

Supplementary Information

Naphthalene diimides carrying two β -cyclodextrins prefer Telomere RNA G-quadruplex recognition

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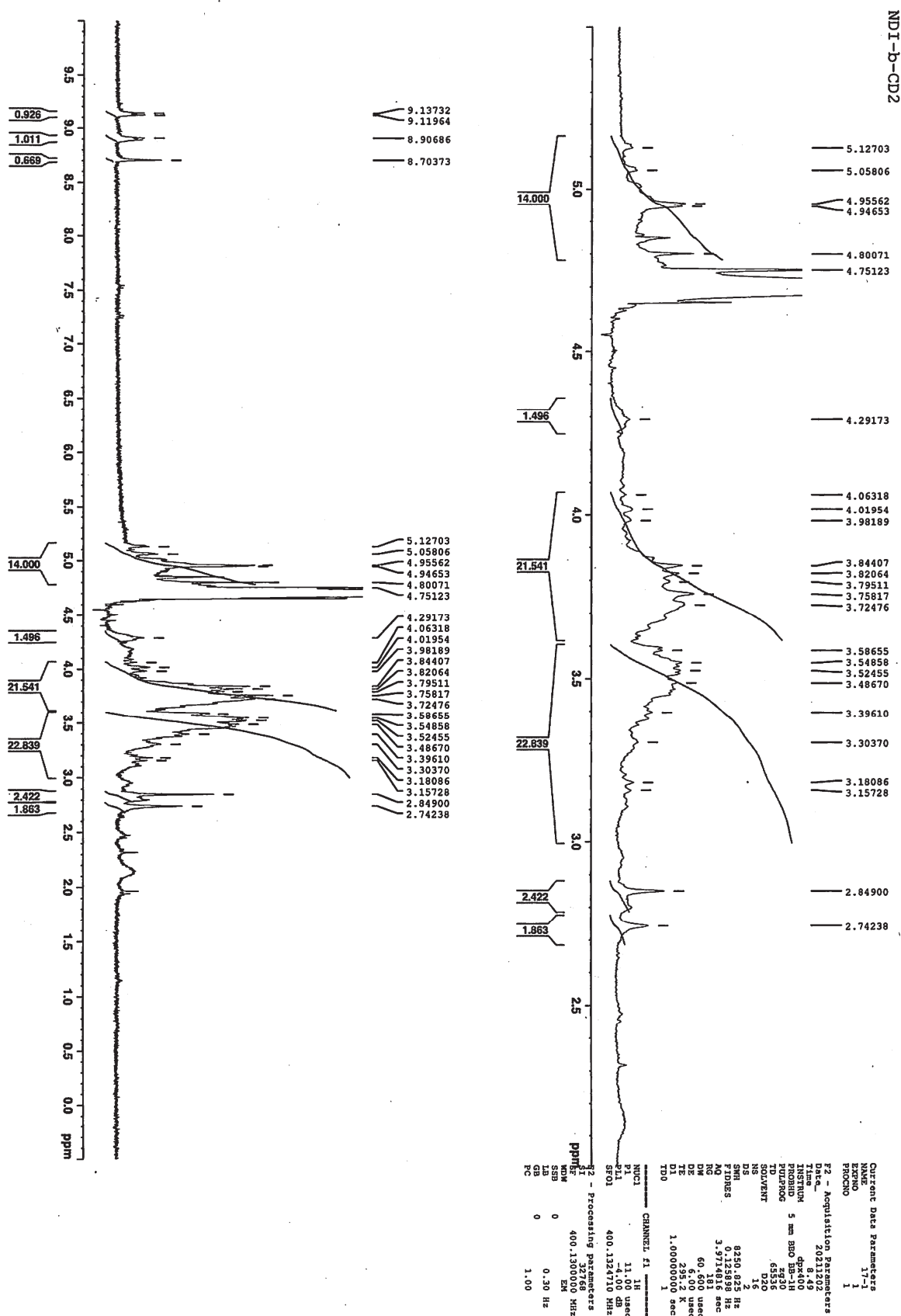


Figure S1. ¹H-NMR spectra of 1 in D₂O.

