

Supplementary Material

DNA-binding properties of cytotoxic naphthindolizinedione-carboxamides acting as type II Topoisomerase inhibitors. A combined *in silico* and experimental study

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Figure S1. View by docking calculation of compound **3** interacting inside the topoisomerase II (PDB ID: 3QX3) cleavage site: in free base form (a) and in protonated form (b). The tyrosine residue mediating DNA cut is drawn in magenta, H-bridge is shown by green circles, the phosphate-deoxyribose bond broken by tyrosine is depicted in green dashed line.

Figure S2. UV-Vis spectra used for the calculation of molar extinction coefficient, recorded in 240-500 nm range, for buffer solutions (Tris-HCl 10mM, NaCl 20mM, pH= 7.5) of compounds. (a) **1**, (b) **2**, (c) **3**.

Figure S3. DNA thermal denaturation based on melting temperature (T_m) analysis by compound **1-3** in comparison with mitoxantrone.

Figure S4. Overlapped CD spectra of DNA complexes using comparable concentrations of each ligand **1-3** and mitoxantrone, in comparison with the spectrum of free *Calf thymus* DNA.

Figure S5. Inhibition of human topoisomerase II α decatenation assay by compounds **1-3** used at increased concentrations (0.1-1-10-100 μ M); B, blank; C, control (+enzyme).

Figure S6. Inhibition of human topoisomerase II α decatenation assay by mitoxantrone: 200ng of kDNA were incubated with increasing concentrations (0.1-0.5-1-5 -10 μ M) of mitoxantrone and human topoisomerase II α and loaded in 1% agarose gel in TBE (90 mM Tris, 90 mM boric acid, 20 mM EDTA); running buffer TBE; gel run was 1.5 h at 4 V/cm. B, blank; C, control (+enzyme). Staining with ethidium bromide 0.5 μ g/mL.

Table S1. Equilibrium constants and T_m values for the indicated *Calf thymus* DNA complexes, in 10 mM TRIS buffer solution, using each ligand as hydrochloride salts.

Table S2. Inhibition of human topoisomerase II α decatenation assay IC₅₀ values for **1-3** and mitoxantrone.

1. General Experimental Methods

Deoxyribonucleic acid sodium salt from *Calf thymus* (CG = 41.9%) and dihydrochloride mitoxantrone were from Sigma. Synthetic CGCGAATTCGCG oligonucleotide (GC = 66.7%) was bought from Thermo Fisher Scientific. Reagents and solvents were used without purification. All evaporations were carried out at room temperature at reduced pressure. UV-Visible spectra were recorded with a double beam spectrophotometer UVIKON 923. The DNA melting point has been obtained by JASCO V-550 UV/Vis Spectrophotometer equipped with a thermostatic unit. CD spectra were recorded with a Jasco J-710 spectrophotometer. pH Measurements were obtained by pH-meter Delta OHM H2256.

2. Preparation and use of (CGCGAATTCGCG)₂ oligonucleotide

Single strand oligonucleotide, dissolved in DNase/RNase free 10mM Tris, was heated at 85°C for 5 minutes and slowly cooled during 12 hours till room temperature. It was used in UV analysis of the complex with compound **2** at a fixed concentration (18.3 μ M) and at increasing oligonucleotide amounts (from 1.4 to 5.6 μ M).

3. Spectroscopic Analyses

3.1. Binding constants evaluation by UV/Vis spectroscopy

A stock solution of *Calf thymus* DNA was diluted in 10 mM Tris (pH 5.8) and the exact concentration was determined by UV/vis spectroscopic analysis ($\lambda = 260$ nm, $\epsilon = 13200$ M⁻¹ cm⁻¹ established relatively to base pairs). The same procedure has been applied for protonated salts of compounds **1** ($\lambda = 331$ nm, $\epsilon = 8791$ M⁻¹ cm⁻¹), **2** ($\lambda = 323$ nm, $\epsilon = 12208$ M⁻¹ cm⁻¹), **3** ($\lambda = 339$ nm, $\epsilon = 8710$ M⁻¹ cm⁻¹) and mitoxantrone ($\lambda = 660$ nm, $\epsilon = 19684$ M⁻¹ cm⁻¹). DNA solution (in the range 1,6 ÷ 16 μ M) was incubated individually with compound **1**, **2**, **3** or mitoxantrone at 20 μ M for 90 minutes at room temperature in the dark. The complexes and the controls were subjected to spectroscopic analysis by a double beam spectrophotometer UVIKON 923 equipped with quartz cuvettes. The calculation of binding constants was achieved following two different approaches reported in literature[1,2], obtaining comparable results (Figure S2).

