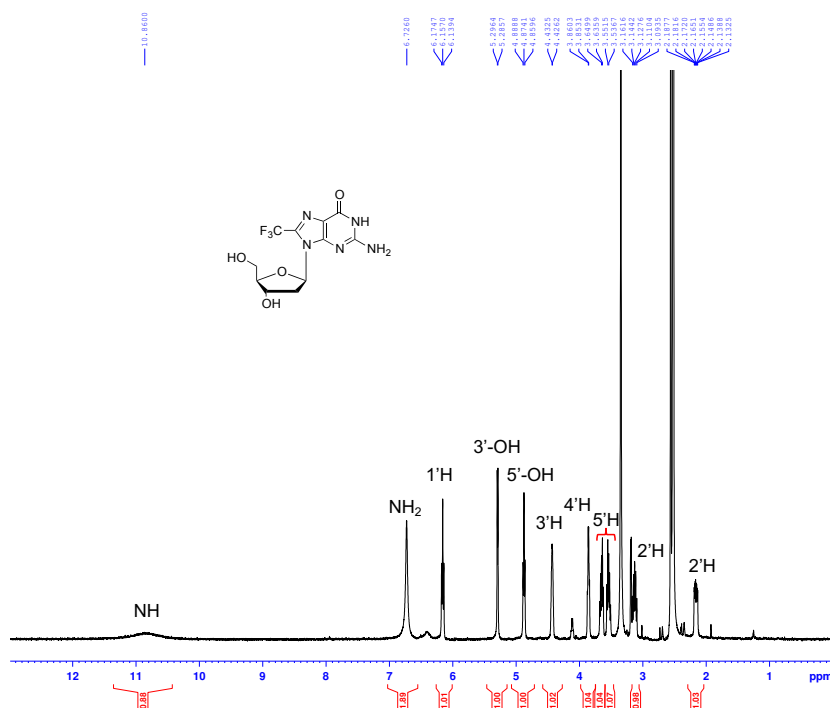


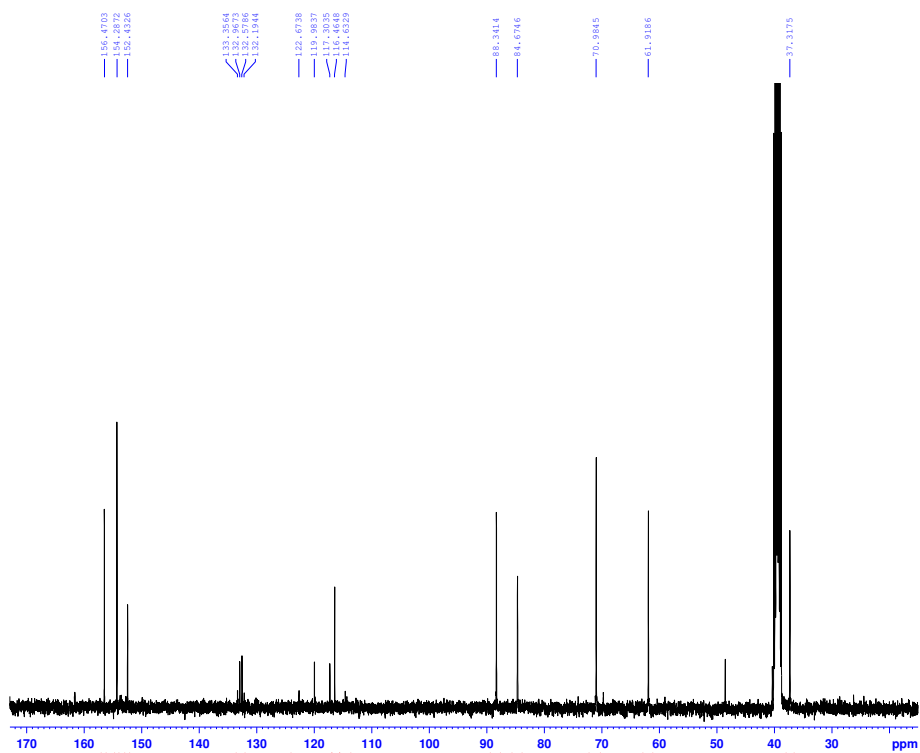
## Supporting Information

### General.

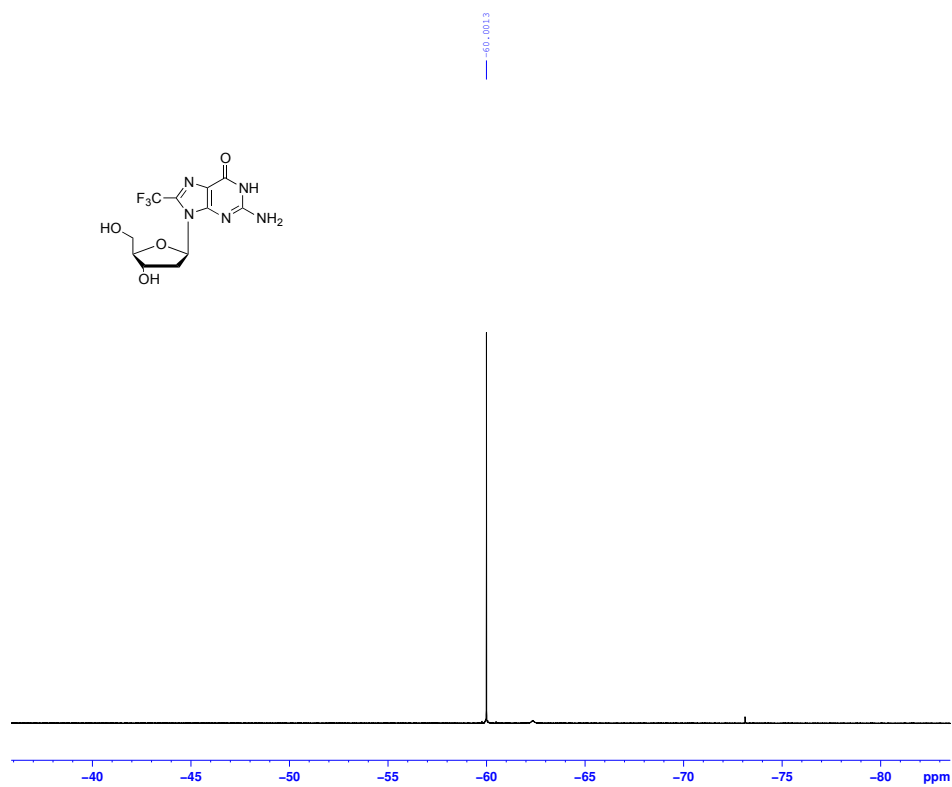
$^1\text{H}$ -NMR,  $^{13}\text{C}$ -NMR,  $^{19}\text{F}$ -NMR and  $^{31}\text{P}$ -NMR spectra were recorded on a BRUKER (AV-400M) magnetic resonance spectrometer. DMSO- $d_6$  and  $\text{CDCl}_3$  were used as the solvents. Coupling constants (J) values are given in Hz and are correct to within 0.5 Hz. Signal patterns are indicated as br, broad; s, singlet; d, doublet; t, triplet; m, multiplet. All reagents were purchased from Aldrich, TCI (Tokyo Chemical Industry Co., Ltd.) or Wako (Wako Pure Chemical Industries, Ltd.). Thin layer chromatography was performed using TLC Silica gel 60 F254 (Merck). Compounds were visualized by staining with a potassium permanganate solution. High-resolution mass spectra (HRMS) were recorded by electrospray ionization (ESI) on a Thermo Scientific Q Exactive instrument.



