

Supporting Informations

Electronic Absorption and EPR Spectra.

The complex $[\text{Cu}(5,6\text{-dmp})_3]^{2+}$ **2** in 10% DMF/5 mM Tris-HCl/50 mM NaCl aqueous buffer solution shows a broad ligand field band around 680 nm (Table S1), which is in agreement with the square based geometry observed in its X-ray crystal structure. The phen (**1**) and dpq (**3**) analogues also show a single LF spectral feature around 712 and 728 nm (Table S1). The higher LF energy of **2** is expected of its stronger Cu-N bond (cf. above). The ligand-based bands are observed around 267 (**1**), 280 (**2**) and 296 nm (**3**). The analogues Zn(II) complexes also exhibit ligand-based bands around values closer to their Cu(II) analogues. The EPR spectra of **1** – **3** in frozen $\text{CH}_3\text{CN}:\text{acetone}$ solutions are axial $[g_{\parallel} > g_{\perp} > 2.023, G = (g_{\parallel}-2)/(g_{\perp}-2) \approx 3.8]$ ¹, which is in agreement with the absorption spectral data and the X-ray crystal structures of **1** and **2**. The g_{\parallel} and A_{\parallel} values expected for a CuN_4 chromophore¹ are respectively 2.200 and $200 \times 10^{-4} \text{ cm}^{-1}$ and the g_{\parallel} value is expected to increase and the A_{\parallel} value to decrease on strong axial interaction by nitrogen. Thus the EPR spectral parameters of the present complexes (g_{\parallel} 2.264 – 2.278; A_{\parallel} 161 – $174 \times 10^{-4} \text{ cm}^{-1}$) are consistent with the tetragonal distortion observed in the X-ray structure and the value of the structural index $g_{\parallel}/A_{\parallel}$ quotient (**1**, 140; **2**, 130; **3**, 141 cm) is consistent with the planarity of CuN_4 basal plane of the complexes.

Reference

1. M. Palaniandavar, T. Pandian, M. Lakshminarayanan and H. Manohar, *J. C. S. Dalton Trans.*, 1995, 455.

Table S1. Electronic absorption and EPR spectral properties ^a of Cu(II) and Zn(II) complexes

Complex		λ_{max} in nm (ϵ , $\text{M}^{-1} \text{cm}^{-1}$)		EPR spectra ^d	
		Ligand field ^b	Ligand based ^c		
[Cu(phen) ₃] ²⁺	1	712 (108)	267 (90150) 294 sh	g_{\parallel} A_{\parallel}	2.278 164
				g_{\perp}	2.073
[Cu(5,6-dmp) ₃] ²⁺	2	680 (58)	280 (679840) 351 sh	g_{\parallel} A_{\parallel}	2.264 174
				g_{\perp}	2.084
[Cu(dpq) ₃] ²⁺	3	728 (99)	254 (60858) 296 sh	g_{\parallel} A_{\parallel}	2.270 161
				g_{\perp}	2.084
[Zn(phen) ₃] ²⁺	4	—	267 (30500) 230 (32300)	—	—
[Zn(5,6-dmp) ₃] ²⁺	5	—	278 (91260) 298 (95460)	—	—
[Zn(dpq) ₃] ²⁺	6	—	255(55465)	—	—

^aIn DMF - buffer solution. ^bConcentration, $5 \times 10^{-3} \text{ mol.dm}^{-3}$. ^c Concentration, $2 \times 10^{-5} \text{ mol.dm}^{-3}$. ^d Acetonitrile/acetone (4:1 V/V) glass at 77 K, A_{\parallel} in 10^{-4} cm^{-1} .

Table S2. Electrochemical behaviour of ^a the copper(II) complexes on interaction with CT DNA
 at R = [DNA]/[Cu] = 3

Complexes	R	E_{pa} (V)	E_{pc} (V)	$E_{1/2}$ (V)		ΔE_p (mV)	K_{2+}/K_+
				CV	DPV ^b		
[Cu(phen) ₃] ²⁺ 1	0	0.040	-0.084	0.022	0.021	124	1.17
	3	0.038	-0.066	0.014	0.017	104	
[Cu(5,6-dmp) ₃] ²⁺ 2	0	-0.004	-0.098	-0.051	-0.031	94	3.49
	3	-0.014	-0.102	-0.063	-0.063	88	
[Cu(dpq) ₃] ²⁺ 3	0	0.118	0.006	0.062	0.065	112	1.78
	3	0.110	0.036	0.073	0.072	146	

^a Measured vs. sat. calomel electrode, using glassy carbon electrode scan rate, 50 mV⁻¹; supporting electrolyte 10% DMF : 5 mM Tris-HCl/50 mM NaCl; Complex concentration: 5 x 10⁻⁴ mol.dm³ ;

^b Differential pulse voltammetry (DPV), scan rate 1 mV s⁻¹, pulse height 50 mV.

Table S3. Concentration dependent oxidative cleavage data of SC pUC19 DNA (40 μM , in base pair) by complexes **2** in the presence of ascorbic acid.

Serial no	Reaction conditions	Form (%)	
		SC	NC
1	DNA control	91.8	8.2
2	DNA + H_2A	90.8	9.2
3	DNA + 2 (10 μM)	86.8	13.2
4	DNA + 2 (20 μM)	83.4	16.6
5	DNA + 2 (30 μM)	52.8	47.2
6	DNA + 2 (40 μM)	45.8	54.2
7	DNA + 2 (50 μM)	43.0	57.0

Table S4. Concentration dependent oxidative cleavage data of SC pUC19 DNA (40 μM , in base pair) by complexes **3** (30 μM) in the presence of ascorbic acid.