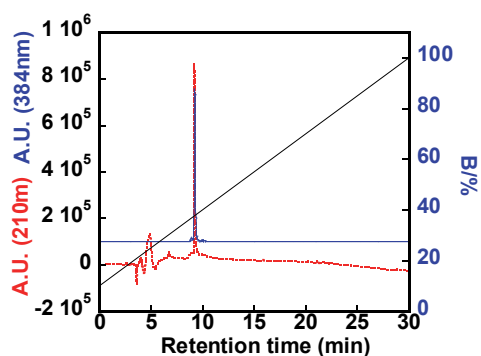


Figure S2. RT-HPLC of **1**. HPLC conditions; the concentration of solution B was changed (from 10% to 100%, 30 min) at 40 °C. Solution A: 0.1% trifluoroacetic acid, Solution B: 70% acetonitrile, 0.1% trifluoroacetic acid. Flow rate: 1 mL/min. Column: Inertsil ODS-4 (GL Sciences Inc., Tokyo, Japan).

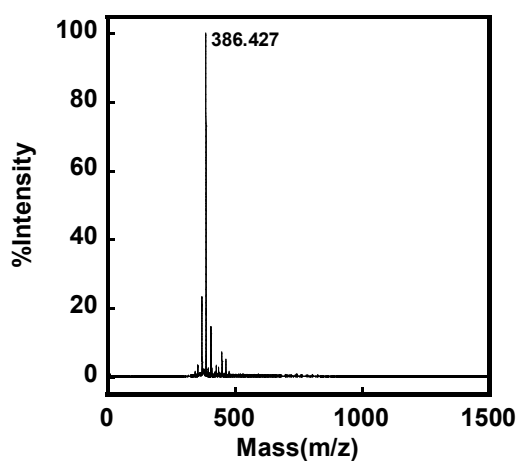


Figure S3. MALDI-TOF-MS spectra of **4** under positive mode (Matrix: DHBA). $m/z = 386.427$ (calculated value of $C_{18}H_{10}N_2O_8 + H^+ = 383.289$).