3.2. Thermal denaturation analysis

DNA melting experiments were carried out by monitoring the absorbance (260 nm) of *Calf thymus* DNA (14 μ M base-pairs) at different temperatures, ranged between 40 and 90 °C, in the absence and presence of the complex, in a 1:1 DNA/complex ratio. Measurements were performed by a JASCO V-550 UV/Vis spectrophotometer equipped with a thermostatic unit. The solution containing the complex and *Calf thymus* DNA in 10 mM Tris buffer (pH 5.8) was heated with a rate of temperature increase of 1°C/ min and reading the absorbance at $\Delta T = 0.25$ °C- (Figure S3).

3.3. Circular dichroism

It was performed using a Jasco J-710 spectrophotometer. The spectra of *Calf thymus* DNA (60 μ M) or synthetic oligonucleotide in 10 mM Tris pH 5.8 and in the absence and presence of various concentrations of each compound **1-3** and mitoxantrone was recorded in the range of 220– 305 nm with a spectral resolution of 1 nm. The scan speed was 20 nm/min and the response time was 0.3330 sec with a band width of 1 nm. Quartz cell with a path length of 10 mm was used and all measurements were carried out at 25°C. Results are expressed as molar ellipticity $[\theta]$, in deg \times cm² \times dmol⁻¹ (Figure S4).

4. Human topoisomerase II α inhibition

The assay was performed according to the reported protocol [3] with minor modifications. The reaction was made in a final volume of 20 μ l; each sample was constituted by 200ng of kDNA, assay buffer (containing Tris HCl 50mM pH=7.5, NaCl 125mM, MgCl₂ 10mM, DTT 5mM and albumin 100 μ g/ml), ATP 10mM, 1U of human topoisomerase II α (provided by Inspiralis Ltd, UK) and 2 μ l of compound solution, to have the desired final concentration. Samples were incubated for 60 min at 37°C and then stopped with 4 μ l of gel loading buffer made with 49% TE (10mM Tris, 1mM EDTA, 20mM NaCl, pH 7.4), 49% glycerol, 2% SDS, 0.025% bromophenol blue, and 0.025% xylene cyanol. Samples are resolved by electrophoresis on a 1% agarose gel running in 1X TBE

buffer (90mM Tris, 90mM boric acid, 20mM EDTA); the arrays for the electrophoresis are the Sub-Cell GT type provided by Bio-Rad. Gel was stained post run with an ethidium bromide solution 0.5 μ g/ml for 30 min. The bands were detected using the Geliance 600 Imaging System and densitometric analysis was performed using the software Gene tools from Perkin-Elmer. In each gel a blank control containing only kDNA, which correspond to 0% of activity, and a control containing kDNA end enzyme, corresponding to 100% of topoisomerase II α activity, were added (Figures S5 and S6). The concentration of the inhibitor preventing 50% of the substrate (knotted DNA) from being converted into the reaction product (unknotted DNA) was determined from a standard sigmoid curve. By averaging at least three experiments, the IC₅₀ values and the standard deviation were determined.

