

SUPPLEMENTARY MATERIAL:

The phenanthridine biguanides efficiently differentiate between dGdC, dAdT and rArU sequences by two independent, sensitive spectroscopic methods.

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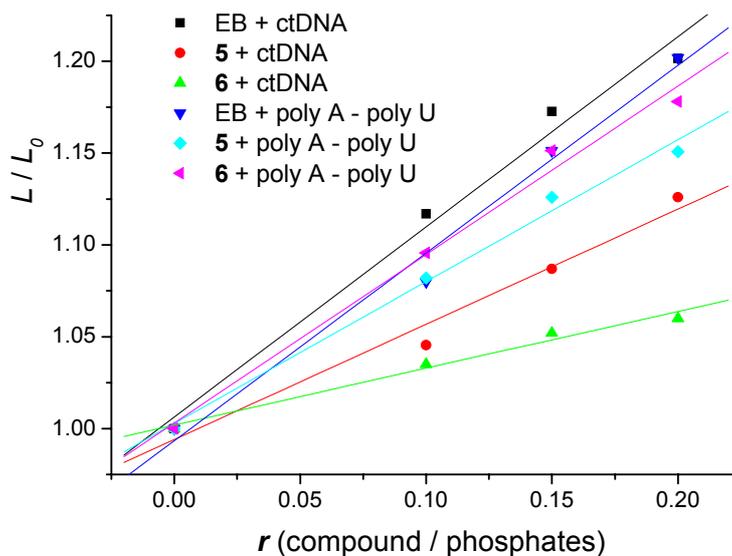


Figure S1. Relative ctDNA and poly A – poly U helix length extension (L/L_0) vs. ratio $r_{[\text{compound}] / [\text{polynucleotide}]}$ plot for **EB**, **5** and **6** at pH 7.0, sodium cacodylate buffer, $I = 0.05 \text{ mol dm}^{-3}$. Results: $\alpha = 0.62 \pm 0.06$ (for **5**), 0.31 ± 0.03 (for **6**) and 1.04 ± 0.09 (**EB**) with ctDNA; $\alpha = 0.77 \pm 0.05$ (for **5**), 0.91 ± 0.06 (for **6**) and 1.02 ± 0.08 (**EB**) with poly A – poly U.

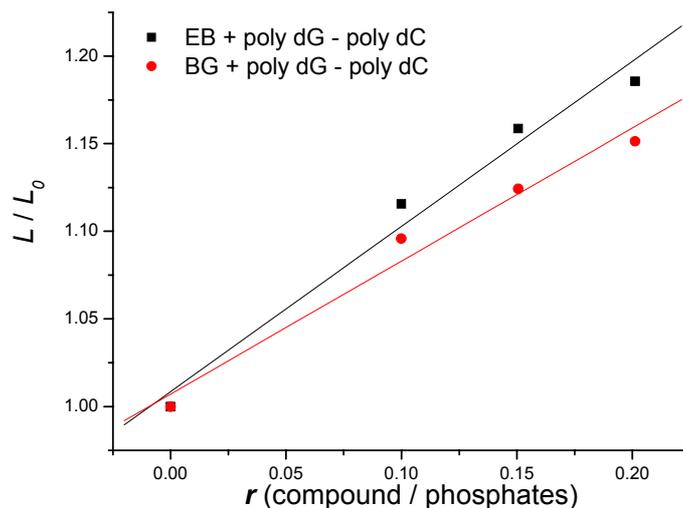


Figure S2. Relative poly dG - poly dC helix length extension (L/L_0) vs. ratio $r_{[\text{compound}]} / [\text{polynucleotide}]$ plot for **EB** and **5** at pH 7.0, sodium cacodylate buffer, $I = 0.05 \text{ mol dm}^{-3}$. Result: $\alpha_5/\alpha_{\text{EB}} = 0.805$.

