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Cite this: DOI: 10.1039/c0xx00000x

www.rsc.org/xxxxxx

ARTICLE TYPE

Enhancement of antiproliferative activity by phototautomerization of anthrylphenols

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Received (in XXX, XXX) Xth XXXXXXXXX 20XX, Accepted Xth XXXXXXXXX 20XX

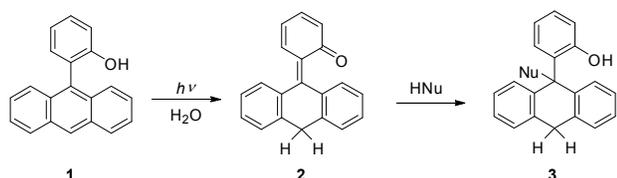
DOI: 10.1039/b000000x

Antiproliferative investigation has been conducted on 3 human cancer cell lines HCT 116 (colon), MCF-7 (breast), and H 460 (lung), on a series of 4 anthrylphenols in dark, and upon exposure to light (350 nm). 9-(2-Hydroxyphenyl)anthracene (**1**) moderately inhibits the proliferation, but the irradiation considerably enhances the effect. The other investigated anthracenes **4-6** exhibit antiproliferative activity in dark, which was not enhanced upon irradiation. The enhancement of the antiproliferative effect on irradiation of **1** was rationalized as being due to the formation of quinone methide (QM **2**) by excited state proton transfer. QM **2** acts as an electrophilic species capable of reacting with biological molecules. Although QM **2** reacts with nucleotides, adducts could not be isolated. On the contrary, cysteine adduct **8** has been isolated and characterized, whereas adducts with glycine, serine and tripeptide glutathione have been characterized by MS. Non-covalent binding of **1** to DNA and bovine serum albumin was demonstrated by UV-vis, fluorescence and CD spectroscopy. However, straightforward conclusion about the DNA or protein alkylating (damaging) ability of **2** could not be drawn. The results obtained by the irradiations of **1** in the presence of DNA, amino acids and peptides, the cell cycle perturbation analysis, and *in vitro* translation of GFP suggest that the effect is not only due to the damage of DNA but also the impact on the cellular proteins. Considering that up to date all QM agents were assumed to target dominantly DNA, this is an important finding with impact to the further development of anticancer agents based on QMs.

Introduction

Quinone methides (QMs) are ubiquitous intermediates in chemistry of phenols.¹ The interest in chemistry of QMs has been initiated owing to their applications in chemical synthesis^{2,3} and biological activity.^{4,5} It has been demonstrated that QMs react with some enzymes, such as tyrosine hydroxylases,⁶ β -lactamase,⁷ β -glucosidases,⁸ phosphatase^{9,10} or ribonuclease-A.¹¹ Furthermore, QMs react with nucleosides¹²⁻¹⁵ and induce alkylation of DNA.¹⁶⁻¹⁹ Some antineoplastic agents such as mitomycin,²⁰⁻²² exert their antiproliferative action on metabolic formation of QMs that alkylate DNA. Moreover, selectivity of some naphthalene diimide-QM derivatives towards guanine-quadruplex structures has recently been demonstrated²³⁻²⁵ and reversible DNA alkylation abilities of QMs reviewed.²⁶ QMs can be formed in mild conditions in the photochemical reactions^{27,28} of dehydration,²⁹ or deamination^{30,31} from the appropriately substituted phenol derivatives. Furthermore, Wan and co-workers have demonstrated that QMs can be formed from 2-phenylphenol in excited state intramolecular proton transfer reactions (ESIPT) from phenolic OH to a carbon atom of the adjacent phenyl ring.^{32,33} The scope of the reaction has been extended to ESIPT in naphthylphenols,³⁴ BINOLs³⁵ and anthrylphenols.³⁶⁻³⁸ Thus, H₂O-assisted ESIPT to the anthracene