Serial no	Reaction conditions	Form(%)	
		SC	NC
1	DNA control	91.8	8.2
2	DNA + H_2A	90.8	9.2
3	DNA + 3 (10 μM)	2.3	97.7
4	DNA + 3 (20 μM)	1.3	98.7
5	DNA + 3 (30 μM)	0.3	99.7
6	DNA + 3 (40 μM)	-	-
7	DNA + 3 (50 μM)	-	-

Figure S1. The effect of CT DNA on the absorption spectra of complexes of **1 - 6** ($R = [\text{DNA}]/[\text{complex}] = 25$).

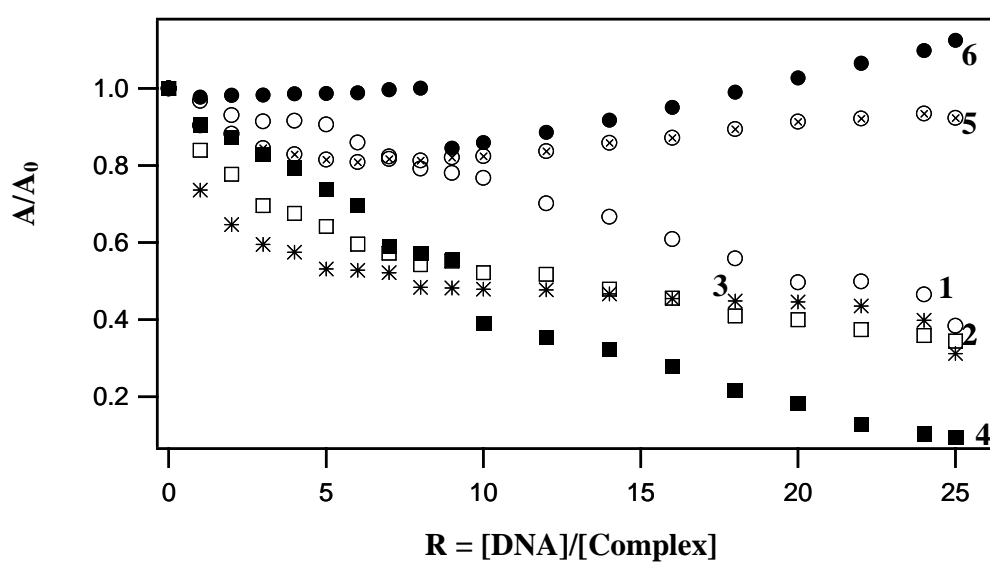


Figure S2. Effect of addition of complex **2** and **5** on the CD intensity of the CT DNA at different complex concentrations in a 5mM Tris-HCl/ 50 mM NaCl buffer at pH = 7.1 at 25 °C

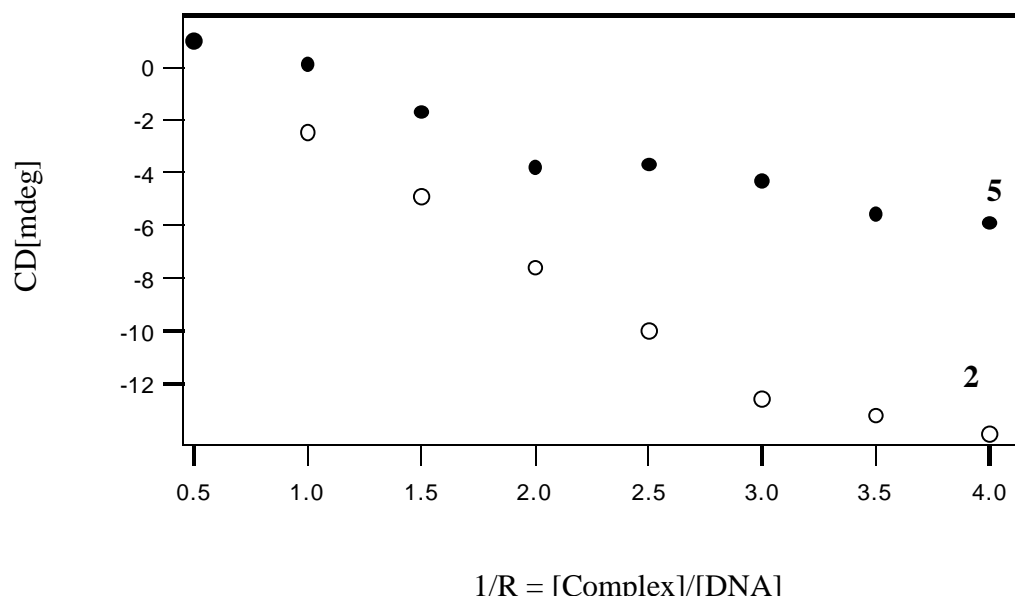


Figure S3A Circular Dichroism spectra of CT DNA in the absence (a) and presence of (b) $[\text{Cu}(\text{phen})_3]^{2+}$ ($1/R = 2$); Conc. of CT DNA = $2 \times 10^{-5} \text{ mol.dm}^3$