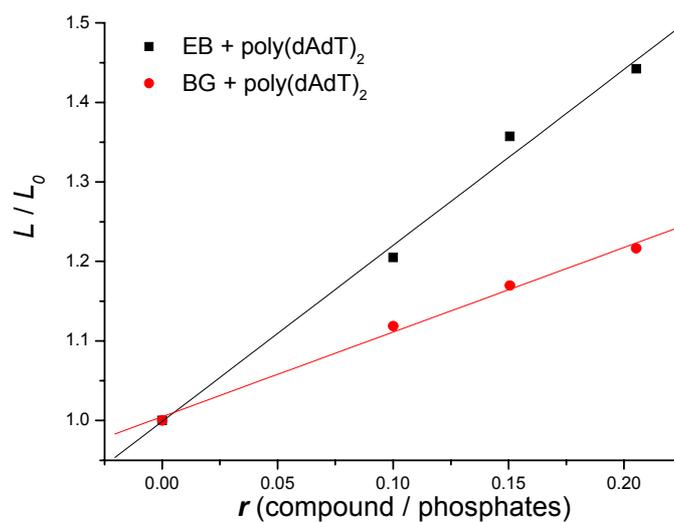


Figure S3. Relative poly(dAdT)₂ helix length extension (L/L_0) vs. ratio $r_{[\text{compound}]} / [\text{polynucleotide}]$ plot for **EB** and **5** at pH 7.0, sodium cacodylate buffer, $I = 0.05 \text{ mol dm}^{-3}$. Result: $\alpha_5/\alpha_{\text{EB}} = 0.48$.

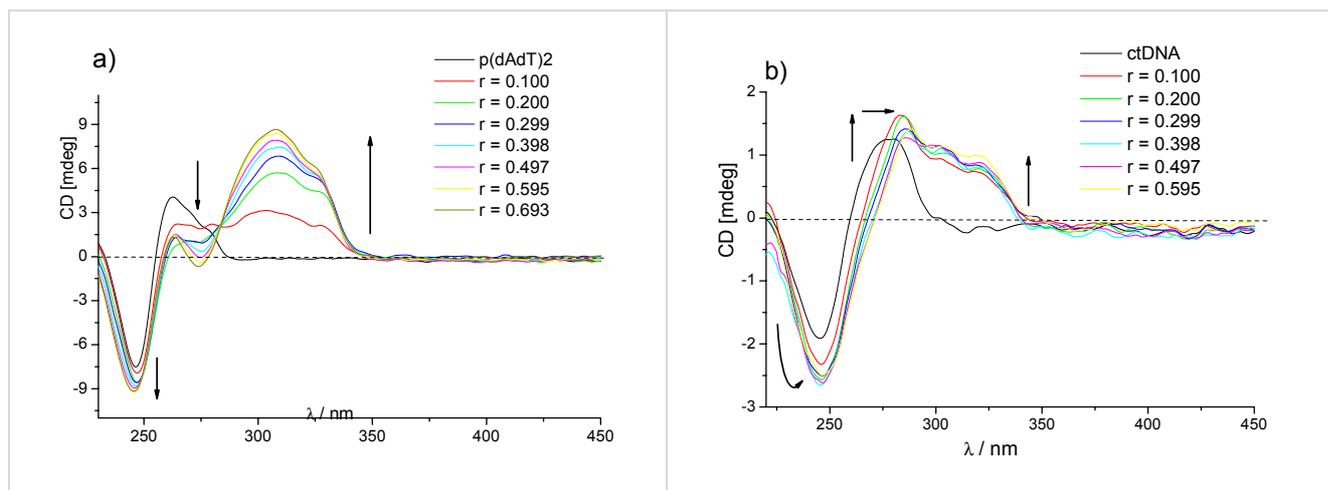


Figure S4. CD titrations of a) poly (dAdT)₂ and b) ctDNA ($c = 3.0 \times 10^{-5} \text{ mol dm}^{-3}$) with **6** at molar ratios $r = [\text{compound}] / [\text{polynucleotide}]$ (pH 5.0, citric acid/NaOH buffer, $I = 0.03 \text{ mol dm}^{-3}$).

