SUPPORTING INFORMATION

Impact of Bistrand Abasic Sites and Proximate Orientation on DNA Global Structure and Duplex Energetics

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Acronym	ΔH ^o _{shape} (kcal·mol ⁻¹)	TΔS ^o _{shape} (kcal·mol ⁻¹)	ΔG ^o _{shape} (kcal·mol ⁻¹)
CCC/GGG	113.7	91.1	22.5
FCC/GGG	69.9	56.7	13.2
CFC/GGG	86.7	72.2	14.5
CCF/GGG	70.8	57.6	13.2
CCC/GFG	79.0	65.1	13.9
FCC/GFG	85.8	71.2	14.5
CFC/GFG	81.9	68.2	13.6
CCF/GFG	77.0	64.1	12.9

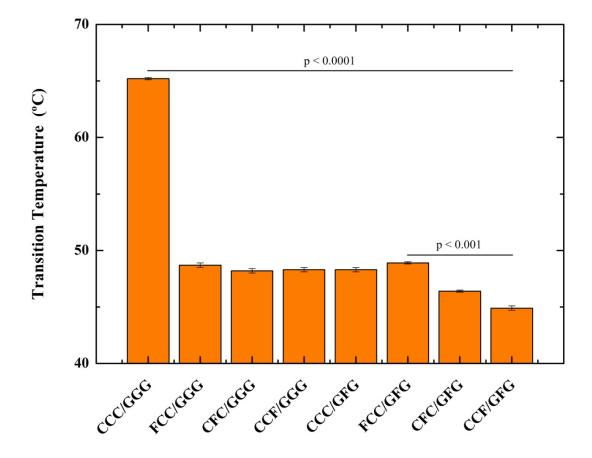
TABLE S1.Summary of van't Hoff Thermodynamic Parameters derived via Shape Analysis
of Temperature-dependent Profiles acquired for the Tridecameric Duplexes.

Thermodynamic data correspond to a DNA duplex of 45.4 μ M total strand concentration (C_T). Standard errors for ΔG° , ΔH° , and T ΔS° are within 2.0, 4.5, and 5.0 kcal·mol⁻¹, respectively.

Acronym	ΔH° _{25 °C} (kcal·mol ⁻¹)	T∆S° _{25 °C} (kcal·mol ⁻¹)	$\Delta G^{\circ}_{25^{\circ}C} (\text{kcal·mol}^{-1})$
CCC/GGG	79.32	63.3	16.0
FCC/GGG	80.56	69.2	11.4
CFC/GGG	74.16	63.4	10.8
CCF/GGG	76.07	65.1	10.9
CCC/GFG	72.51	61.8	10.7
FCC/GFG	60.65	50.7	10.0
CFC/GFG	74.20	63.9	10.3
CCF/GFG	58.55	49.6	8.9

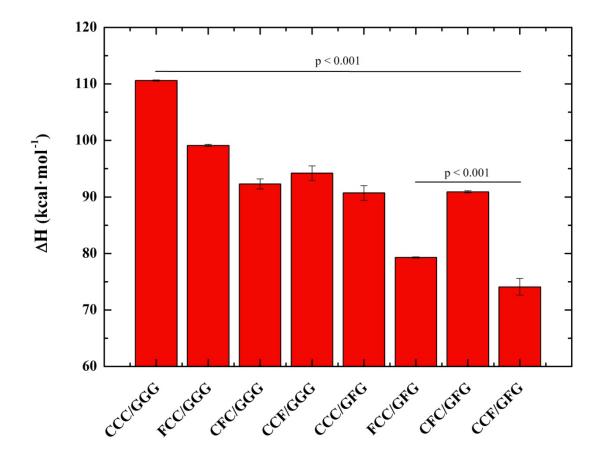
 TABLE S2.
 Heat-Capacity Corrected Thermodynamic Dissociation Parameters.

Values extrapolated to 25 °C using a ΔC_p of 780 cal·mol⁻¹·deg ⁻¹ (i.e., 60 cal·mol⁻¹·deg ⁻¹·bp⁻¹) applying the following relations: $\Delta H(T) = \Delta H(T_m) - \Delta C_p (T_m - T)$; $\Delta S(T) = \Delta H/T_m + \Delta C_p \ln(T/T_m) + R \ln (C_T/4)$; $\Delta G(T) = \Delta H(T_m) (1 - T/T_m) + \Delta C_p [T - T_m - T \ln (T/T_m)] - RT \ln (C_T/4)$, whereby T = 25 °C.