5. References

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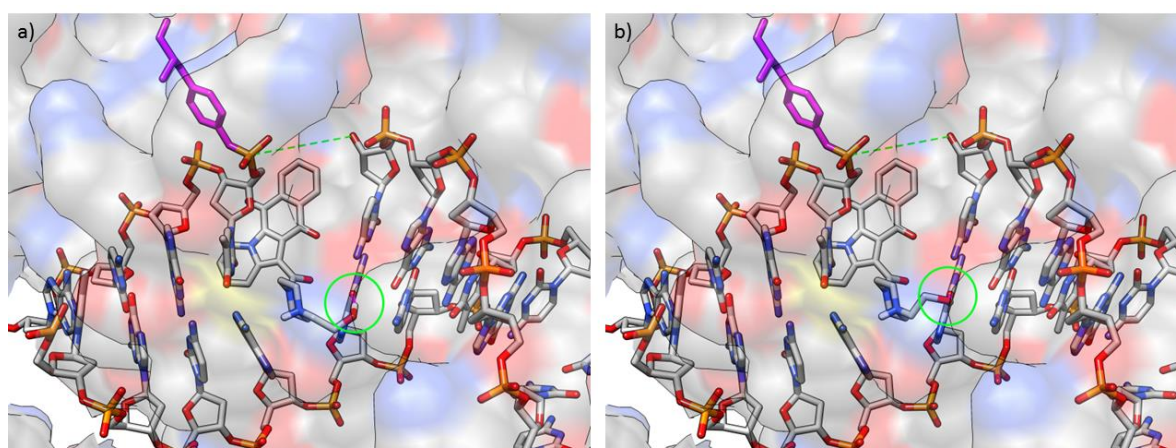


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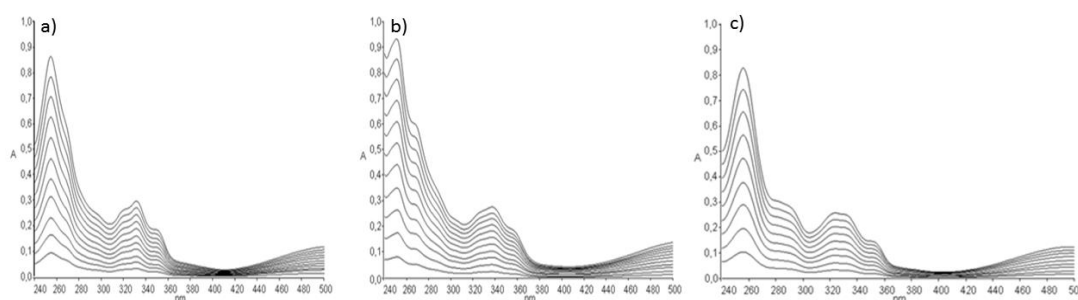


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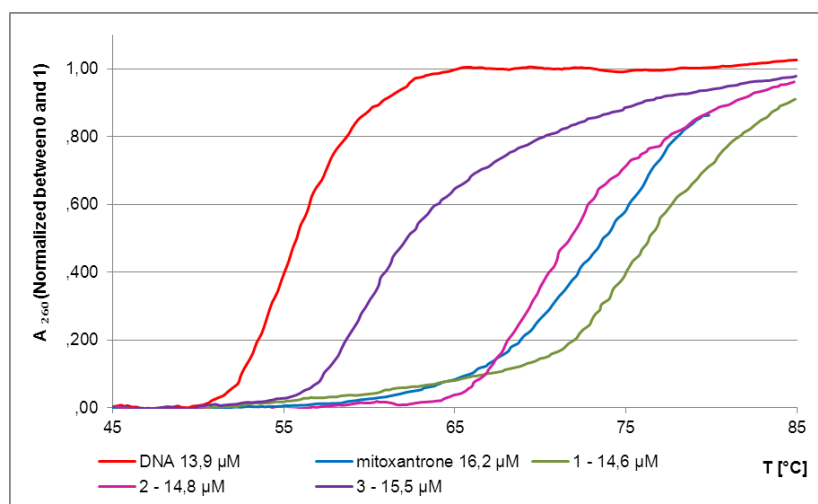


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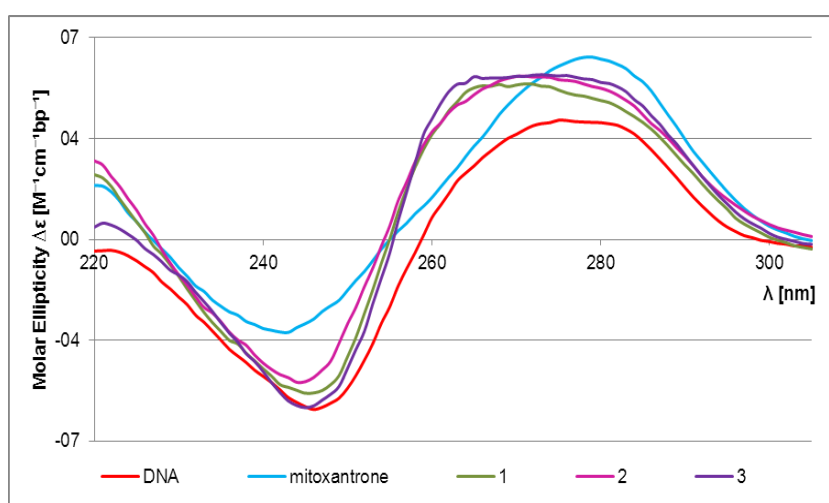