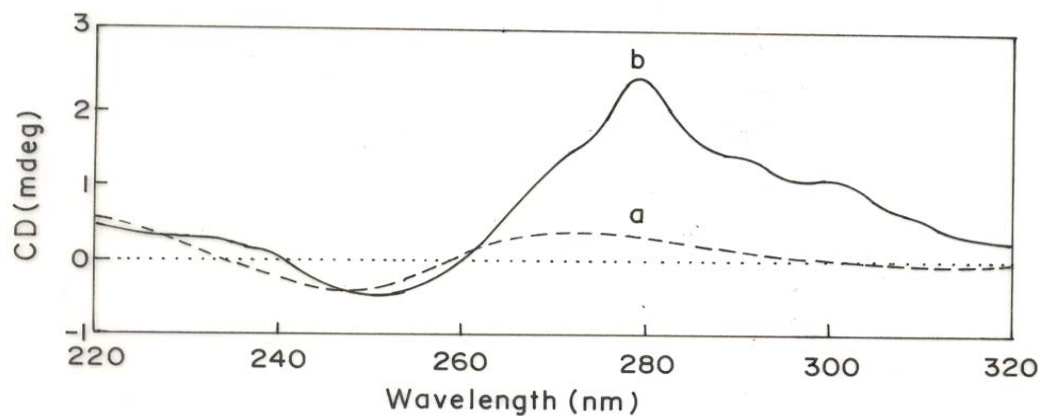


Figure S3B Circular Dichroism spectra of CT DNA in the absence (a) and presence of (b) $[\text{Zn}(\text{phen})_3]^{2+}$ ($1/R = 2$); Conc. of CT DNA = $2 \times 10^{-5} \text{ mol.dm}^3$

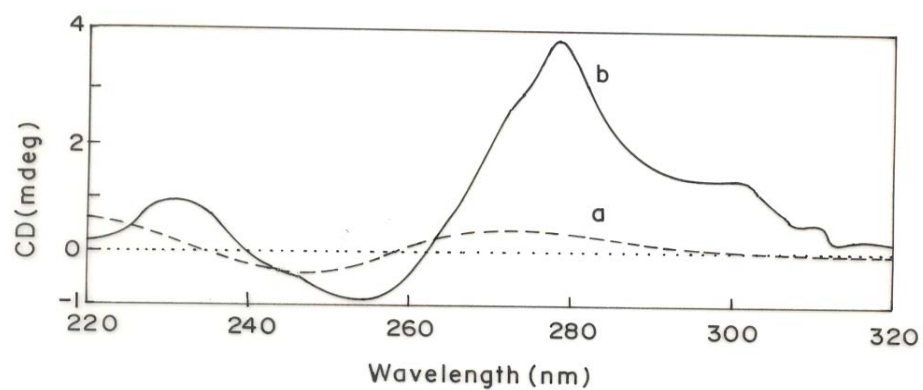


Figure S4. Circular Dichroism spectra of CT DNA, d(AT)₁₂, d(GC)₁₂ and d(GTCGAC)₂ in the absence (a) and presence of the *rac*-[Zn(5,6-dmp)₃]²⁺ (b) Conc of the oligonucleotides = 2×10^{-5} mol.dm³ ; Cell length = 0.2 cm

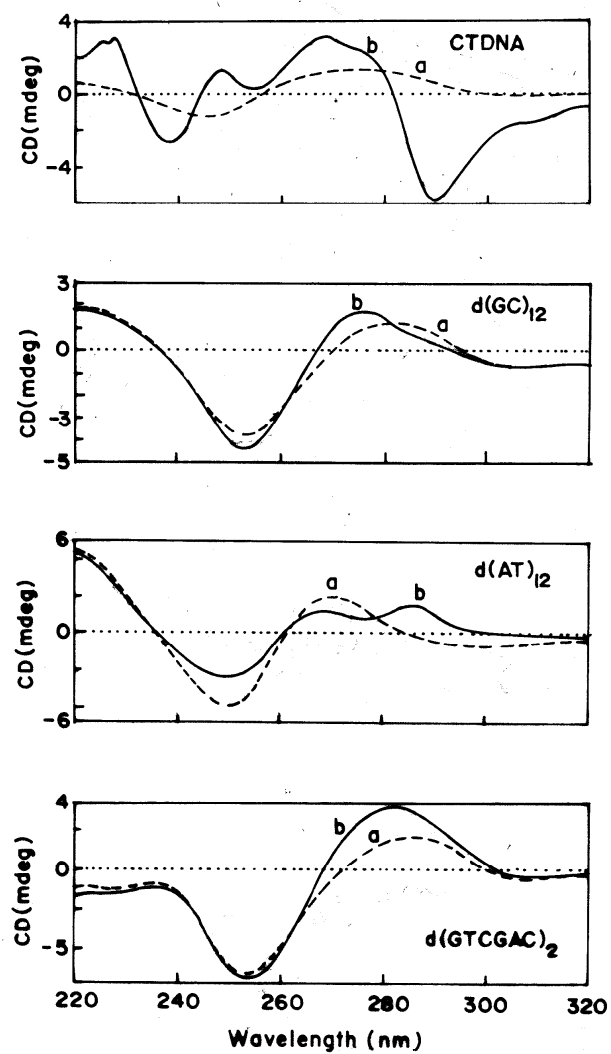


Figure S5A. Circular Dichroism spectra of CT DNA in the absence (a) and presence of *rac*-[Cu(5,6-dmp)₃]²⁺ at 1/R = 2 (b) and of CT DNA incubated with EthBr in the presence of *rac*-[Cu(5,6-dmp)₃]²⁺ at 1/R = 2 (c). Conc. of CT DNA = 2 x 10⁻⁵ mol.dm³.

