin, Z.; Zhou, L.; Chudin, E.; Garcia, E. W.; Wu, B.; Doucet, D.; Thomas, N. J.; Wang, Y.; Vollmer, E.; Goldmann, T.; Seifart, C.; Jiang, W.; Barker, D. L.; Chee, M. S.; Floros, J.; Fan, J.-B. *Genome Res.* **2006**, *16*, 383.