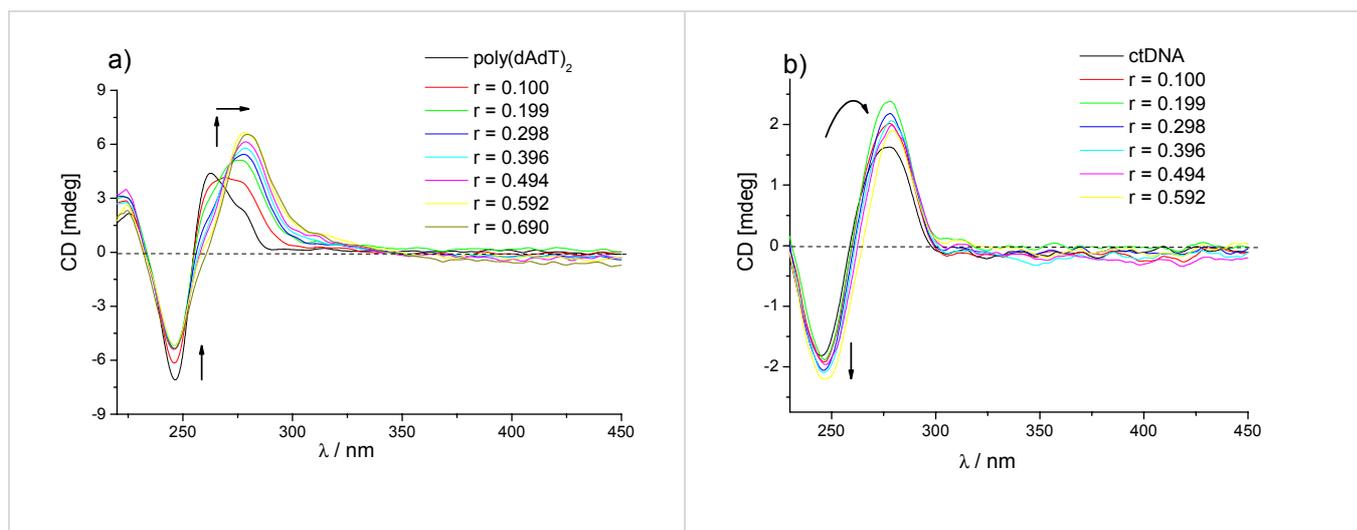


Figure S5. CD titrations of a) poly (dAdT)₂ and b) ctDNA ($c = 3.0 \times 10^{-5} \text{ mol dm}^{-3}$) with **5** at molar ratios $r = [\text{compound}] / [\text{polynucleotide}]$ (pH 5.0, citric acid/NaOH buffer, $I = 0.03 \text{ mol dm}^{-3}$).

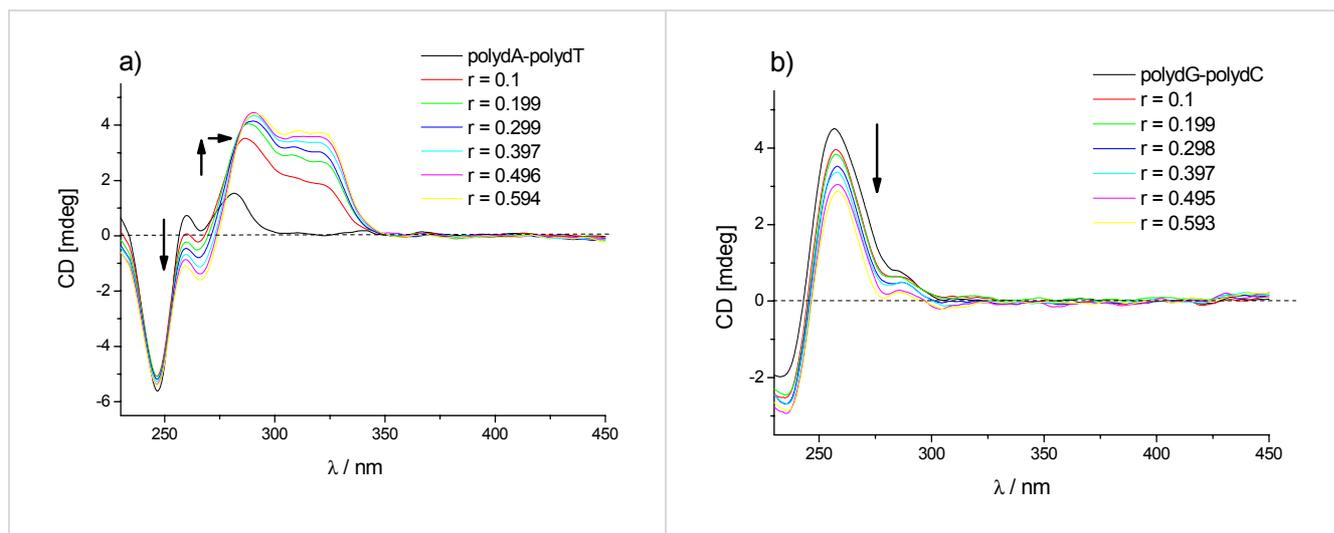


Figure S6. CD titrations of a) poly dA – poly dT and b) poly dG – poly dC ($c = 3.0 \times 10^{-5}$ mol dm⁻³) with **6** at molar ratios $r = [\text{compound}] / [\text{polynucleotide}]$ (pH 7.0, buffer sodium cacodylate, $I=0.05$ moldm⁻³).

