Supporting Information for:

Alternative DNA structure formation in the mutagenic human *c-MYC* promoter

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SI-1. Predicted folding topologies under triplex/H-DNA-forming conditions.



SI-2. Similar gel mobilities were observed with R2 independent of DNA concentration.



PAGE gel of R2 and R2-ino with Mg²⁺.

1) MWM (10 bp ladder)
2) R2-ino (1 μM)
3) R2 (1 μM)
4) R2 (0.2 μM)
5) R2 (1 μM)
6) R2 (20 μM)
7) R2 (99 μM)

SI- 3. K⁺ titration into known G4-DNA-forming sequences initially annealed under H-DNA-forming conditions (G4-Htel27, antiparallel G4-DNA; G4-cMyc22, parallel G4-DNA).



G4-Htel27 : 5'-TTA GGG TTA GGG TTA GGG TTA GGG TTA-3' G4-cMyc22 : 5'-TGA GGG TGG GTA GGG TGG GTT A-3'

SI- 4. CD structures obtained at 25°C are structure-related. The structures at 90°C are similarly in random coil/denatured states.



SI- 5. Native gel and CD spectroscopy results under Dulbecco's Phosphate Buffered Saline (DPBS) showing that motifs **A**) also folded intramolecularly and **B**) possess the triplex CD signature.



SI- 6. Comparison of the temperature-dependent fluorescence of AP-A33hg under G4-DNAforming conditions with 2AP-substituted, control G4-DNA-forming sequences (AP12-myc22, AP16-myc22, AP-Tel). AP-A33hg was A) only 30% as intense but B) had a similar temperaturedependent profile as that of AP16-myc22. Intensity comparison suggests that 2AP may participate in a 2-nt loop.



AP16-myc22: 5'-TGA GGG T GGG TA GGG 2AP GGG TT A-3' AP12-myc22: 5'-TGA GGG T GGG T2AP GGG T GGG TT A-3' 2-nt loop AP-Tel: 5'-GGGG TT GGGG T2AP <u>G</u>GGG TT GGGG-3' AP-33hg: CCCCTCCC TTTTT GGGAGGGG CGCTTAT GGG<u>G</u>2AP<u>G</u>GG

2-4-nt loop 1-2-nt loop **SI-7.** A representative denaturing gel image after chemical modification of R2-5' (primary data) under H-DNA/triplex- or G4-DNA-forming conditions. Lane 1=untreated; lanes 2-5=triplex conditions; lanes 6-7=G4 conditions; lanes 2 and 4 (underlined)=boiled samples. Average % protection was calculated as described in the Methods section.



SI- 8. Polyacrylamide gel electrophoresis of structure-forming DNA in the **A**) presence and **B**) absence of Mg^{2+} . **C**) CD spectroscopy results show the presence of the triplex signature (negative peak at ~220 nm) in the presence and absence of Mg^{2+} .