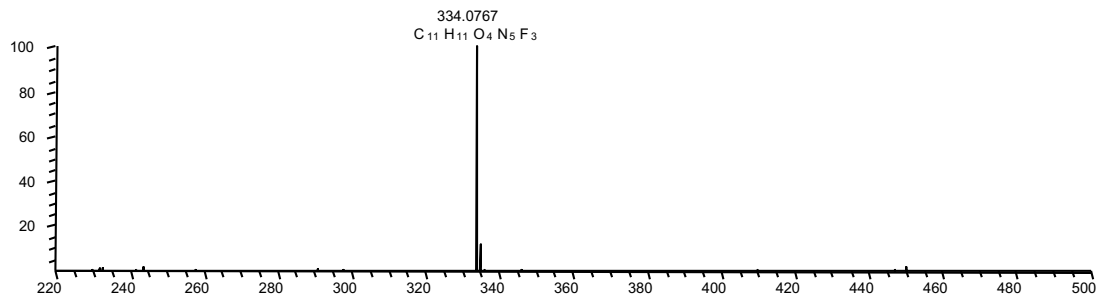
Supplementary Figure S1.  $^1\text{H}$  NMR spectrum of  $^{\text{F}}\text{G}$ .



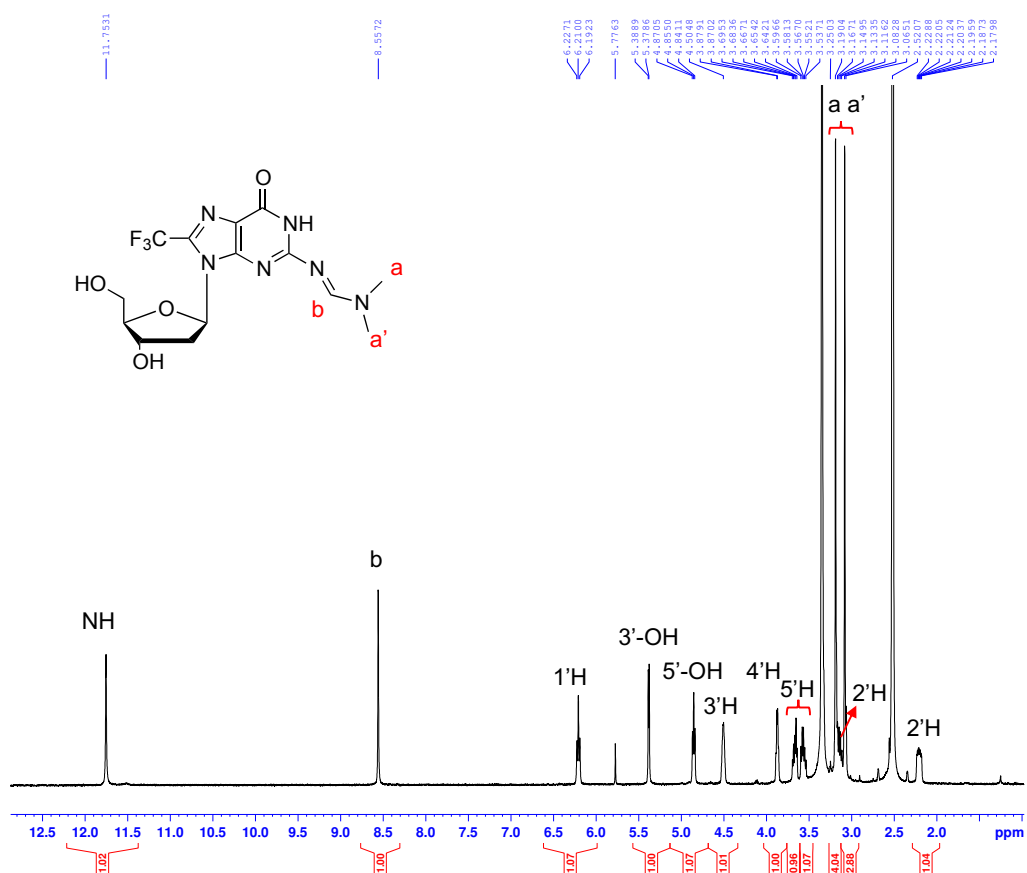
Supplementary Figure S2.  $^{13}\text{C}$  NMR spectrum of  $^{\text{F}}\text{G}$ .