ndi-gly

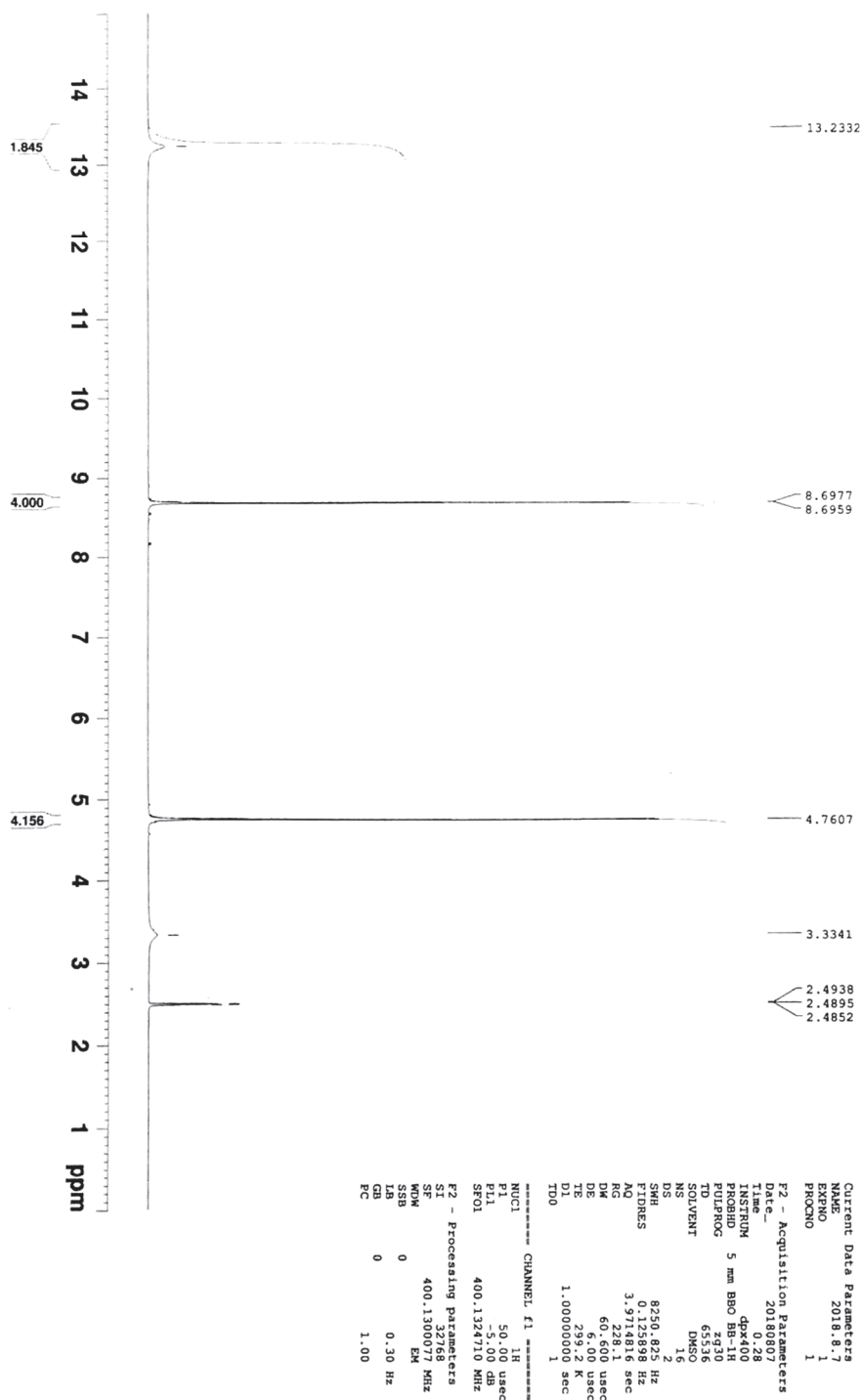


Figure S4. ¹H-NMR spectra of **4** in DMSO.

