Electronic supplementary information

Design and synthesis of purine nucleoside analogues for the formation of stable anti-paralleltype triplex DNA with duplex DNA bearing the ^{5m}CG base pair

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Figure S1 ¹H-NMR and ¹³C-NMR spectra of new compounds 5-9.













Figure S2 ¹H-NMR, ¹³C-NMR and ³¹P-NMR spectra of new compounds 10, 11 and 12.

Table S1. MALDI-TOF MS results for synthesized TFOs.^a

dAN derivatives	sequences	calcd. [M-H] ⁻	found
Aminoethyl-dAN	3'-GZA-5'	5791.06	5794.29
	3'-GZG-5'	5807.05	5809.70
	3'-AZG-5'	5791.06	5792.84
	3'-AZA-5'	5775.06	5777.47
	3'-GZA-5'	5840.04	5840.21
2 Dhonol dAN	3'-GZG-5'	5856.04	5855.39
5-FIIEIIOI-UAN	3'-AZG-5'	5840.04	5840.13
	3'-AZA-5'	5824.05	5822.79
	3'-GZA-5'	5840.04	5841.41
2 Dhonol dAN	3'-GZG-5'	5856.04	5856.98
2-Phenoi-uAn	3'-AZG-5'	5840.04	5841.94
	3'-AZA-5'	5824.05	5824.54

^a The structural integrity of all synthesized TFOs (3' GGA AGG *NZN'* GAG GAG GGA 5') was analyzed by MALDI-TOF MS (m/z). *NZN'* = GZA, GZG, AZG and AZA.

Figure S3 MALDI-TOF MS charts of synthesized TFOs having dAN derivatives.

3'-G Aminothyl-dAN A-5'

3'-G 3Phenol-dAN A-5'

3'-G 3Phenol-dAN G-5'

3'-A 3Phenol-dAN G-5'

3'-A 3Phenol-dAN A-5'

Figure S5 Gel results of confirmation of triplex formation.

Conditions: The mixture of FAM-labeled Py-strand (24 mer; 100 nM) or Pu-strand (24 mer; 100 nM) or TFO ((A) and (B): 18 mer; 0-1000 nM or (C): 500 nM) was incubated in buffer containing 20 mM Tris-HCl and 20 mM MgCl₂ at 37°C and pH 7.5. Electrophoresis was performed using a 10% non-denatured polyacrylamide gel. (A) Target duplex containing mismatch sites in the natural DNA triplet. (B) Target purine strand oriented parallel to TFO. (C) The mixture of each component one by one. (D) The mixture of each component using reverse-oriented Py-strand and Pu-strand.

Figure S6 Optimized structures of the dAN-derivatives/^{5m}CG base triplet.

(A) aminoethyl-dAN/5mCG, (B) 3-phenol-dAN/5mCG and (C) 2-phenol-dAN/5mCG, DFT at

B3LYP/6-31 level.

Figure S7 Molecular modeling of triplex DNA formation.

This triplex DNA includes aminoethyl-dAN/^{5m}CG or CG base triplet under Amber99 force field using HyperChem (7.52). At the end of the calculation, three triplet base pairs are shown as snapshots. The part of the triplet base pair of the artificial nucleoside analogue is shown by the CPK model, and the base triplet structures on both sides are shown in green. (A) Complex containing Aminoethyl-dAN and ^{5m}CG base pair. (B) Complex containing Aminoethyl-dAN and CG base pair.

(A) Aminoethyl-dAN/^{5m}CG (5'-AZG-3' sequence)

(B) Aminoethyl-dAN/CG (5'-AZG-3' sequence)