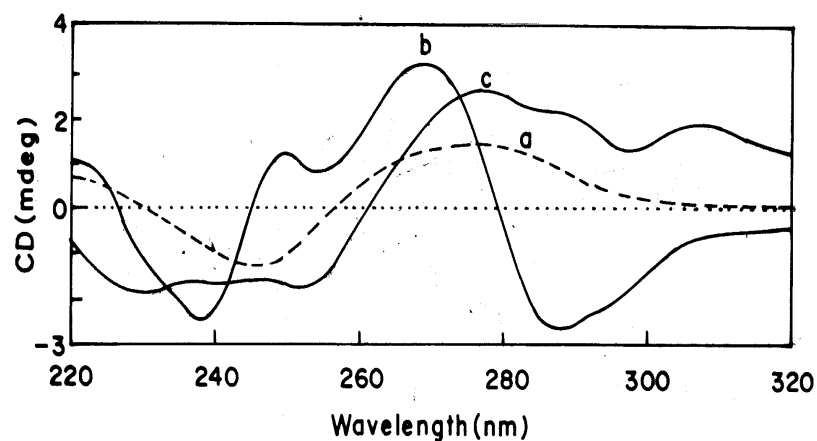


Figure S5B. Circular Dichroism spectra of CT DNA in the absence (a) and presence of *rac*-[Zn(5,6-dmp)₃]²⁺ at 1/R = 2 (b) and of CT DNA incubated with EthBr in the presence of *rac*-[Zn(5,6-dmp)₃]²⁺ at 1/R = 2 (c). Conc. of CT DNA = 2 x 10⁻⁵ mol.dm³.

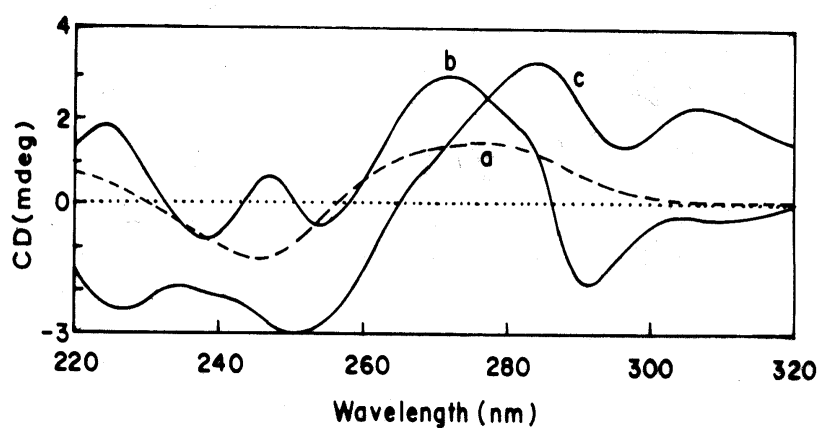


Figure S6A. Circular Dichroism spectra of CT DNA in the absence (a) and presence of *rac*-[Cu(5,6-dmp)₃]²⁺ at 1/R = 2 (b) and of CT DNA incubated with *rac*-[Cu(phen)₃]²⁺ in the presence of *rac*-[Cu(5,6-dmp)₃]²⁺ at 1/R = 2 (c). Conc. of CT DNA = 2 x 10⁻⁵ mol.dm³.

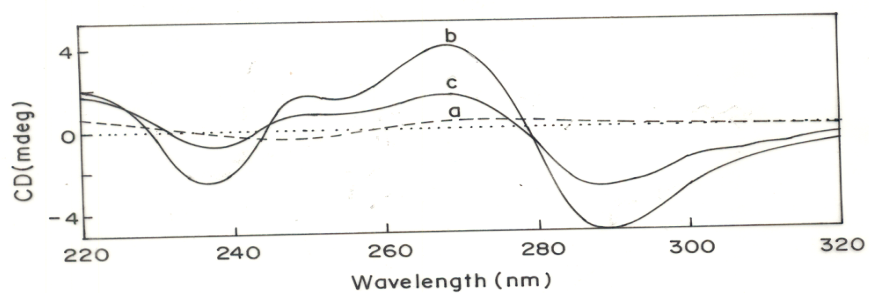


Figure S6B. Circular Dichroism spectra of CT DNA in the absence (a) and presence of *rac*-[Cu(phen)₃]²⁺ at 1/R = 2 (b) and of CT DNA incubated with *rac*-[Cu(5,6-dmp)₃]²⁺ in the presence of *rac*-[Cu(phen)₃]²⁺ at 1/R = 2 (c). Conc. of CT DNA = 2 x 10⁻⁵ mol.dm³.

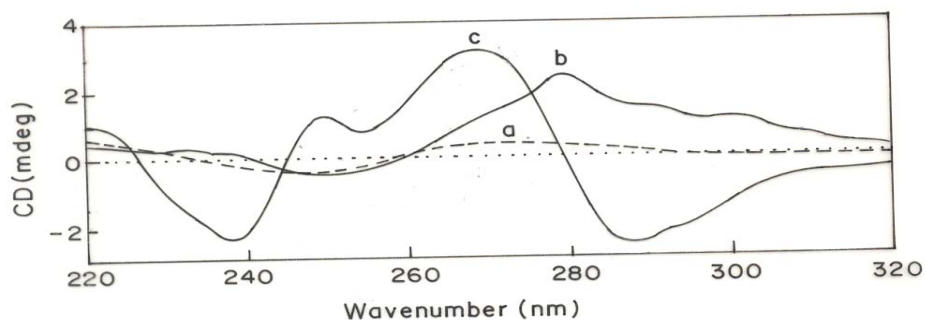


Figure S7A. Circular Dichroism spectra of CT DNA in the absence (a) and presence of *rac*-[Zn(5,6-dmp)₃]²⁺ at 1/R = 2 (b) and of CT DNA incubated with *rac*-[Zn(phen)₃]²⁺ in the presence of *rac*-[Zn(5,6-dmp)₃]²⁺ at 1/R = 2 (c). Conc. of CT DNA = 2 x 10⁻⁵ mol.dm³.