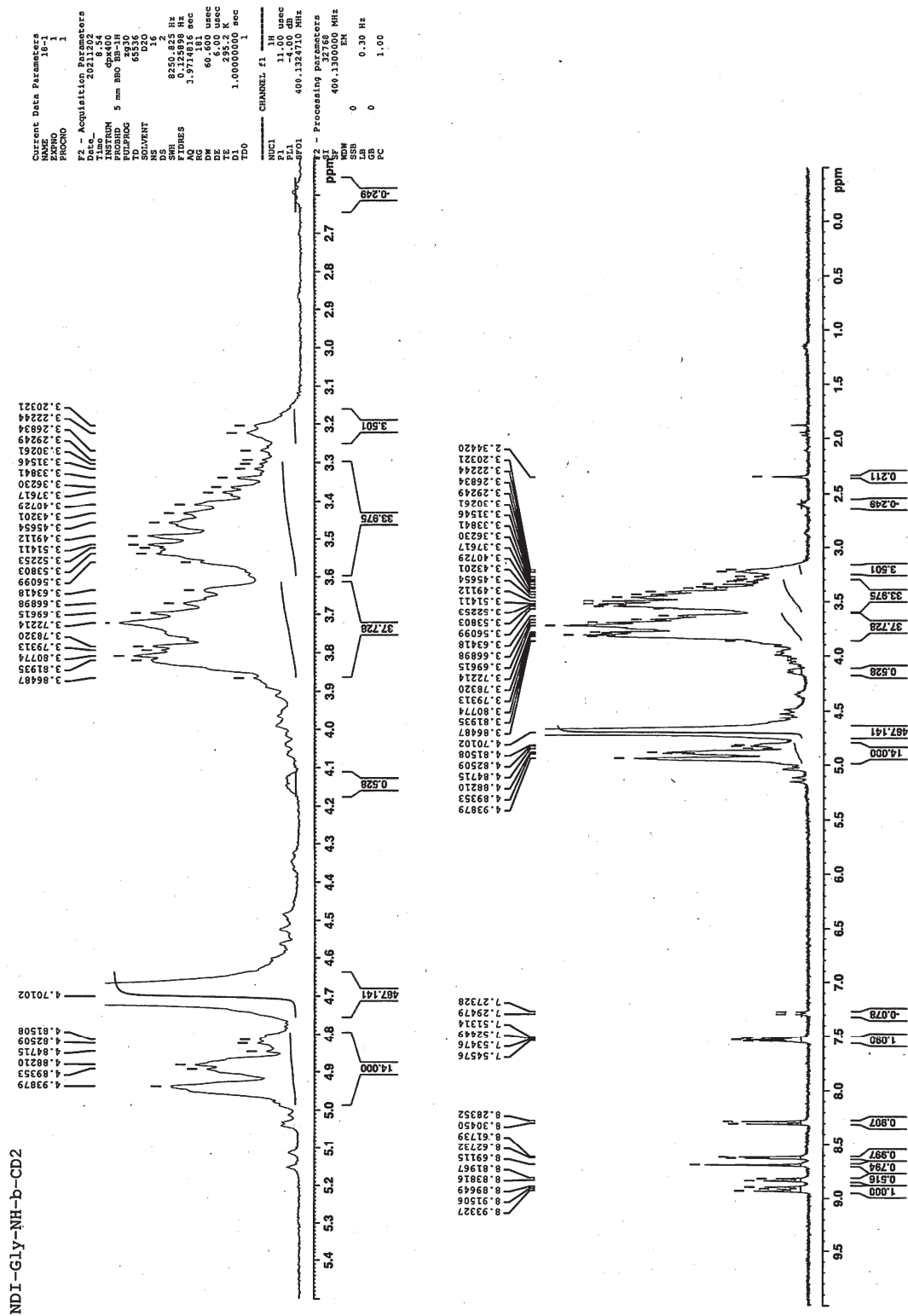


Figure S5. ^1H -NMR spectra of **2** in D_2O .

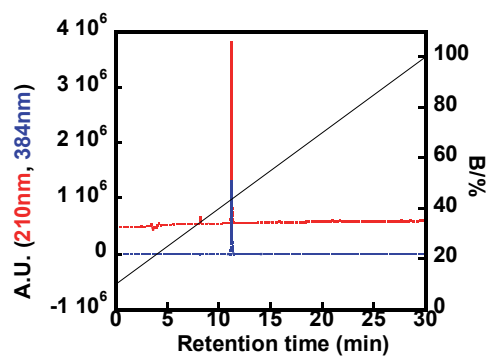


Figure S6. RT-HPLC of 2. HPLC conditions; the concentration of solution B was changed (from 10% to 100%, 30 min) at 40 °C. Solution A: 0.1% trifluoroacetic acid, Solution B: 70% acetonitrile, 0.1% trifluoroacetic acid. Flow rate: 1 mL/min. Column: Inertsil ODS-4 (GL Sciences Inc., Tokyo, Japan).