position 10 in **1** gives QM **2** that reacts with nucleophiles (H₂O, alcohols, amines) and give addition products **3** (Scheme 1).³⁶ Since reactions of the photogenerated QMs can be applied in the biological systems for the cross-linking of DNA,³⁹⁻⁴³ and for the increase of antiproliferative effect,^{41,43-46} we probed if QMs formed by ESIPT to carbon can enhance antiproliferative activity. Herein we report an investigation of antiproliferative activity on a series of anthrylphenols **1**, and **4-6** on three human cancer cell lines HCT 116 (colon), MCF-7 (breast), and H 460 (lung) with and without exposure to irradiation, and compared to the activity of psoralen **7**. An enhancement of the antiproliferative effect was observed for **1**, so additional experiments were performed to elucidate the mechanism of the antiproliferative action including assessment of non-covalent binding to DNA and proteins, alkaline DNA electrophoresis, inhibition of green fluorescence protein (GFP) translation *in vitro* and influence on cell cycle of tumor cells. The results presented herein demonstrate that DNA is not the only target of QMs leading to antiproliferative activity, as it has been postulated in most of the reports. Proteins⁴⁷ and enzymes⁶⁻¹¹ are also viable QM targets.



Scheme 1 Photochemical formation of QM 2 by H_2O -assisted ESIP and reaction with nucleophiles.

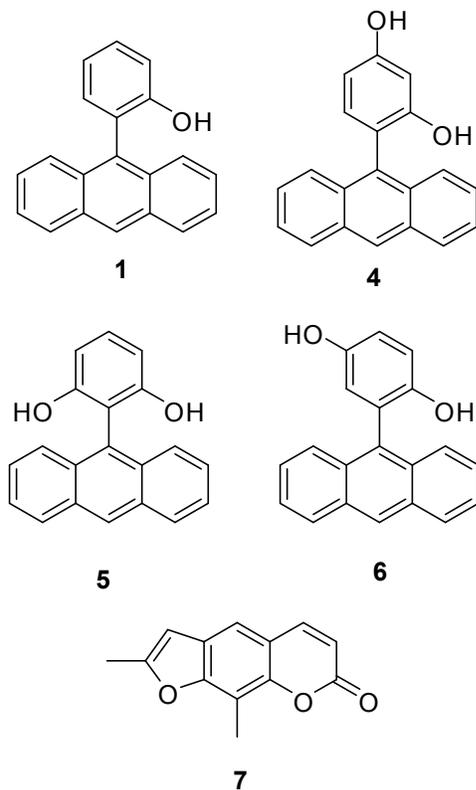


Fig. 1 Structures of anthrylphenols **1**, **4–6** undergoing H_2O -assisted ESIP and psoralene derivative **7**.

Experimental

General. Compounds **1** and **4–6** were prepared according to the previously described procedure.^{36–38} Chemicals for the synthesis were purchased from the usual suppliers, whereas solvents for the synthesis and chromatographic separations were purified by distillation, or used as received (p.a. grade). Prior to spectroscopic investigations **1** was additionally purified by crystallization from dichloromethane/cyclohexane. ^1H and ^{13}C NMR spectra were recorded on a Bruker AV-300, 500 or 600 MHz. The NMR spectra were taken in CDCl_3 or $\text{DMSO}-d_6$ at rt using TMS as a reference. For the sample analysis a Shimadzu HPLC equipped with a Diode-Array detector and a Phenomenex Luna 3u C18(2) column was used. Mobile phase was $\text{CH}_3\text{OH}-\text{H}_2\text{O}$ (20 %). Alternatively, an HPLC Agilent 1200 Series and triple quadrupole mass spectrometer Agilent 6410 were used. MS were obtained also on a Amazon ETD, Bruker Daltonik (ESI-MS or MALDI-TOF). For the chromatographic separations silica gel (Merck 0.05–0.2mm) or aluminum oxide (activity IV/V) were used. Analytical thin layer chromatography was performed on

Polygram® SILG/UV₂₅₄ (Machery-Nagel) plates. In the irradiation experiments, CH_3CN was of HPLC purity and mQ- H_2O (Millipore) was used. ct-DNA and BSA were purchased from Aldrich.