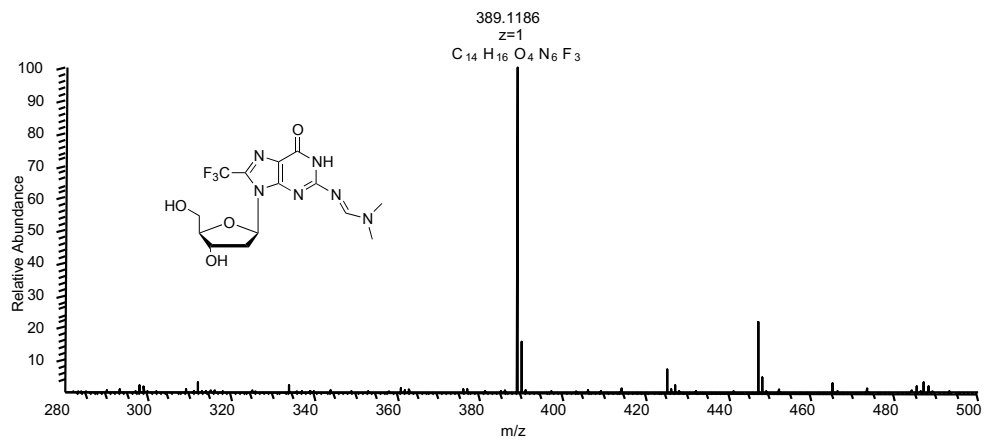
Supplementary Figure S3.  $^{19}\text{F}$  NMR spectrum of  $^{\text{F}}\text{G}$ .



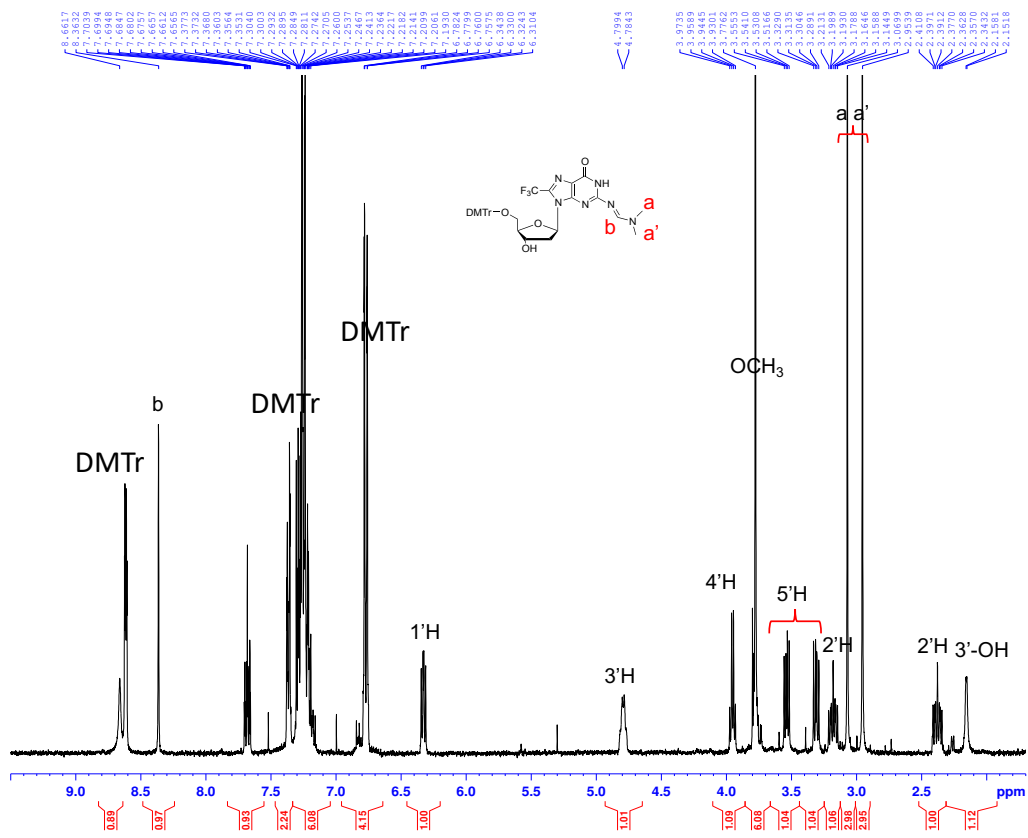
Supplementary Figure S4. HRMS spectrum of <sup>1</sup>G.