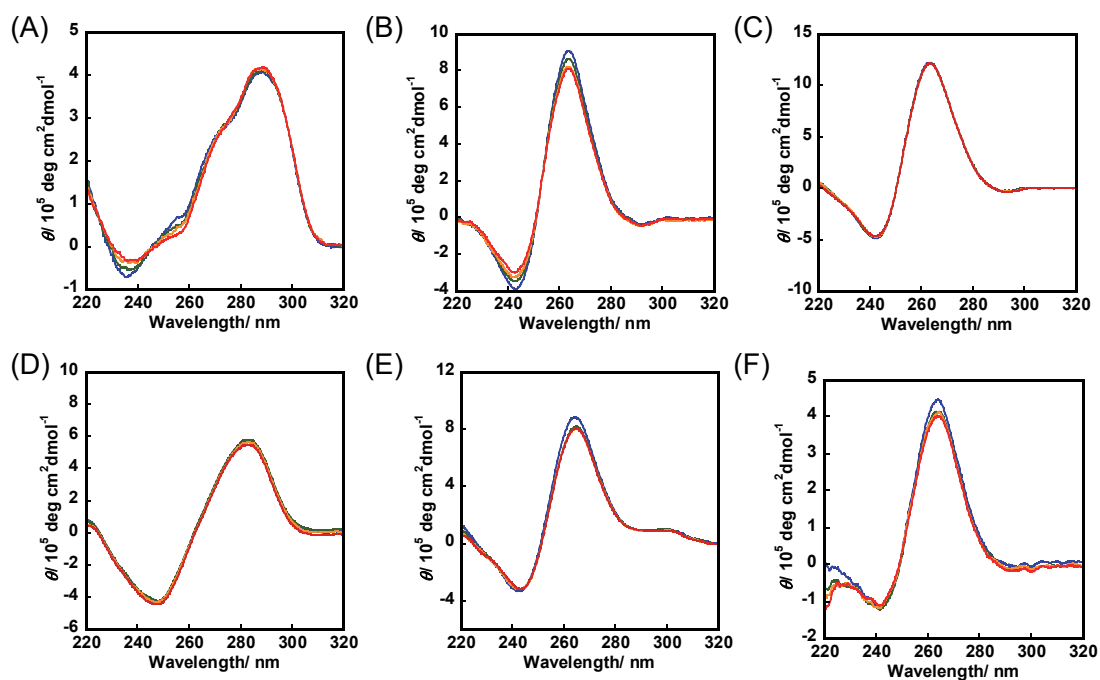


Figure S7. CD spectra of (A) Telomere G1, (B) c-myc, (C) c-kit, (D) ds-oligo, (E) G1-r23t, and (F) G1-r12t after adding 2. 1.5 μ M DNA, 0-4.5 μ M 2, 50 mM Tris-HCl buffer (pH 7.4) with 100 mM KCl at 25 $^{\circ}$ C. [DNA] : [2]=1:0 (blue), 1:1 (green), 1:2 (orange), 1:3 (red).

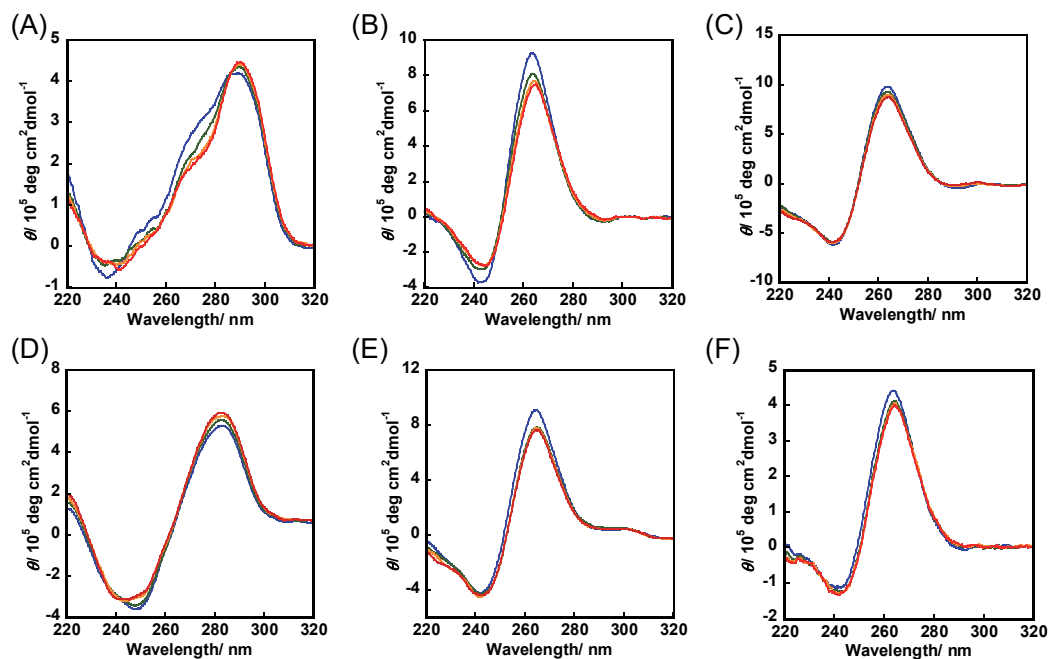


Figure S8. CD spectra of (A) Telomere G1, (B) c-myc, (C) c-kit, (D) ds-oligo, (E) G1-r23t, and (F) G1-r12t after adding 3. 1.5 μM DNA, 0-4.5 μM 3, 50 mM Tris-HCl buffer (pH 7.4) with 100 mM KCl at 25 $^{\circ}\text{C}$. [DNA] : [3]=1:0 (blue), 1:1 (green), 1:2 (orange), 1:3 (red).