Analytical Irradiation Experiments in the presence of DNA bases. Anthracene **1** (15 mg, 0.055 mmol) was dissolved in CH_3CN (60 mL) and divided into four cuvettes (4×15 mL) to which aqueous solutions of sodium phosphate buffer (4×2 mL, pH = 6.8, 0.1M) were added. To the first cuvette, adenine (2mg, 0.015 mmol) was added, to the second cytosine (2mg, 0.018 mmol) and to the third guanine (2 mg, 0.013 mmol). The solutions were purged with Ar for 30 min, sealed with a septum and simultaneously irradiated in a Luzchem reactor equipped with 8 lamps with the output at 350 nm over 1 h. The composition of the irradiated solutions was analyzed by HPLC (see the supplement).

Preparative irradiation in the presence of cytosine. Anthracene **1** (20 mg, 0.074 mmol) was dissolved in CH_3CN (30 mL) and mixed with an aqueous solution of sodium phosphate buffer (20 mL, pH = 6.8, 0.1M) and cytosine (20 mg, 0.18 mmol) suspended in CH_3CN (100 mL). The resulting mixture was purged with Ar for 30 min and irradiated in a Rayonet reactor equipped with 11 lamps with the output at 350 nm during 1 h. The course of the reaction was followed by HPLC. During the irradiation the solution was continuously purged with Ar and cooled with a tap- H_2O finger-condenser. After the irradiation, H_2O (50 mL) was added and extraction with CH_2Cl_2 (2×75 mL) and EtOAc (2×75 mL) was carried out. The organic extracts were combined and dried over anhydrous MgSO_4 . After filtration, the solvent was removed on a rotary evaporator and the residue chromatographed on a thin layer of silica using $\text{CH}_2\text{Cl}_2/\text{EtOAc}$ (10%) as eluent. In addition to **1** that is recovered, a fraction (5 mg) was isolated that contained mixture of compounds with possible cytosine adducts (see the supplement).

Irradiation in the presence of glycine ethyl ester

Anthracene **1** (100 mg, 0.37 mmol) was dissolved in CH_3CN (150 mL) and mixed with a solution of glycine ethyl ester (2.0 g, 14 mmol) in aqueous phosphate buffer (50 mL, pH = 7, $c = 0.05$ M), or H_2O (50 mL) to which pH was adjusted to 9.5 with NaOH prior to the mixing with the solution of **1**. The resulting $\text{CH}_3\text{CN}-\text{H}_2\text{O}$ solutions were purged with Ar for 30 min and irradiated in a Rayonet reactor equipped with 11 lamps with the output at 350 nm during 2 h. After the irradiation, the solution was analyzed by ESI-MS. H_2O (100 mL) was added and the extraction with CH_2Cl_2 (3×75 mL) was carried out. The organic extracts were dried over anhydrous MgSO_4 , filtered and the solvent was removed on a rotary evaporator. The residue was chromatographed on a silica gel or alumina column using $\text{CH}_2\text{Cl}_2/\text{EtOAc}$ (0–100%) as eluent. Chromatographic separation furnished starting compound **1** (40–50 mg), and a mixture of unidentified rearrangement products.

Irradiation in the presence of serine or glutathione

Anthracene **1** (15 mg, 0.055 mmol) was dissolved in CH_3CN (10 mL) and mixed with a solution of (\pm) serine (120 mg, 1.14 mmol)

or L-glutathione (150 mg, 0.49 mmol) in H₂O (10 mL). The solutions were poured into cuvettes, sealed with a septum, purged with Ar for 30 min, and irradiated in a Luzchem reactor equipped with 8 lamps with the output at 350 nm over 1 h. After the irradiation the composition of the solutions was analyzed by ESI-MS.

Preparative irradiation in the presence of protected cysteine.

Anthracene **1** (70 mg, 0.26 mmol) was dissolved in CH₃CN (70 mL) and mixed with a solution of methyl ester of *N*-acetyl-L-cysteine (350 mg, 2 mmol in 80 mL CH₃CN) and H₂O (30 mL). The resulting solution was purged with Ar for 30 min and irradiated in a Rayonet reactor equipped with 11 lamps with the output at 350 nm during 2 h. During the irradiation the solution was continuously purged with Ar and cooled with a tap-H₂O finger-condenser. After the irradiation the solvent was removed on a rotary evaporator and the residue chromatographed on a column of silica using CH₂Cl₂/EtOAc (10%) as eluent, and rechromatographed on a TLC using the same eluent. The chromatography furnished 20 mg (28%) of the starting material, 40 mg (34%) of the cysteine adduct and 10 mg of a mixture containing H₂O-adduct **3** and some unidentified products.