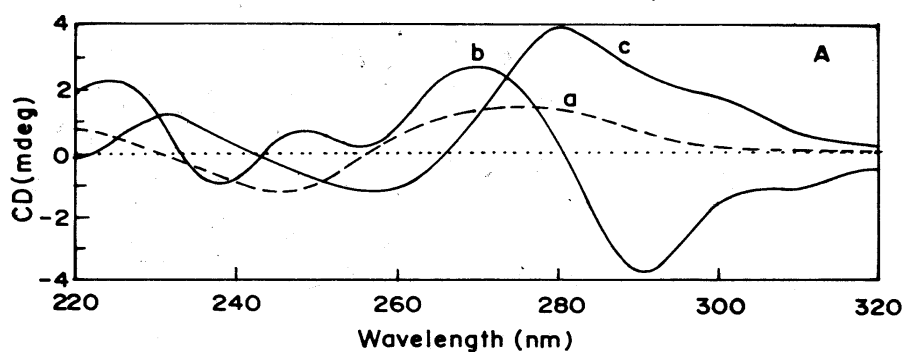


Figure S7B. Circular Dichroism spectra of CT DNA in the absence (a) and presence of *rac*-[Zn(phen)₃]²⁺ at 1/R = 2 (b) and of CT DNA incubated with *rac*-[Zn(5,6-dmp)₃]²⁺ in the presence of *rac*-[Zn(phen)₃]²⁺ at 1/R = 2 (c). Conc. of CT DNA = 2 x 10⁻⁵ mol.dm³.

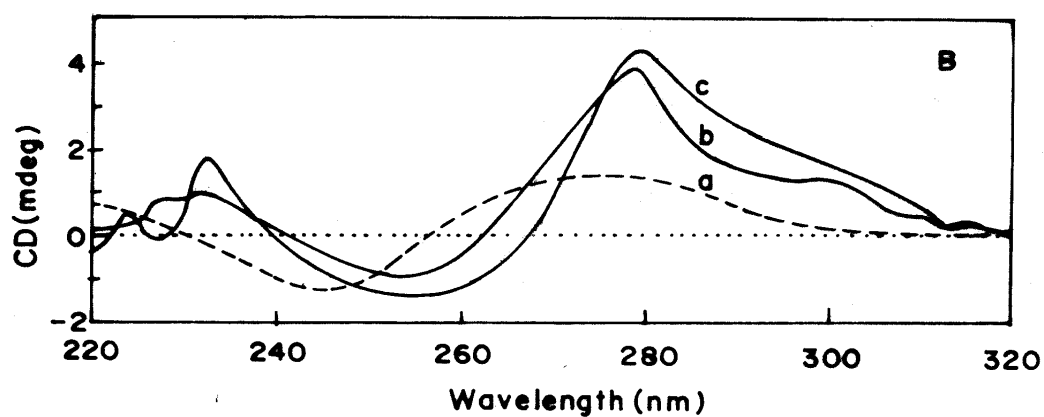


Figure S8A. Circular Dichroism spectra of CT DNA in the absence (a) and presence of *rac*-[Cu(5,6-dmp)₃]²⁺ at 1/R = 2 (b) and of CT DNA incubated with EDTA in the presence of *rac*-[Cu(5,6-dmp)₃]²⁺ at 1/R = 2 (c). Conc. of CT DNA = 2 x 10⁻⁵ mol.dm³.

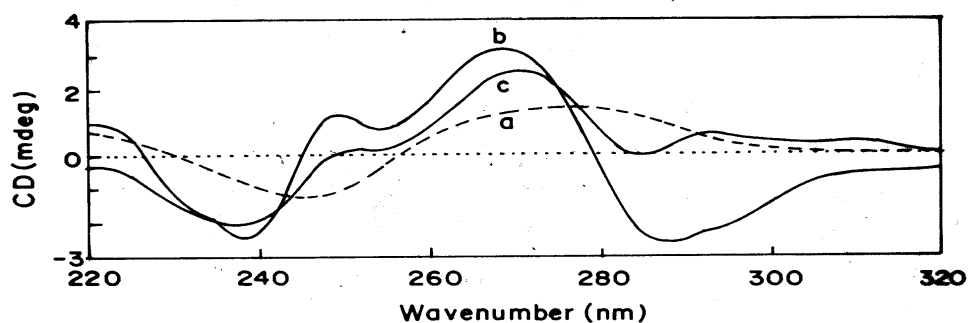


Figure S8B. Circular Dichroism spectra of CT DNA in the absence (a) and presence of *rac*-[Zn(5,6-dmp)₃]²⁺ at 1/R = 2 (b) and of CT DNA incubated with EDTA in the presence of *rac*-[Zn(5,6-dmp)₃]²⁺ at 1/R = 2 (c). Conc. of CT DNA = 2 x 10⁻⁵ mol.dm³.