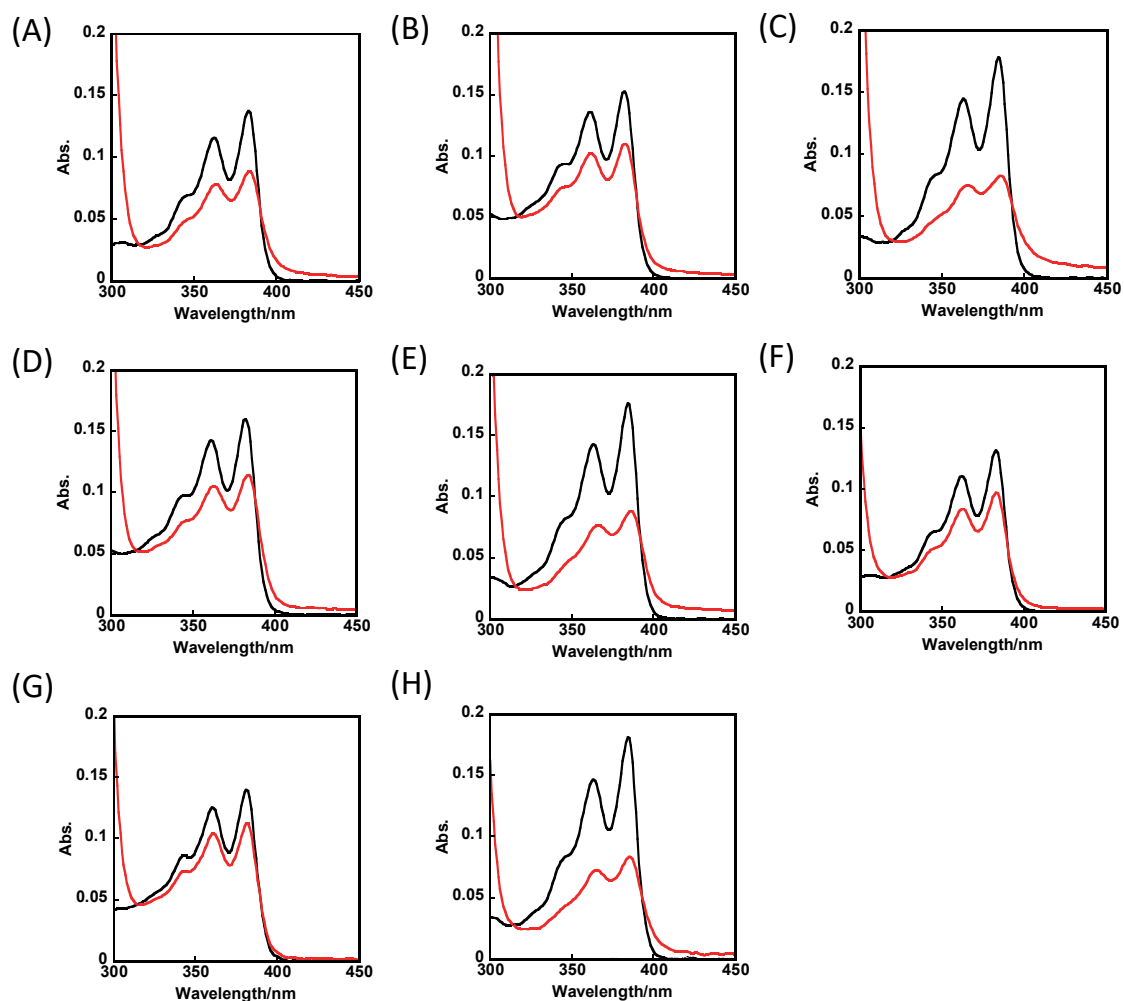


Figure S9. UV-Vis spectra of (A) **1** with Telomere G1, (B) **2** with Telomere G1, (C) **3** with Telomere G1, (D) **2** with c-myc, (E) **3** with c-myc, (F) **1** with ds-oligo, (G) **2** with ds-oligo, and (H) **3** with ds-oligo in 50 mM Tris-HCl buffer (pH 7.4) with 100 mM KCl at 25 °C (Before adding DNA: black line, after adding DNA: red line).