(R)-methyl 2-acetamido-3-[9-(2-hydroxyphenyl)-9,10-dihydroanthracen-9-ylthio]propanoate (**8**)

40 mg (34%); yellowish oil, ¹H NMR (300 MHz, CDCl₃) δ/ppm 8.05 (dd, 1H, *J* = 1.5 Hz, *J* = 7.8 Hz), 7.34-7.42 (m, 3H), 7.26-7.34 (m, 2H), 7.08-7.18 (m, 3H), 6.78-7.85 (m, 3H), 5.62 (d, 1H, *J* = 7.7 Hz), 4.28-4.44 (m, 3H), 4.14 (br s, 1H), 3.60 (s, 3H, OCH₃), 2.79 (dd, 1H, *J* = 4.5 Hz, *J* = 13.0 Hz), 2.66 (dd, 1H, *J* = 6.1 Hz, *J* = 13.0 Hz), 1.85 (s, 3H, COCH₃); ¹³C NMR (75 MHz, CDCl₃) δ/ppm 170.7 (s), 169.4 (s), 153.5 (s), 137.2 (s), 137.1 (s), 136.0 (s), 135.9 (s), 131.2 (d), 129.8 (d), 128.2 (d), 128.1 (d, 2C), 128.0 (d), 127.6 (s), 127.2 (s), 127.1 (s), 126.7 (d), 126.6 (d), 120.4 (d), 117.9 (d), 65.4 (s), 52.4 (d), 51.1 (q), 35.4 (t), 34.0 (t), 22.9 (q), 14.1 (q); ESI-MS (*m/z*) 470 (M⁺+Na); HRMS (MALDI) calculated for C₂₀H₁₅O 271.1117, found 271.1118; calculated for C₆H₁₀NO₃S 176.0376 found 176.0330.

Spectroscopic measurements. The stock solutions of anthracene **1** in DMSO (*c* = 0.01 M), ct-DNA (*c* = 0.01 M) in aqueous sodium cacodylate buffer (pH = 7, *c* = 0.05 M), and bovine serum albumin (BSA, *c* = 0.15 mM) in potassium phosphate buffer (pH = 7, *c* = 1 mM) were prepared. The solutions were further diluted with the cacodylate or the phosphate buffer, respectively. UV-VIS spectra were recorded on a Varian Cary 100 Bio spectrophotometer at rt. Fluorescence spectra were obtained on a Varian Cary Eclipse fluorometer and the concentrations were adjusted to absorbances of less than 0.1 at the excitation wavelengths of 270, 280, 290, 295, 346 or 365 nm. The solutions were not purged.

In the study of interaction with DNA, UV-vis spectra of **1** (1 × 10⁻⁵ M) in the cacodylate buffer were recorded in the presence of ct-DNA (1 × 10⁻⁴ M) at different temperatures (25-75 °C). Thermal melting curves for DNA and complexes with studied compounds (*c*(DNA) = 2 × 10⁻⁵ M; compound/DNA ratio *r*=0.2) were determined as previously described⁴³ by following the absorption change at 270 nm as a function of temperature. The *T_m* values are

the inflection points of the transition curves determined from the maximum of the first derivative, and were checked graphically by the tangent method. The Δ*T_m* corresponds for the difference of the *T_m* of the free nucleic acid *T_m* of the complex. CD spectra were recorded on a Jasco J-815 spectrometer in the cacodylate buffer with ct-DNA (*c* = 4 × 10⁻⁵ M) and **1** (*c* = 1 × 10⁻⁵ M) in 3 mL cuvettes at rt.

In the study of interaction with bovine serum albumin (BSA), UV-vis, fluorescence and CD spectra of **1** (1 × 10⁻⁵ M) in the phosphate buffer were recorded in the presence of BSA (3 × 10⁻⁶ M) in 3 mL cuvettes at 25 °C.