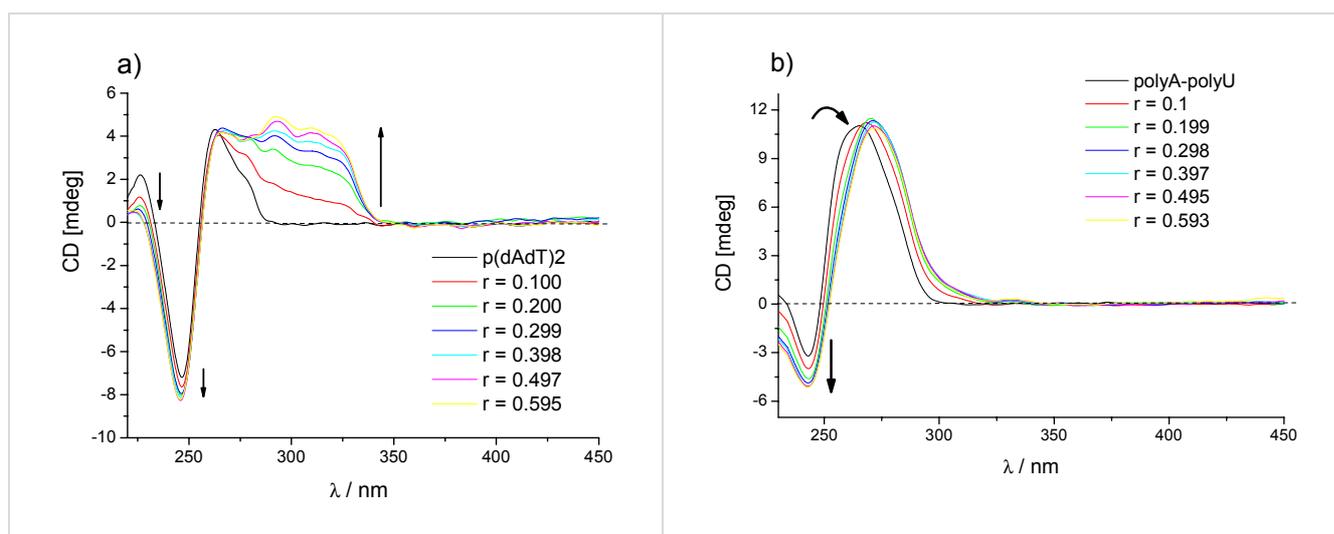


Figure S7. CD titrations of a) poly (dAdT)₂ and b) poly A – poly U ($c = 3.0 \times 10^{-5}$ mol dm⁻³) with **6** at molar ratios $r = [\text{compound}] / [\text{polynucleotide}]$ (pH 7.0, buffer sodium cacodylate, $I=0.05$ moldm⁻³).