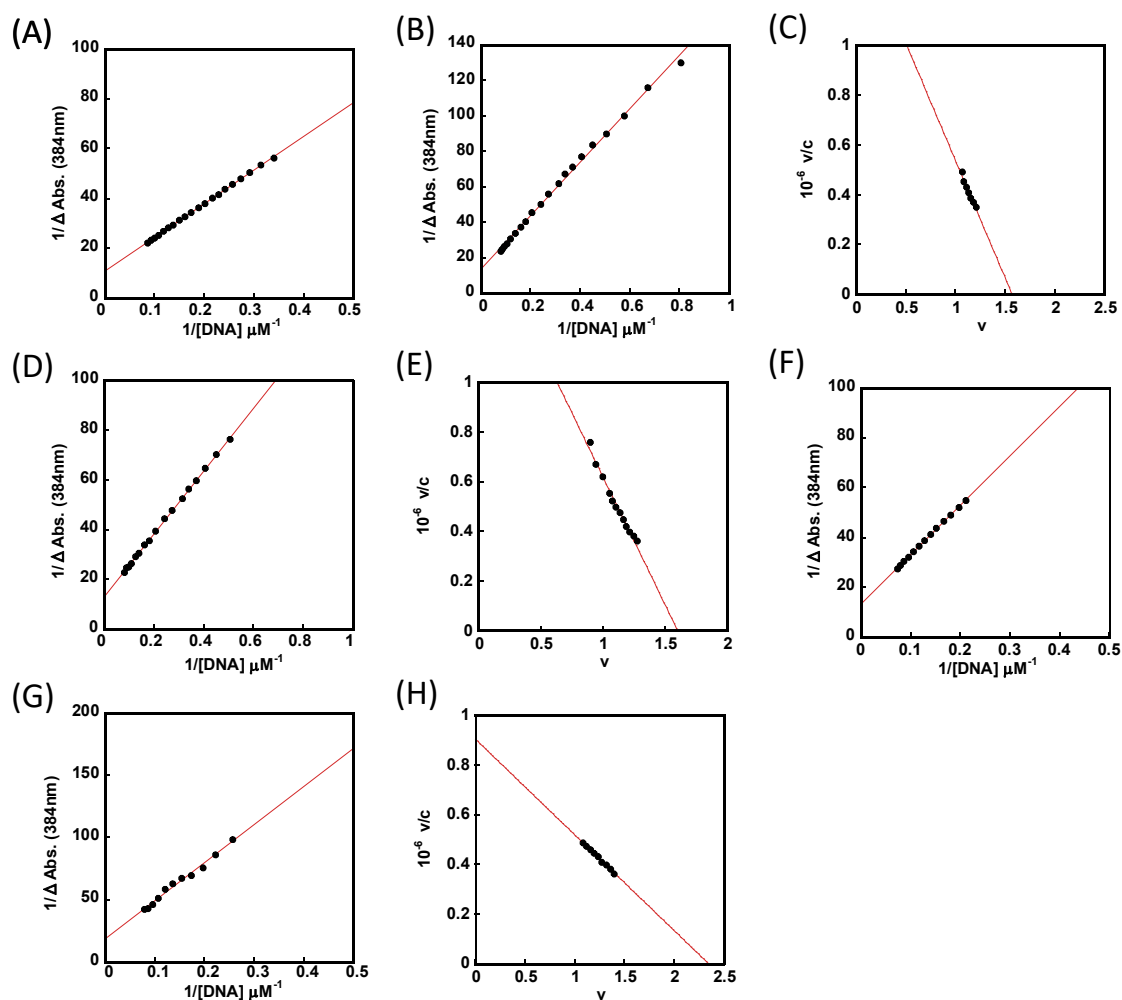


Figure S10. Binding analysis of (A) **1** with Telomere G1 (Benesi-Hildebrand analysis), (B) **2** with Telomere G1 (Benesi-Hildebrand analysis), (C) **3** with Telomere G1 (Scatchard analysis), (D) **2** with c-myc (Benesi-Hildebrand analysis), (E) **3** with c-myc (Scatchard analysis), (F) **1** with ds-oligo (Benesi-Hildebrand analysis), (G) **2** with ds-oligo (Benesi-Hildebrand analysis), and (H) **3** with ds-oligo (Scatchard analysis) in 50 mM Tris-HCl buffer (pH 7.4) with 100 mM KCl at 25 °C.

