

Supplementary Information

DNA binding by pixantrone

Najia Adnan^a, Damian P. Buck^a, Benny J. Evison^b, Suzanne M. Cutts^{b*},
Don R. Phillips^b and J. Grant Collins^{a*}

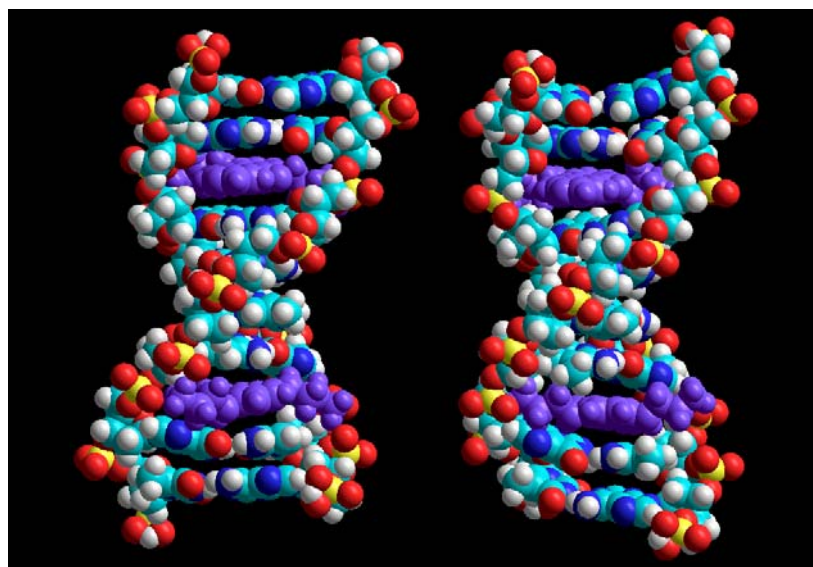
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| Supplementary Information Table 1 | ¹ H NMR assignments of the resonances of free d(ACGATCGT) ₂ and with added pixantrone at R = 0.6 |
| Supplementary Information Table 2 | ¹ H NMR assignments of the resonances of free d(A ^{5Me} CGAT ^{5Me} CGT) ₂ and with added pixantrone at R = 2.0 |
| Supplementary Information Figure 1 | Models of two pixantrone molecules intercalated into the symmetric 5'-CG or 5'- ^{5Me} CG sites in d(ACGATCGT) ₂ and d(A ^{5Me} CGAT ^{5Me} CGT) ₂ . |

Supplementary Information Table 1 ^1H NMR assignment of the DNA resonances of $d(\text{ACGATCGT})_2$ with added pixantrone, at $R = 0.6$ molar ratio in pH 7 phosphate buffer at 25 °C. Changes in chemical shift upon pixantrone binding are given in brackets, with a negative sign indicating an upfield shift.

Residue	H8/H6	H2/H5	H1'	H2'	H2''	Methyl
A₁	8.17 (-0.05)	7.87 (-0.16)	6.13 (-0.10)	2.58 (-0.05)	2.76 (-0.01)	
C₂	7.38 (-0.02)	5.35 (-0.10)	5.35 (-0.02)	2.11 (0.01)	2.32 (-0.03)	
G₃	7.88 (-0.02)		5.62 (-0.07)	2.68 (-0.03)	2.79 (-0.03)	
A₄	8.18 (-0.04)	7.77 (-0.05)	6.23 (-0.06)	2.60 (-0.05)	2.92 (-0.06)	
T₅	7.12 (-0.04)		5.89 (-0.04)	1.99 (-0.04)	2.42 (-0.01)	1.30 (-0.06)
C₆	7.40 (-0.02)	5.54 (-0.06)	5.72 (-0.01)	1.94 (-0.01)	2.36 (0.00)	
G₇	7.94 (-0.03)		5.98 (-0.09)	2.62 (-0.06)	2.72 (0.03)	
T₈	7.34 (-0.07)		6.18 (-0.05)	2.26 (-0.01)	2.26 (-0.01)	1.54 (-0.10)

Supplementary Information Table 2 ^1H NMR assignment of the DNA resonances of $d(\text{A}^{5\text{Me}}\text{CGAT}^{5\text{Me}}\text{CGT})_2$ with added pixantrone at $R = 2$ in pH 7 phosphate buffer at 25 °C. Changes in chemical shift upon pixantrone binding are given in brackets, with a negative sign indicating an upfield shift.

	H8/H6	H2	H1'	H2'	H2''	Methyl
A₁	8.18 (-0.02)	7.99 (-0.04)	6.11 (-0.10)	2.48 (-0.08)	2.76 (-0.05)	
C₂Me	7.21 (-0.02)		5.80 (0.43)	2.21 (-0.10)	2.30 (-0.04)	1.43 (-0.15)
G₃	7.92 (0.03)		5.45 (-0.18)	2.67 (-0.06)	2.72 (-0.00)	
A₄	8.21 (-0.03)	7.78 (-0.07)	6.20 (-0.10)	2.64 (-0.03)	2.84 (-0.11)	
T₅	7.07 (-0.11)		5.88 (-0.07)	1.67 (-0.26)	2.39 (-0.13)	1.23 (-0.14)
C₆Me	7.29 (0.02)		5.83 (0.20)	2.08 (-0.01)	2.36 (0.04)	1.59 (-0.14)
G₇	7.96 (-0.02)		5.96 (-0.11)	2.67 (-0.04)	2.76 (0.02)	
T₈	7.30 (-0.10)		6.18 (-0.07)	2.23 (-0.02)	2.23 (-0.02)	1.43 (-0.18)



Supplementary Information Figure 1 Energy minimised HyperChem models of two pixantrone molecules intercalated into the symmetric 5'-CG or 5'-^{5Me}CG sites in d(ACGATCGT)₂ (left-hand side) and d(A^{5Me}CGAT^{5Me}CGT)₂ (right-hand side). The pixantrone intercalative-binding is shown from the minor groove (top) and the major groove (bottom).