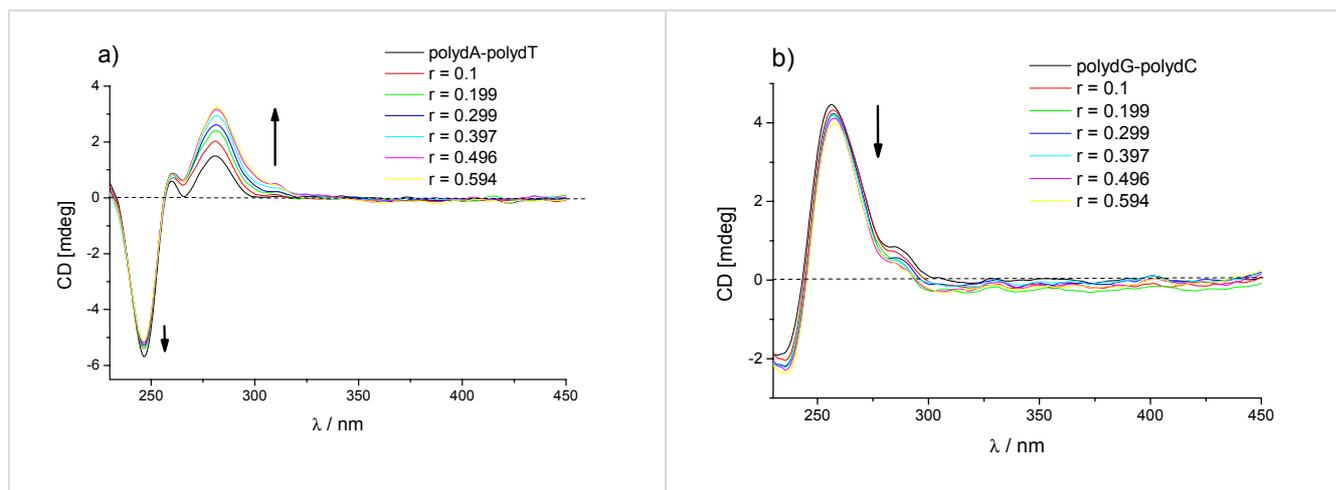


Figure S8. CD titrations of a) poly dA – poly dT b) poly dG – poly dC ($c = 3.0 \times 10^{-5}$ mol dm⁻³) with **5** at molar ratios $r = [\text{compound}] / [\text{polynucleotide}]$ (pH 7.0, buffer sodium cacodylate, $I=0.05$ mol dm⁻³).

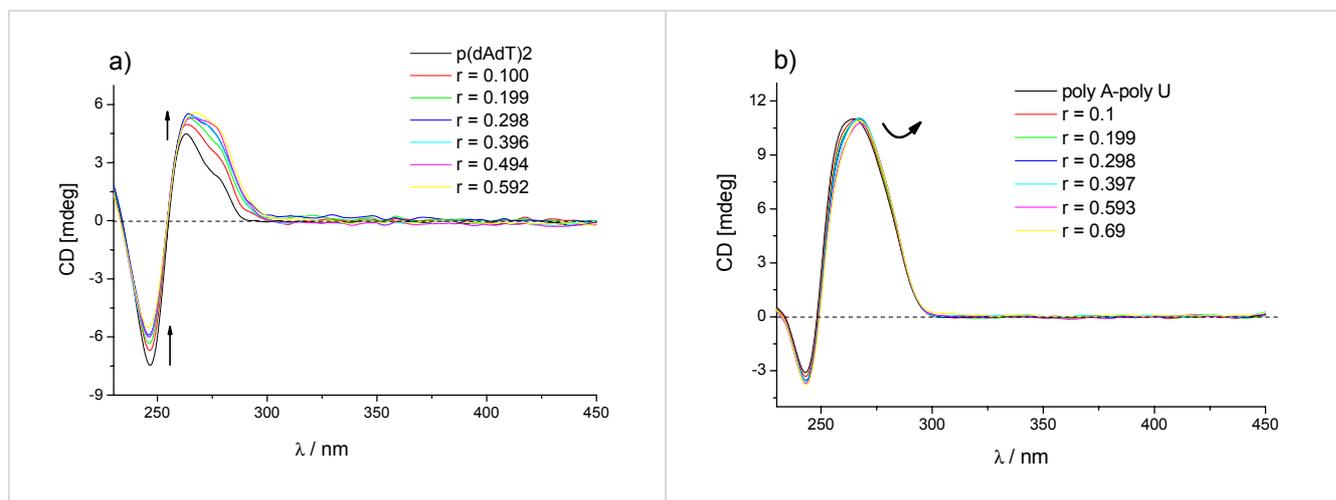


Figure S9. CD titrations of a) poly (dAdT)₂ and b) poly A – poly U ($c = 3.0 \times 10^{-5}$ mol dm⁻³) with **5** at molar ratios $r = [\text{compound}] / [\text{polynucleotide}]$ (pH 7.0, buffer sodium cacodylate, $I=0.05$ mol dm⁻³).