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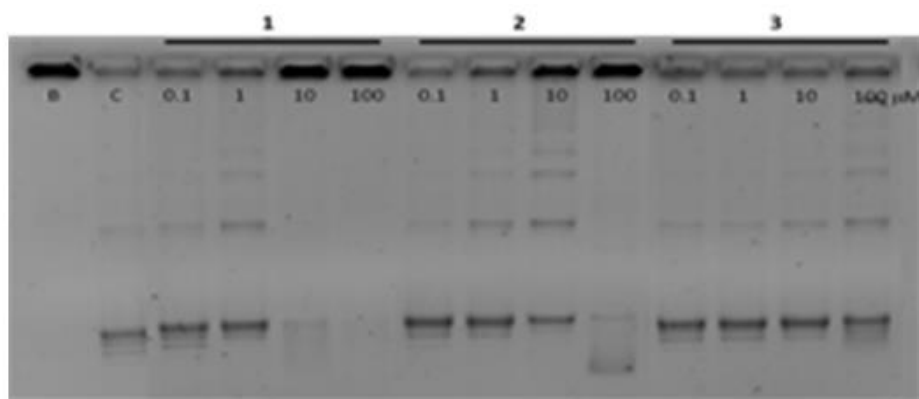


Figure S5. Inhibition of human topoisomerase II α decatenation assay by compounds 1-3 used at increased concentrations (0.1-1-10-100 μ M); B, blank; C, control (+enzyme).

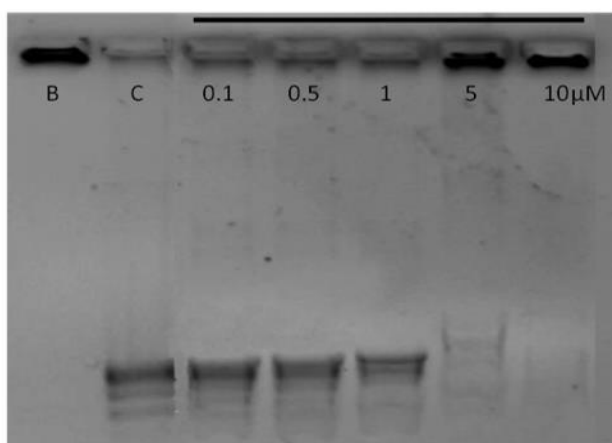


Figure S6. Inhibition of human topoisomerase II α decatenation assay by mitoxantrone: 200ng of kDNA were incubated with increasing concentrations (0.1-0.5-1-5 -10 μ M) of mitoxantrone and human topoisomerase II α and loaded in 1% agarose gel in TBE (90 mM Tris, 90 mM boric acid, 20 mM EDTA); running buffer TBE; gel run was 1.5 h at 4 V/cm. B, blank; C, control (+enzyme). Staining with ethidium bromide 0.5 μ g/mL.

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System	K [M ⁻¹]	T _m [°C]	ΔT _m [°C]
DNA	-	55.8±0.1	-
DNA/1	(2,2 ± 0,3)·10 ⁶	76.3±0.1	20.5±0.2
DNA/2	(3,2 ± 0,3)·10 ⁶	71.7±0.1	15.9±0.2
DNA/3	(3,6 ± 0,1)·10 ⁵	62.2±0.1	6.4±0.2
DNA/Mitoxantrone	(2,3 ± 0,1)·10 ⁵	73.6±0.1	17.8±0.2

Table S2. Inhibition of human topoisomerase IIα decatenation assay: IC₅₀ values for 1-3 and mitoxantrone

Compound	IC ₅₀ (μM)
1	1.78±0.32
2	7.83±3.33
3	>100
Mitoxantrone	2.54±0.25