Antiproliferative investigation. The experiments were carried out on three human carcinoma cell lines HCT 116 (colon), MCF-7 (breast) and H 460 (lung). Cells were cultured as monolayers and maintained in Dulbecco's modified Eagle medium (DMEM) supplemented with 10% fetal bovine serum (FBS), 2mM L-glutamine, 100 U/mL penicillin, and 100 μg/mL streptomycin in a humidified atmosphere with 5% CO₂ at 37 °C.

The cells were inoculated in parallel on three 96-well microtiter plates on day 0, at 1 × 10⁴ (HCT 116 and H 460) or 3 × 10⁴ cells/ml (MCF-7), depending on the doubling times of a specific cell line.

Test agents were added in ten-fold dilutions (10⁻⁸ to 10⁻⁴ M) the next day and incubated for further 72 h. Working dilutions were freshly prepared on the day of testing. One of the plates was left in the dark, the other one was irradiated in a reactor (6 lamps 350 nm, 1 min) 4, 24, and 48 hours after the addition of the compounds (3 × 1 min), while the third one was irradiated in a reactor (6 lamps 350 nm, 5 min) 4, 24, and 48 hours after the addition of the compounds (3 × 5 min). After 72 h of incubation the cell growth rate was evaluated by performing the modified MTT assay⁴⁴⁻⁴⁶ (for the irradiated and non-irradiated cells) which detects dehydrogenase activity in viable cells. The absorbance (*A*) was measured on a microplate reader at 570 nm. The absorbance is directly proportional to the number of living, metabolically active cells. The percentage of growth (PG) of the cell lines was calculated according to one or the other of the following two expressions:

If (mean *A_{test}* - mean *A_{lzero}*) ≥ 0, then PG = 100 × (mean *A_{test}* - mean *A_{lzero}*) / (mean *A_{ctrl}* - mean *A_{lzero}*).

If (mean *A_{test}* - mean *A_{lzero}*) < 0, then: PG = 100 × (mean *A_{test}* - mean *A_{lzero}*) / *A_{lzero}*, where the mean *A_{lzero}* is the average of absorbance measurements before exposure of cells to the test compound, the mean *A_{test}* is the average of absorbance measurements after the desired period of time and the mean *A_{ctrl}* is the average of absorbance measurements after the desired period of time with no exposure of cells to the test compound. In the experiments where the cells were irradiated, *A_{ctrl}* represents irradiated control cells. After irradiation at 350 nm (3 × 5 min) up to 25% growth inhibition compared to *A_{ctrl}* without irradiation was observed. The results are expressed as IC₅₀, which is the concentration necessary for 50% of inhibition. The IC₅₀ values are calculated from concentration-response curves using linear regression analysis by fitting the test concentrations that give PG values above and below the reference value (*i.e.* 50%). Each test was performed in quadruplicate in at least two individual experiments.

Cell Cycle Analysis. Colon cancer cells HCT 116 were seeded into 6-well plates (2×10^5 per well). After 24 hours the tested compounds were added at various concentrations (as shown in the Results section). One of the plates was left in the dark, while the other was irradiated in a reactor (6 lamps 350 nm, 5 min) 4, 24 and 72 hours after the addition of the compounds. After the desired length of time (48 or 72 hours) the attached cells were trypsinized, combined with floating cells, washed with phosphate buffer saline (PBS), fixed with 70% ethanol and stored at -20°C . Immediately before the analysis, the cells were washed with PBS and stained with $50\mu\text{g/ml}$ of propidium iodide (PI) with the addition of $0.2\mu\text{g}/\mu\text{l}$ of RNase A. The stained cells were then analyzed with Becton Dickinson FACScalibur (Becton Dickinson) flow cytometer (20 000 counts were measured). The percentage of the cells in each cell cycle phase was determined using the ModFit LT™ software (Verity Software House) based on the DNA histograms. The tests were performed in duplicates and repeated at least twice.