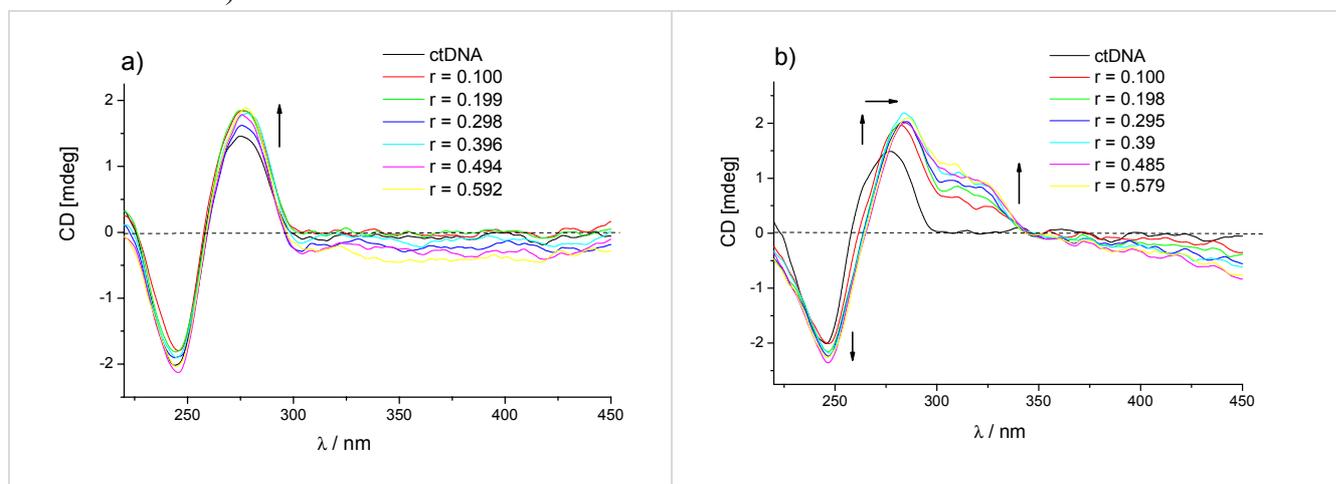


Figure S10. CD titrations of ctDNA ($c = 3.0 \times 10^{-5}$ mol dm⁻³) with a) **5** and b) **6** at molar ratios $r = [\text{compound}] / [\text{polynucleotide}]$ (pH 7.0, buffer sodium cacodylate, $I=0.05$ mol dm⁻³).

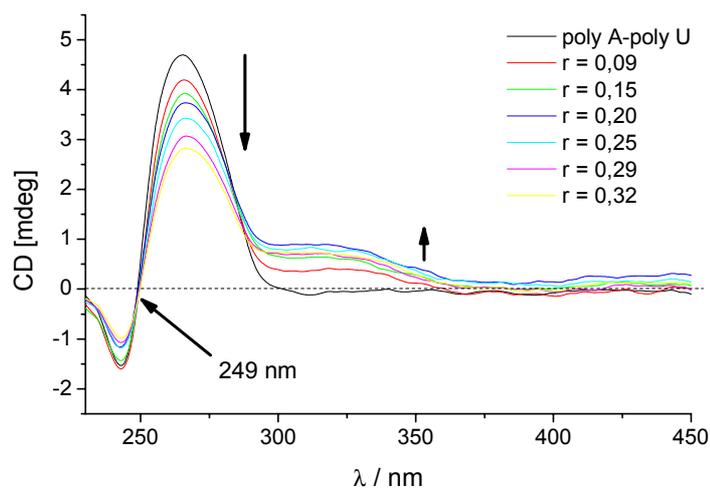


Figure S11. CD titration¹ of poly A-poly U ($c = 1.5 \times 10^{-5} \text{ mol dm}^{-3}$) with **9** at molar ratios $r = [\text{compound}] / [\text{polynucleotide}]$ (pH 5.0, citric acid buffer, $I = 0.03 \text{ mol dm}^{-3}$).