Supplementary Figure S5. <sup>1</sup>H NMR spectrum of compound 1

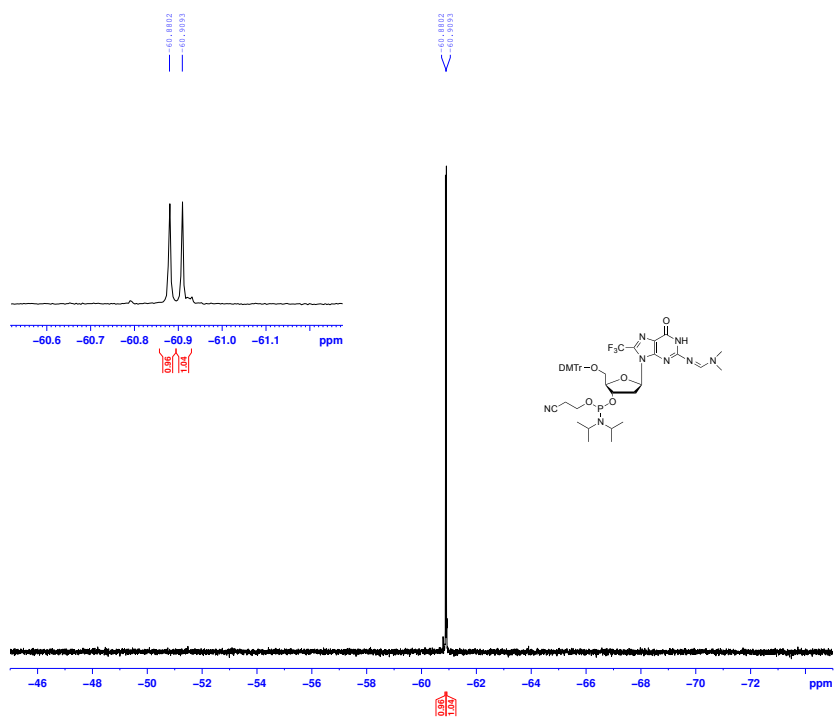


Supplementary Figure S6. HRMS spectrum of compound 1

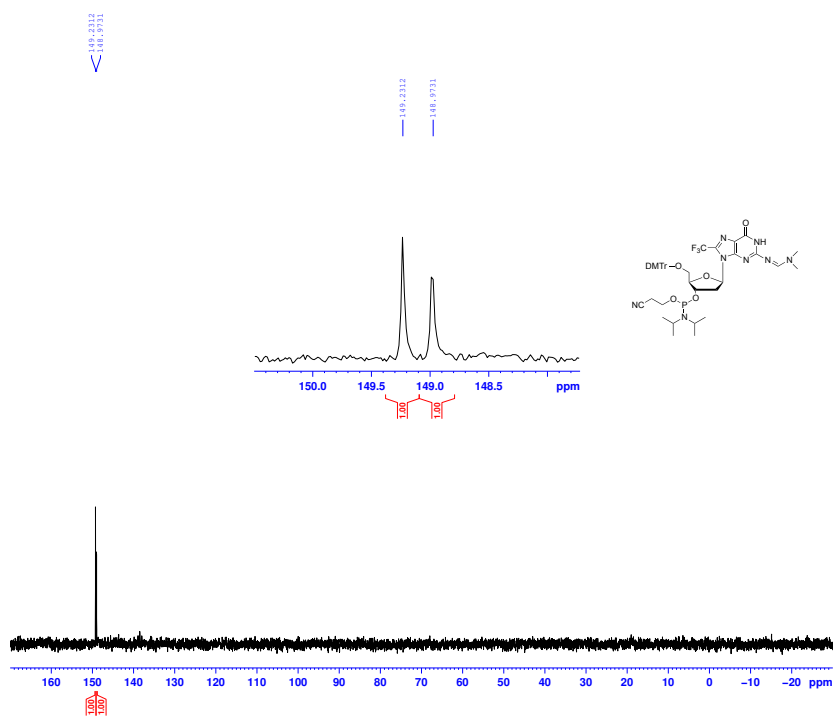