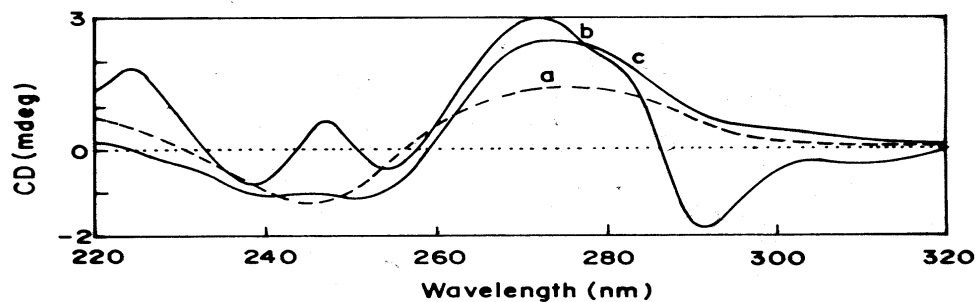


Figure S9. Cleavage of supercoiled pUC 19 DNA (40 μM) by the copper(II) complexes in 5% DMF : 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^{\circ}\text{C}$ in the presence of ascorbic acid (H_2A , 10 μM) at 37 $^{\circ}\text{C}$. Lane 1, DNA ; lane 2 + DNA + H_2A ; lane 3, DNA + $[\text{Cu}(\text{phen})_2\text{Cl}]\text{Cl}$ + H_2A ; lane 4, DNA + **1** + H_2A ; lane 5, DNA + $[\text{Cu}(5,6\text{-dmp})_2\text{Cl}]\text{Cl}$ + H_2A ; lane 6, DNA + **2** + H_2A ; Complex concentration is 30 μM for lanes 2-7. Forms I and II are supercoiled and nicked circular forms of DNA respectively.

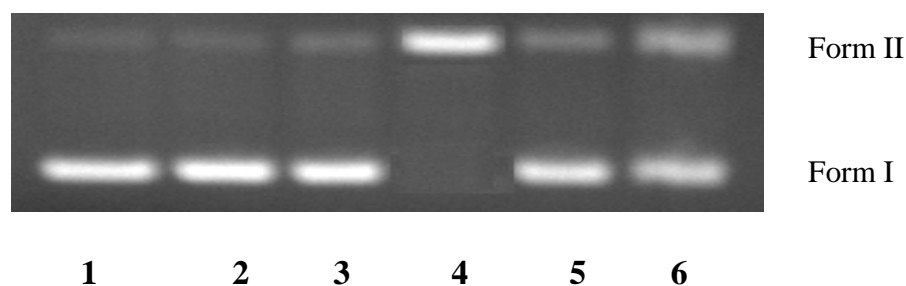


Figure S10A. Cleavage of supercoiled pUC19 DNA (40 μM) by complex **1** (10 – 30 μM) in 10% DMF / 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^{\circ}\text{C}$ in the presence of ascorbic acid (H_2A , 10 μM) at 37 $^{\circ}\text{C}$. Lane 1, + DNA + H_2A ; lane 2, DNA + **1** (10 μM) + H_2A ; lane 3, DNA + **1** (20 μM) + H_2A ; lane 4, DNA + **1** (30 μM) + H_2A ; lane 5, DNA + **1** (30 μM) + H_2A + 2 μL DMSO; Lane 6, DNA + **1** (30 μM) + H_2A + Distamycin. Forms I and II are supercoiled and nicked circular forms of DNA respectively.

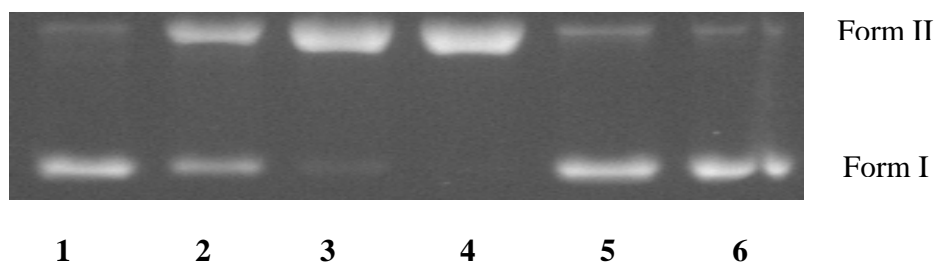


Figure S10B. Concentration dependent DNA (40 μM) cleavage by complex **1** (10 – 50 μM) in 10% DMF / 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^{\circ}\text{C}$ in the presence of ascorbic acid (H_2A , 10 μM) at 37 $^{\circ}\text{C}$. Lane 1, DNA ; lane 2, DNA + H_2A ; lane 3, DNA + **1** (10 μM) + H_2A ; lane 4, DNA + **1** (20 μM) + H_2A ; lane 5, DNA + **1** (30 μM) + H_2A ; lane 6, DNA + **1** (40 μM) + H_2A ; lane 7, DNA + **1** (50 μM) + H_2A ; Forms I and II are supercoiled and nicked circular forms of DNA respectively