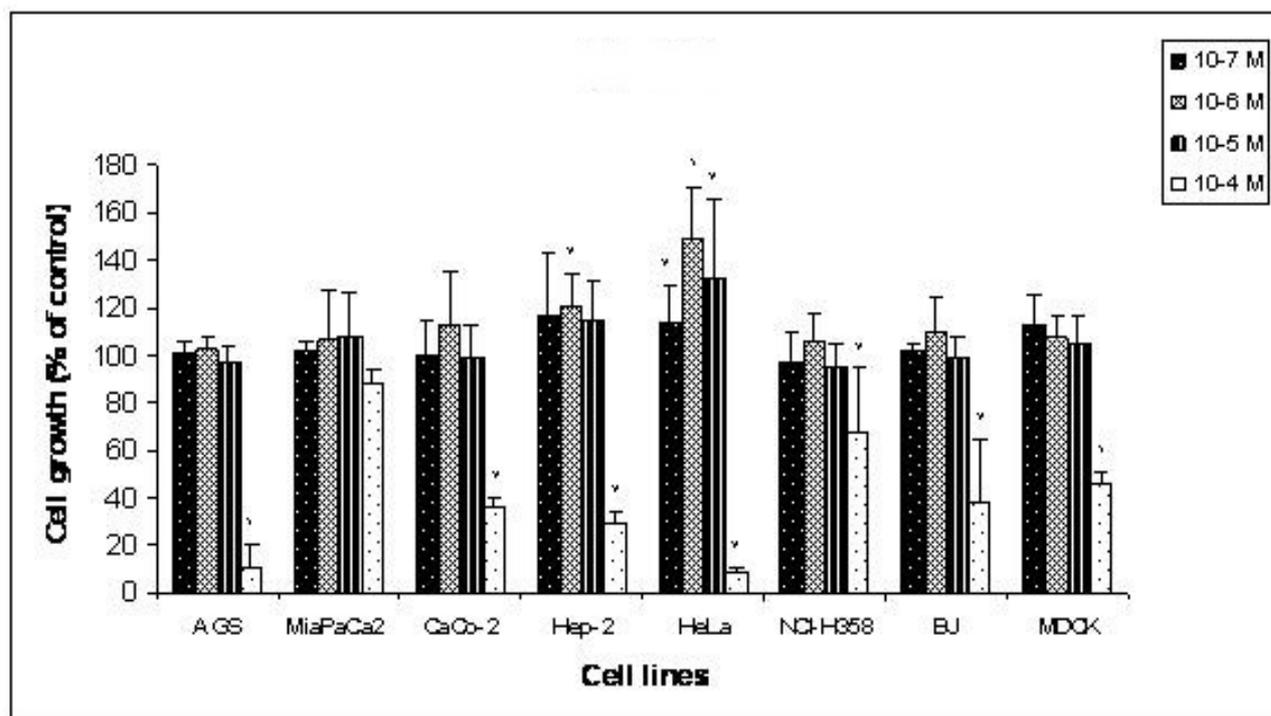


Figure S12: Cytotoxic effects of compound **5** on tumor and normal cell lines growth after 72 h of incubation in final concentration range (10^{-4} – 10^{-7} mol/dm^3). Data are presented as mean value \pm SD of three independent experiments done in three plicate. Statistically significant change ($p < 0.05$) is presented by (*).