Supplementary Figure S7. <sup>1</sup>H NMR spectrum of compound 2

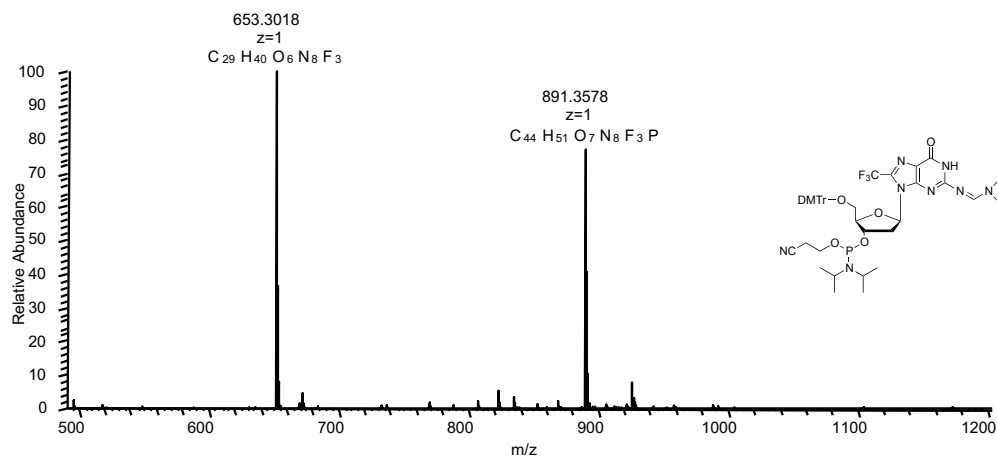




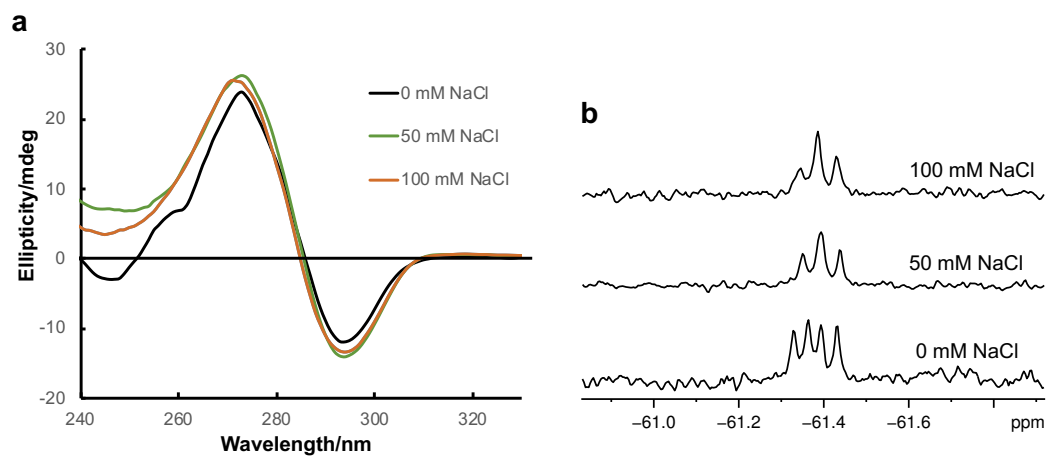
Supplementary Figure S10.  $^{19}\text{F}$  NMR spectrum of compound 3