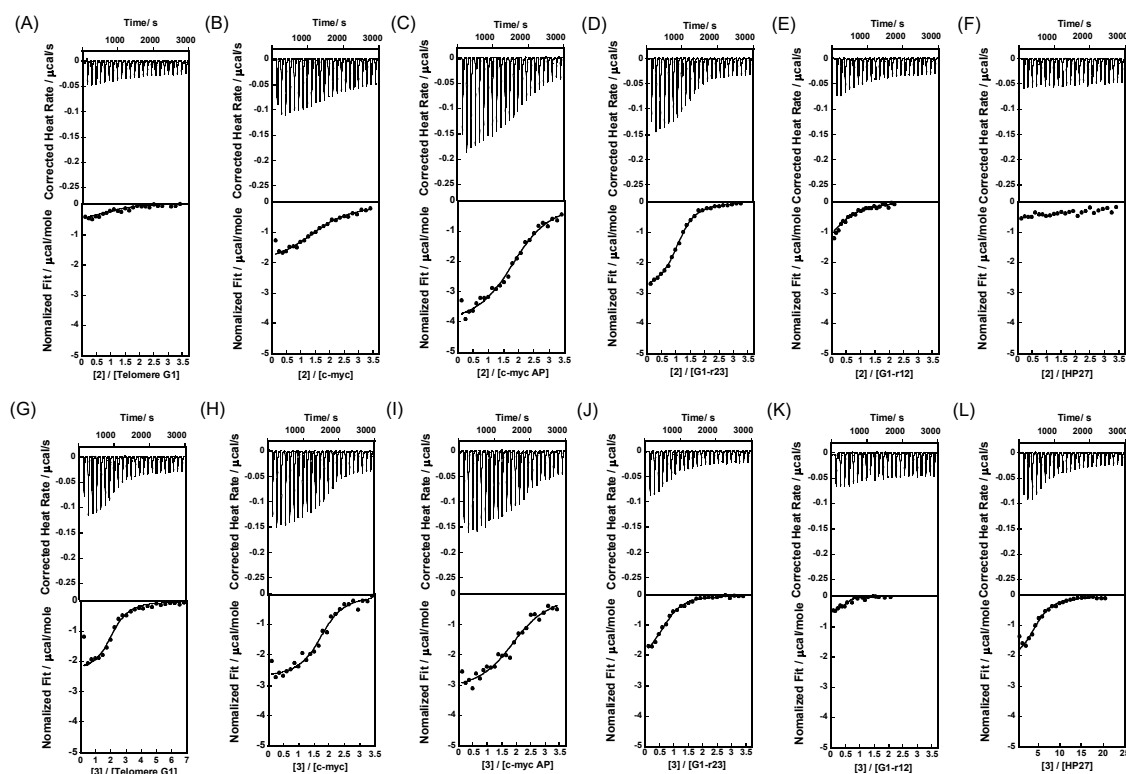


Figure S11. ITC measurement of (A, G) Telomere G1, (B, H) c-myc, (C, I) c-myc AP site, (D, J) HP27, (E, K) G1-r23nt and (F, L) HP27 with adding **2** (A-F) or **3** (G-L) in 50 mM KH_2PO_4 - K_2HPO_4 buffer (pH 7.0), 25 °C.

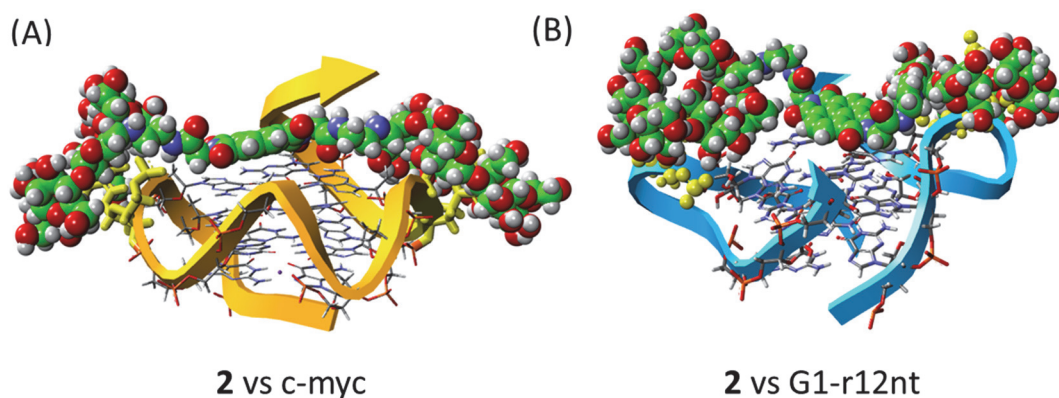


Figure S12. Binding model simulation of NDI- β -CyDs conjugator **2** with (A) *c-myc* G-quadruplex (PDB:1xav); (B) 12nt of telomere RNA G-quadruplex (PDB: 2kbp).