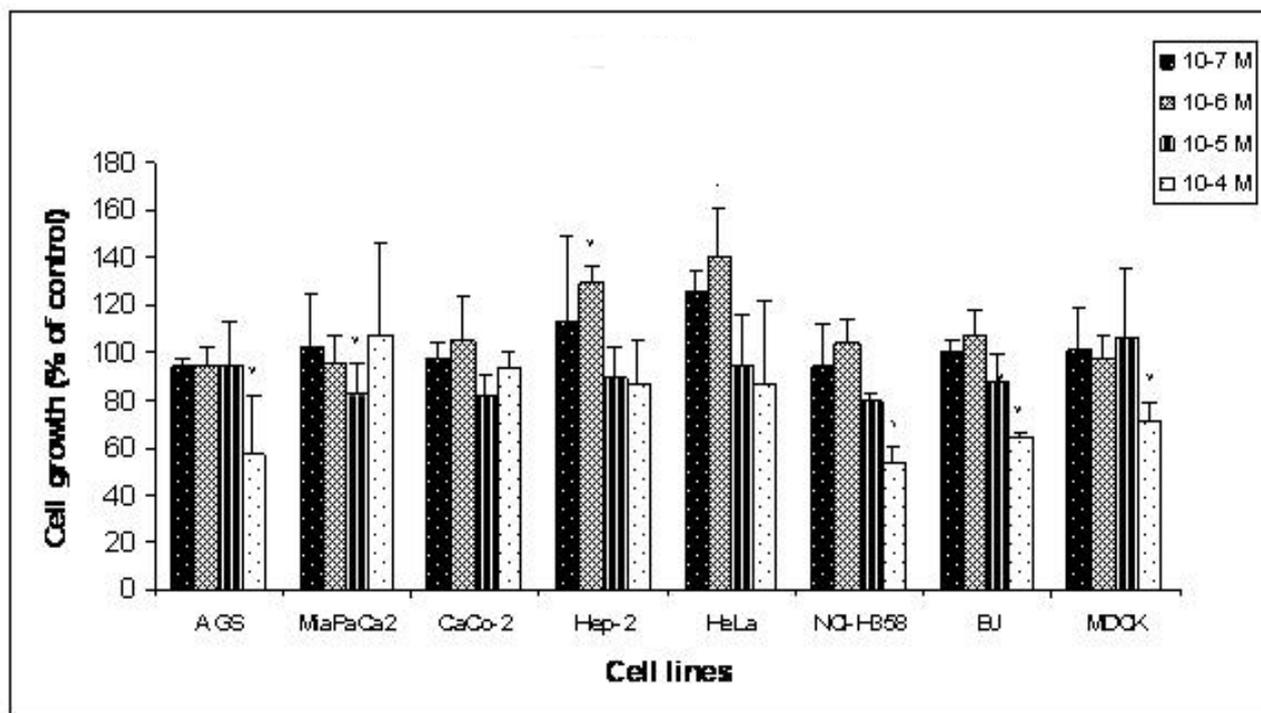


Figure S13: Cytotoxic effects of compound **6** on tumor and normal cell lines growth after 72 h of incubation in final concentration range (10^{-4} – 10^{-7} mol/dm³). Data are presented as mean value \pm SD of three independent experiments done in three plicate. Statistically significant change ($p < 0.05$) is presented by (*).

Table S1. Preliminary assignment of the vibrational bands observed in the SERS spectra of the **5** and **6** and their complexes with ctDNA, [**5,6**]/[DNA] 1/20

Wavenumber / cm ⁻¹				Assignment
5	5 /DNA 1/20	6	6 /DNA 1/20	
1623	1624	1621	1620	v ring (v CC/CN), v CN
1616 sh	1612			v CN, δ NH ₂
1582	1583	1591 sh	1592 sh	v ring (v CC/CN)
1536	1537	1543	1542	v ring (v CC)
1467	1466	1465 sh	1465 sh	δ_{as} CH ₃
		1449	1448	v ring (v CC)
1422	1419 sh	1426	1419 sh	v _s CNC, v CN ₃ , δ NH ₂
			1414	v ring (v CC)
1391	1392	1389	1388	v ring (v CC)
1365	1366	1361	1362	v ring (v CC), δ_s CH ₃
1345	1346	1328 sh	1328 sh	v ring (v CC)
1298	1300	1277	1284	δ_{ip} CH, δ NH
1258	1257			δ_{ip} CH
1229	1227	1221	1222	δ_{ip} CH, v _{as} CNC, v CN ₃

1210	1212			δ_{ip} CH, δ NH
		1185		δ_{ip} CH
1154	1157	1154	1154	δ_{ip} CH
		1138	1137	δ_{ip} CH
1076	1081	1079	1080	δ NH ₂ , ν CN ₃
1039	1039			ν ring (ν CC)
1007	1004			ν CN ₃ , δ NH ₂
		971	966 sh	δ ring
952	952	952 sh	954	δ ring
923	924	911	911	δ CNC, ν CN, δ NH ₂
893	891			ν ring (ν CC)
869	870	870	867	ν ring (ν CC), δ_{oop} CH
837				δ ring
		797	797	δ ring, δ CNC
727	729	722	729	δ ring
		706 sh	708	δ ring
		696	696	δ ring
655	655	679	679	δ ring
617	618 sh			δ ring
588	600			δ NH ₂ , δ NH, δ ring
513 sh	516	520	520	δ CNC, δ ring
506	504			δ ring
462	460 sh	463	459	δ_{as} CNC
448	447	435	435	δ ring
421	421	395	395	δ CN ₃ , δ NH ₂
341	339	328	326 sh	δ_s CNC
227	224	227	223	ν Ag–N

Abbreviations: sh, shoulder; ν , stretching; δ , deformation; s, symmetrical; as, antisymmetrical; ip, in plane; oop, out of plane.

¹ M. Radić Stojković, I. Piantanida, Tetrahedron 64 (2008) 7807-7814