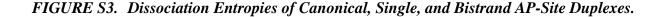


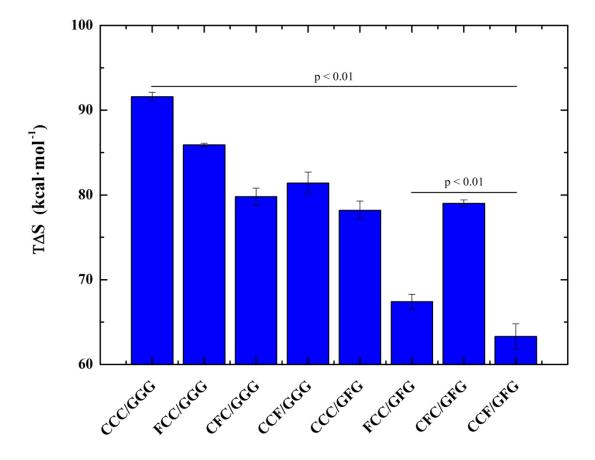
The single and bistrand abasic site duplexes exhibit statistically significant differences in thermal stability (p < 0.0001) relative to the undamaged parent tridecamer. The differential thermal stability of CFC/GFG and CCF/GFG are statistically significant (p < 0.001) when compared to the corresponding single AP-site reference duplexes CFC/GGG and CCF/GGG The bistrand CFC/GFG and CCF/GFG duplexes are thermally destabilized with CCF/GFG > CFC/GFG (p < 0.001). Thermal destabilization of FCC/GFG does *not* differ significantly (p > 0.5) relative to the single AP-site CCC/GFG and FCC/GFG duplexes.

FIGURE S2. Dissociation Enthalpies of Canonical, Single, and Bistrand AP-Site Duplexes.



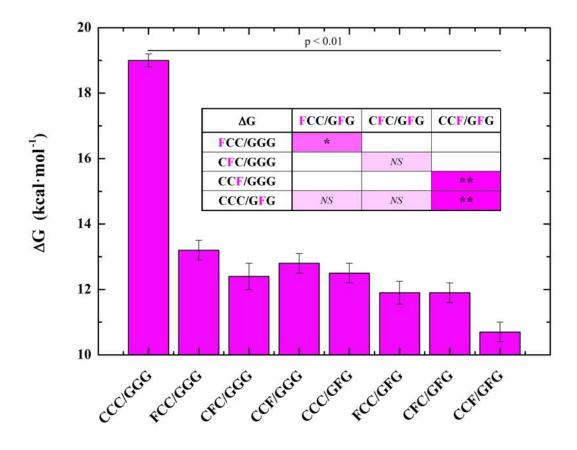
The single and bistrand abasic site duplexes exhibit statistically significant differences in enthalpic stability (p < 0.001) relative to the undamaged parent tridecamer. The differential enthalpic stability of FCC/GFG and CCF/GFG are statistically significant (p < 0.001) when compared to the corresponding single AP-site reference duplexes FCC/GGG and CCF/GGG. Enthalpic destabilization of the bistrand FCC/GFG and CCF/GFG duplexes are comparable (p > 0.5). The dissociation enthalpy of CFC/GFG does *not* differ significantly (p > 0.5) relative to the single AP-site CCC/GFG and CFC/GGG duplexes.





The single and bistrand abasic site duplexes exhibit statistically significant differences in entropic stability (p < 0.01) relative to the undamaged parent tridecamer. The differential entropic stability of FCC/GFG and CCF/GFG are statistically significant (p < 0.001) when compared to the corresponding single AP-site reference duplexes FCC/GGG and CCF/GGG. Entropic stabilization of the bistrand FCC/GFG and CCF/GFG duplexes are comparable (p > 0.5). The dissociation entropy of CFC/GFG is *not* significantly different (p > 0.5) relative to the single AP-site CCC/GFG and CFC/GGG duplexes.





The single and bistrand abasic duplexes exhibit statistically significant differences in thermodynamic stability (p < 0.01) relative to the undamaged parent tridecamer. INSET: Statistical significance of dissociation free energy (ΔG) differences employing ANOVA and Tukey tests. Meaningful comparisons are designated as ** and * signifying that the corresponding means differ at a level of p < 0.05 and p < 0.1, respectively. An *NS* designation indicates that the values differ somewhat albeit not significantly.