Alkaline agarose gel assay. DMSO-stock solutions of compounds **1** and **7** (psoralen) were diluted in H_2O and used for reactions. Plasmid pCI ($0.8\mu\text{g}/\text{sample}$) was mixed with compound dilutions. Reaction mixtures were irradiated for 5 min at 350 nm in a Luzchem reactor. Irradiated solutions were added to alkaline agarose gel loading buffer [50mM NaOH, 1mM ethylenediaminetetraacetic acid (EDTA), 3% Ficoll, and 0.02% bromophenol blue] and loaded on a 1% alkaline agarose gel containing 50mM NaOH and 1mM EDTA. Gels were run in 50mM NaOH and 1mM EDTA at 25 V constant voltage in horizontal electrophoresis system (BIO-RAD, USA), stained with ethidium bromide ($0.5\mu\text{g}/\text{mL}$) for 10 min, after neutralization of the gel with 30mM Tris-HCl (pH 7.5). Resulting products were visualized and documented with UV light at 254 nm (Uvitec, Cambridge).

Inhibition of *in vitro* translation. The compound **1** was tested for the inhibitory effect on translation of Enhanced Green Fluorescence Protein (EGFP) using an *E. coli* derived cell-free protein synthesis system (S30 T7 High-Yield Protein Expression System, Promega, USA) according to manufacturer's protocol as described previously.⁴⁹ This system can produce high levels of recombinant proteins if supplemented with an appropriate expression plasmid, T7 RNA polymerase for transcription, and other necessary components for translation such as amino acids. Briefly, 500ng of pEGFP-C1 plasmid (Clontech, USA) was incubated with water or the tested compound (at $100\mu\text{M}$ concentration) and was irradiated at 350 nm for 5 min, or kept in dark at 24°C . Along with the tested compounds psoralen, a known intercalative agent, which, upon exposure to ultraviolet (UVA) radiation, can form covalent interstrand cross-links (ICL) ($5\mu\text{M}$ concentration, Sigma) was also used for comparison reasons. Positive controls contained the plasmid in sterile water and negative controls did not contain DNA template (plasmid), respectively. After that, the protein expression system was deployed according to the manufacturer recommendations and protocol with slight modification of incubation temperature to ensure optimal EGFP folding. Instead of 37°C the mixture was incubated at 32°C for 3 hours. Afterwards, GFP fluorescence was

recorded on microtiter plates using Fluoroskan Ascent Microplate Fluorometer (excitation at 485 nm, emission at 538 nm; ThermoScientific). The mean fluorescence of negative controls was subtracted from the means of tested and control samples and percentages from control were calculated. Minimum three experiments were performed, and statistical difference was calculated using Microsoft Excel t-test.

Results and discussion

Chemistry

Synthesis of compounds **1** and **4–6** and their photochemical reactivity have been described.^{36–38} They were obtained by Suzuki coupling reactions, followed by a cleavage of the methyl ethers by BBr_3 .

Irradiation of **1** in $\text{CH}_3\text{CN-H}_2\text{O}$ gives H_2O -adduct **3** ($\text{Nu} = \text{OH}$) that can be isolated by chromatography.³⁶ Further, we investigated reactivity of the photogenerated QM **2** with nucleobases. Irradiations of **1** in $\text{CH}_3\text{CN-H}_2\text{O}$ were performed in the presence of adenine (A), guanine (G) and cytosine (C). In a typical photochemical experiment a mixture of **1** (0.1mmol), and a nucleobase ($\approx 0.1\text{mmol}$), were dissolved in a mixture of $\text{CH}_3\text{CN-H}_2\text{O}$ containing phosphate buffer ($c = 0.1\text{M}$, $\text{pH} = 6.8$) and irradiated 1 h at 350 nm. The course of the reaction was followed by HPLC. Irradiations gave H_2O -adduct **3** and additional products that were not detected when the irradiation was performed without the nucleobases. Therefore, we tentatively assigned the structures to adducts of the nucleobases to QM **2** (Fig 2). However, no adduct could be isolated. The photoproducts decomposed after standing in $\text{CH}_3\text{CN-H}_2\text{O}$ solution at rt for several hours. On attempts to purify the photoproducts by preparative TLC on silica, they decompose to **1** and nucleobase, and gave some additional more complex rearrangement products that were not analyzed. Therefore, the site of the attack to the nucleobases could not be determined. The drawings in Fig. 2 correspond to the anticipated sites of attack according to the published reactivity of QMs with deoxynucleosides.^{12–15}

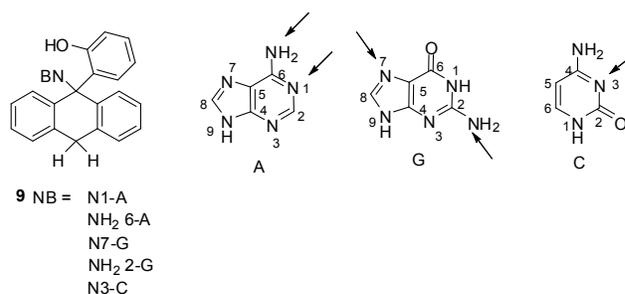


Fig. 2 Tentative structures **9**, adducts of the nucleobases to QM **2**.