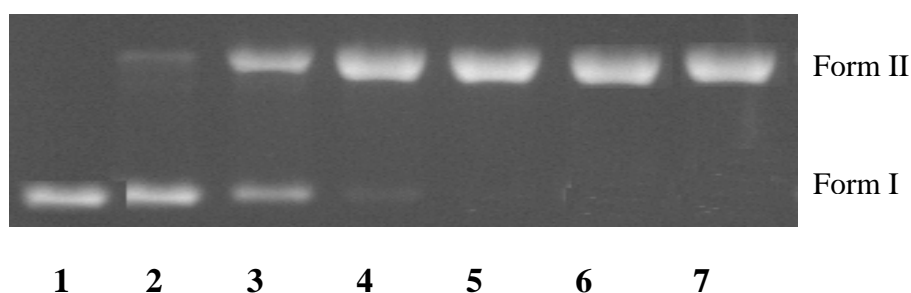


Figure S10C. Concentration dependent DNA (40 μM) cleavage by complex **1** (2 – 10 μM) in 10% DMF / 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^{\circ}\text{C}$ in the presence of ascorbic acid (H_2A , 10 μM) at 37 $^{\circ}\text{C}$. lane 1 + DNA + H_2A ; lane 2, DNA + **1** (2 μM) + H_2A ; lane3, DNA + **1** (4 μM) + H_2A ; lane 4, DNA + **1** (6 μM) + H_2A ; lane 5, DNA + **1** (10 μM) + H_2A ; lane 7, DNA + **1** (8 μM) + H_2A ; Forms I and II are supercoiled and nicked circular forms of DNA respectively

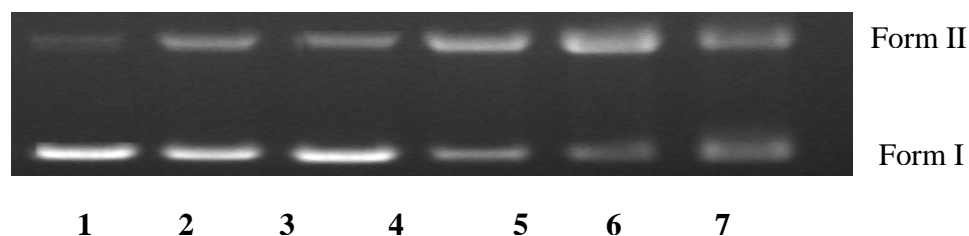


Figure S11. Concentration dependent DNA (40 μM) cleavage by the complex **2** (10 – 50 μM) in 5% DMF : 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^{\circ}\text{C}$ in the presence of ascorbic acid (H_2A , 10 μM) at 37 $^{\circ}\text{C}$. Lane 1, DNA ; lane 2, DNA + H_2A ; lane 3, DNA + **2** (10 μM) + H_2A ; lane 4, DNA + **2** (20 μM) + H_2A ; lane 5, DNA + **2** (30 μM) + H_2A ; lane 6, DNA + **2** (40 μM) + H_2A ; lane 7, DNA + **2** (50 μM) + H_2A ; ; lane 8, DNA + **2** (30 μM) + H_2A + 2 μL DMSO; Lane 9, DNA + **2** (30 μM) + H_2A + Distamycin. Forms I and II are supercoiled and nicked circular forms of DNA respectively.

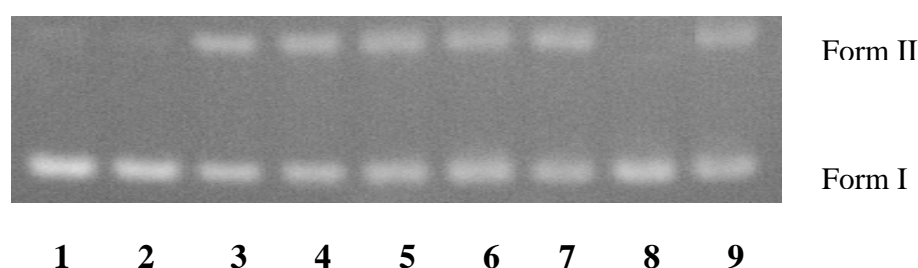


Figure S12. Concentration dependent DNA (40 μM) cleavage by the complex **3** (2 – 10 μM) in 10% DMF 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^{\circ}\text{C}$ in the presence of ascorbic acid (H_2A , 10 μM) at 37 $^{\circ}\text{C}$. Lane 1, DNA ; lane 2, DNA + H_2A ; lane 3, DNA + **3** (2 μM) + H_2A ; lane 4, DNA + **3** (4 μM) + H_2A ; lane 5, DNA + **3** (6 μM) + H_2A ; lane 6, DNA + **3** (8 μM) + H_2A ; lane 7, DNA + **3** (10 μM) + H_2A ; Forms I and II are supercoiled and nicked circular forms of DNA respectively