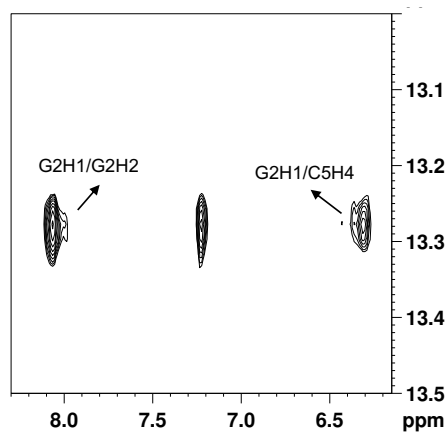
Supplementary Figure S11.  $^{31}\text{P}$  NMR spectrum of compound 3



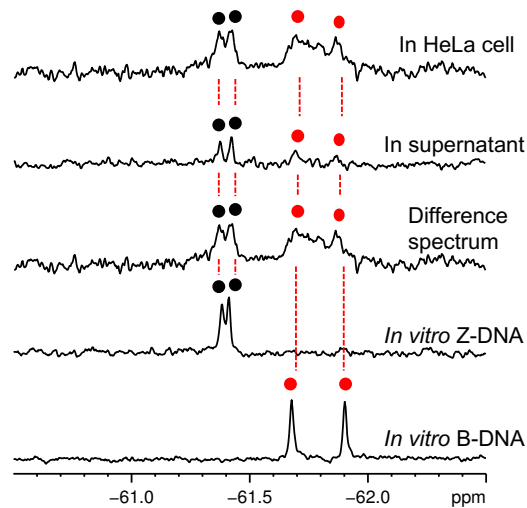
Supplementary Figure S12. HRMS spectrum of compound 3



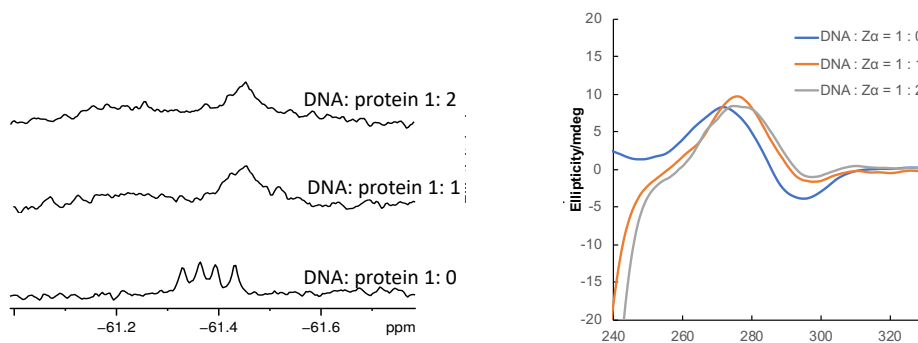
Supplementary Figure S13. a, CD spectra of  $d(C^F GCAC^F GCG)/d(C^F GCGT^F GCG)$ . Condition: 15  $\mu$ M duplex in 1 mM Na-PO<sub>4</sub> buffer (pH 7.0) and various concentration of NaCl. b, <sup>19</sup>F NMR spectra of  $d(C^F GCAC^F GCG)/d(C^F GCGT^F GCG)$ . Condition: 30  $\mu$ M duplex in 1 mM Na-PO<sub>4</sub> buffer (pH 7.0) and various concentration of NaCl.



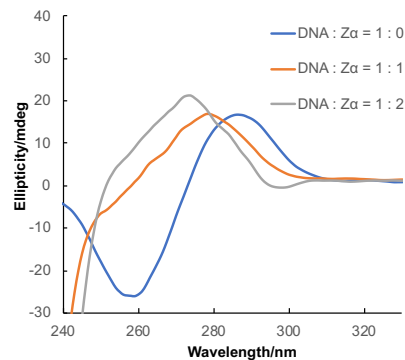
Supplementary Figure S14. The cross peaks of imino proton of G2 and amino proton of C5, indicated Watson–Crick-type base pairs of Z-DNA.