Reactivity of the photogenerated QM **2** was also investigated with amino acids. Irradiation of **1** in $\text{CH}_3\text{CN-H}_2\text{O}$ was performed in the presence of C-protected glycine, serine and *N*- and C-protected cysteine. Irradiation in the presence of glycine ethyl ester was performed at pH 7 and pH 9.5. In the neutral solution no glycine adducts were detected. Chromatographic separation on silica recovered the starting anthracene (40%) and gave H_2O -adduct **3** in addition to a mixture of unidentified products (40%). On the contrary, after the irradiation of the solution at pH 9.5,

Table 1. IC₅₀ values (in μM)^a induced with compounds **1** and **4–7** with and without irradiation at 350 nm.

Comp.	HCT116			MCF-7			H 460		
	Not irradi.	3×1 min	3×5 min	Not irradi.	3×1 min	3×5 min	Not irradi.	3×1 min	3×5 min
1	21±4	12±2	2±0.2	20±4	8±5	2±0.3	20±4	13±1	2±0.2
4	24±2	23±4	19±1	24±2	22±1	17±2	18±1	19±1	15±2
5	18±1	18±2	10±2	17±2	14±3	6±2	18±0.4	18±2	12±1
6	21±1	19±0.1	11±0.02	17±3	15±3	5±0.2	17±3	16±1	9±3
7	39±24	0.1±0.1	<0.01	3±1	0.2±0.1	0.02	≥100	0.03±0.01	<0.01

^a Concentration that causes 50% inhibition of tumor cell growth.

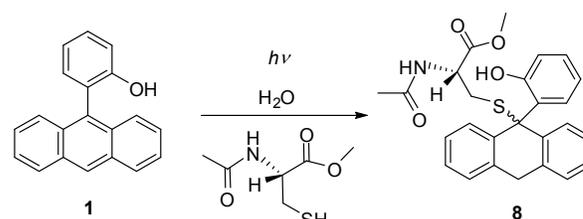
5 corresponding to the pK_a of glycine, adduct **10** (Fig. 3) could be detected by NMR and ESI-MS-MS (see the SI). However, attempts to isolate **10** on silica gel or alumina gave starting compound **1** (50%) and a mixture of H₂O-addcut and some rearranged products (30%). The experiments indicate that amino groups in amino acids or peptides can in principle react with QM **2**. However, the addition is feasible only at high pH values when the amino groups are not protonated.

Methanol is a good nucleophile that can react with QM **2** giving adducts.³⁶ Consequently, it is plausible that QM **2** can react with 15 other alcohols or amino acids and peptides bearing free OH group such as serine. Indeed, irradiation of **1** in CH₃CN-H₂O in the presence of nonprotected serine in neutral solution gave an adduct that was detected by ESI-MS (see the SI). MS-MS analysis of the isolated molecular ion (*m/z* 376) gave major peaks of *m/z* 20 271 and 106, corresponding to the protonated **1** and serine, respectively, in accord with a high aptitude of molecule to decompose to the starting materials. We could not isolate the product, but its structure probably corresponds to ether **11**. At pH 7 the hydroxyl group in serine is much better nucleophile than the 25 protonated amino group.