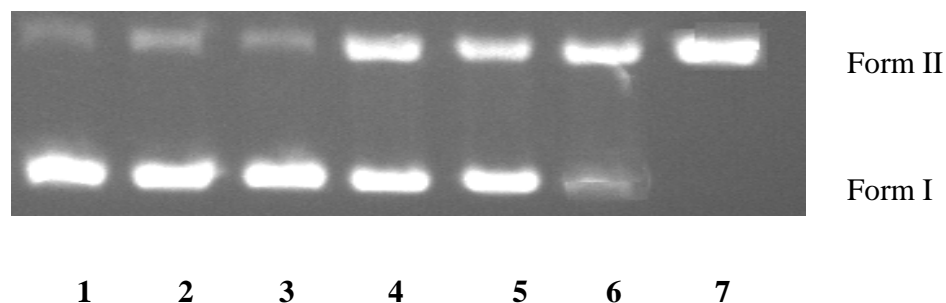


Figure S13. Concentration dependent DNA (40 μM) cleavage by the $[\text{Cu}(\text{dpq})_2\text{Cl}]^{2+}$ complex (2 – 10 μM) in 5% DMF : 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^\circ\text{C}$ in the presence of ascorbic acid (H_2A , 10 μM) at 37 $^\circ\text{C}$. Lane 1, DNA ; lane 2 , DNA + H_2A ; lane 3, DNA + $[\text{Cu}(\text{dpq})_2\text{Cl}]^{2+}$ (2 μM) + H_2A ; lane 4, DNA + $[\text{Cu}(\text{dpq})_2\text{Cl}]^{2+}$ (4 μM) + H_2A ; lane 5, DNA + $[\text{Cu}(\text{dpq})_2\text{Cl}]^{2+}$ (6 μM) + H_2A ; lane 6, DNA + $[\text{Cu}(\text{dpq})_2\text{Cl}]^{2+}$ (8 μM) + H_2A ; lane 7, DNA + $[\text{Cu}(\text{dpq})_2\text{Cl}]^{2+}$ (10 μM) + H_2A .

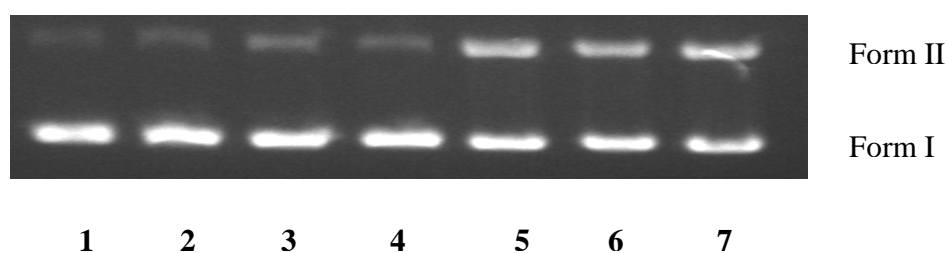


Figure S14. Cleavage of supercoiled pUC19 DNA (40 μM) by 10 μM complex (**1** – **3**) in 10% DMF / 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^\circ\text{C}$ in the presence of H_2O_2 (200 μM) at 37 $^\circ\text{C}$. Lane 1, DNA ; lane 2, DNA + H_2O_2 ; lane 3, DNA + H_2O_2 + **1**; lane 4, DNA + H_2O_2 + **3**; lane 5, DNA + H_2O_2 + **2**. Forms I and II are supercoiled and nicked circular forms of DNA respectively.

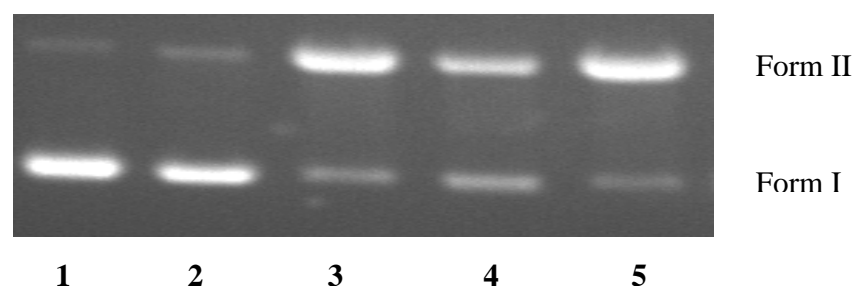


Figure S15. Cleavage of supercoiled pUC19 DNA (40 μM) by the copper(II) complexes in a 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^{\circ}\text{C}$ in the presence of H_2O_2 (200 μM) at 37 $^{\circ}\text{C}$. Lane 1, DNA + 2 μL DMSO; lane 2, DNA + H_2O_2 + 2 μL DMSO; lane 3, DNA + **1** + H_2O_2 + 2 μL DMSO; lane 4, DNA + **3** + H_2O_2 + 2 μL DMSO; lane 5, DNA + **2** + H_2O_2 + 2 μL DMSO; Complex concentration is 10 μM for lanes 3-5. Forms I, II and are supercoiled, nicked circular forms of DNA, respectively.

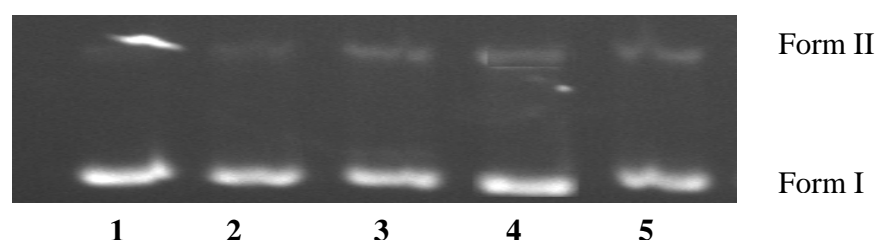


Figure S16. (A) Time dependent DNA (40 μM) cleavage by the complex **2** (10 – 60 min) in 10% DMF / 5 mM Tris-HCl/50 mM NaCl at pH = 7.1 and 37 $^{\circ}\text{C}$ in the presence of H_2O_2 (200 μM) at 37 $^{\circ}\text{C}$. Lane 1, DNA + H_2O_2 ; lane 2, DNA + **2** (10 min) + H_2O_2 ; lane 3, DNA + **2** (20 min) + H_2O_2 ; lane 4, DNA + **2** (30 min) + H_2O_2 ; lane 5, DNA + **2** (40 min) + H_2O_2 ; lane 6, DNA + **4** (50 min) + H_2O_2 ; lane 7, DNA + **3** (60 min) + H_2O_2 ; Forms I and II are supercoiled and nicked circular forms of DNA respectively.