Supplementary Figure S15. In-cell  $^{19}\text{F}$  NMR spectrum of  $d(\text{C}^{\text{F}}\text{GCAC}^{\text{F}}\text{GCG})/d(\text{CGCGTGCG})$ . Comparison of  $^{19}\text{F}$  NMR spectra for *in vitro* B-DNA, *in vitro* Z-DNA, in HeLa cell, in supernatant and difference spectrum between HeLa cell and supernatant.

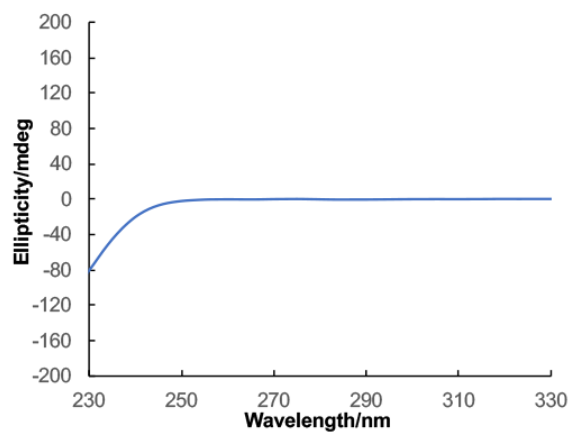


Supplementary Figure S16.  $^{19}\text{F}$  NMR (left) spectra of four positions modified 8-mer DNA  $d(\text{C}^{\text{F}}\text{GCAC}^{\text{F}}\text{GCG})/d(\text{C}^{\text{F}}\text{GCGT}^{\text{F}}\text{GCG})$  and  $Z\alpha$  protein. Condition:  $30\ \mu\text{M}$  DNA in  $1\ \text{mM}$   $\text{Na-PO}_4$  buffer (pH 7.0) and various  $\text{NaCl}$  concentrations. CD spectra (right) of four positions modified 8-mer DNA  $d(\text{C}^{\text{F}}\text{GCAC}^{\text{F}}\text{GCG})/d(\text{C}^{\text{F}}\text{GCGT}^{\text{F}}\text{GCG})$  and  $Z\alpha$  protein. Condition:  $5\ \mu\text{M}$  DNA in  $1\ \text{mM}$   $\text{Na-PO}_4$  buffer (pH 7.0) and various  $\text{NaCl}$  concentrations.



Supplementary Figure S17. CD spectra of native 8-mer DNA  $d(\text{CGACGCG})/d(\text{CGCGTGCG})$  and  $Z\alpha$  protein. Condition:  $10\ \mu\text{M}$  DNA in  $1\ \text{mM}$   $\text{Na-PO}_4$  buffer (pH 7.0) and various  $\text{NaCl}$  concentrations.





Supplementary Figure S18. CD spectrum Z $\alpha$  protein. Condition: 10  $\mu$ M Z $\alpha$  protein.in 1 mM Na-PO<sub>4</sub> buffer (pH 7.0).

Table S1. <sup>1</sup>H NMR chemical shifts (ppm) of r(CGCFGCG)<sub>2</sub> Z-DNA.

	H5	H8/6	H1'	H2'	H2''	H3'	H4'	H5'	H5''	H1	H2a/4a	H2a'/4a'
C1	5.51	7.24	5.64	1.47	2.38	4.47	3.71	2.94	3.53		6.69	6.69
G2		7.65	6.11	2.64	2.64	4.87	4.00	4.08	4.08	13.28	6.38	8.23
C3	5.03	7.24	5.64	1.59	2.51	4.70	3.71	2.41	3.71		6.30	8.07
G4			6.30	2.51	2.75	4.87	4.00	4.12	4.12	13.02	6.30	8.07
C5	5.11	7.31	5.64	1.59	2.51	4.87	3.71	2.75	4.07		6.38	8.23
G6		7.59	6.08	2.31	3.09	4.67	4.00	4.08	4.08	13.02	8.10	8.10