Irradiation of **1** in CH₃CN-H₂O in the presence of *N*- and *C*-protected cysteine, followed by chromatographic separation gave cysteine adduct **8** (34%, Scheme 2), together with the recovered starting material (28%), and some unidentified products. 30 Although adduct **8** can be present in two diastereomeric forms, only one isomer was isolated. The structure assignment of **8** was straightforward from ¹H and ¹³C NMR spectra. However, stereochemistry at the stereogenic center formed in the photochemical reaction could not be assigned. In the aliphatic part of the ¹H NMR spectrum a characteristic signal corresponding to the dihydroanthracene H-atoms were observed in a multiplet at δ 4.2–4.4 ppm, together with the signal of H-atom at the cysteine chiral center. In the aromatic part of the ¹³C NMR spectrum twelve doublets were detected together, with the 40 singlets of two carbonyls and four aromatic C-atoms. The aliphatic part of the ¹³C NMR spectrum revealed two triplets, one doublet and two quartets, all in accordance with the assigned structure.

To demonstrate the ability of QM to react with oligopeptides irradiation of **1** in CH₃CN-H₂O in the presence of glutathione was conducted. After the irradiation the solution was analyzed by ESI-MS (see the SI) wherein we detected presence of an adduct whose structure was tentatively assigned to **12**. The MS-MS analysis of the isolated molecular ion (*m/z* 578) gave major peaks 50 of *m/z* 271, 308, and 449 corresponding to the protonated **1**, glutathione and a fragment after the loss of 2-amino-5-

oxopentanoic acid, respectively. The fragmentation of the molecule clearly indicated that glutathione sulfur is attached to the dihydroanthracene moiety, in accordance with structure **12**.



Scheme 2 Photochemical formation of cysteine adduct **8**.

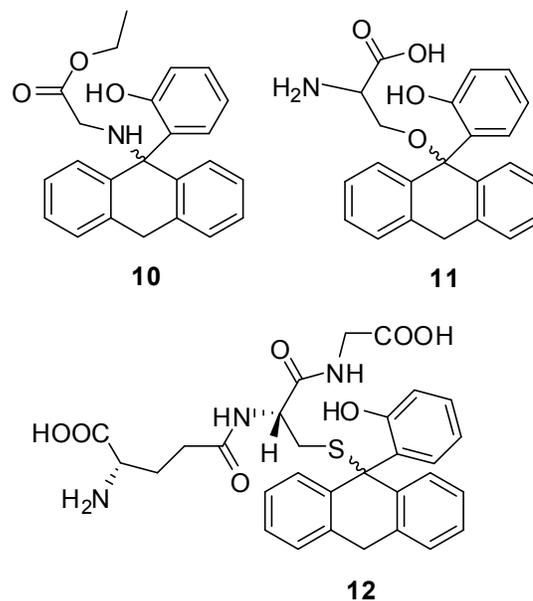


Fig. 3 Tentative structures **10**, **11** and **12**, adducts of the amino acids and glutathione to QM **2**.

Photocytotoxicity study

Antiproliferative effect of the photogenerated QMs on three cancer cell lines HCT 116 (colon carcinoma), MCF-7 (breast carcinoma), and H 460 (lung carcinoma), was investigated with 4 anthracene derivatives. Psoralen derivative **7** was used as a positive reference compound that is known to exhibit antiproliferative activity on photoactivation.⁴³ Cells were incubated with the compounds and kept in dark, or irradiated at 70 350 nm 3×1min or 3×5min. The activities expressed as IC₅₀

(concentration that causes 50% inhibition of the cell growth) are compiled in Table 1. Without the irradiation, all compounds exert mild antiproliferative activity in micromolar concentration range, without a specific effect with respect to the cell type. Irradiation induced enhancement of the activity for derivatives **1** and **6**, and the effect could clearly be correlated to the irradiation time, probably due to the formation of the corresponding QMs. Up to 10-fold and 3-fold increase of the activity can be achieved for **1** and **6**, respectively. Although the enhancement for psoralene derivative **7** is more pronounced (more than 100-fold), **1** was chosen as a candidate compound for further studies for the elucidation of the antiproliferation mechanism.

Non-covalent binding to DNA

Anthracene derivatives are generally good candidates for non-covalent binding to DNA, either by intercalation or binding to a minor groove. Non-covalent binding of **1** to calf thymus DNA (ct DNA) was investigated by UV-vis and CD spectroscopy. Absorption spectrum of **1** ($c = 1 \times 10^{-5}$ M) in aqueous solution in the presence of cacodylate buffer (50 mM, pH = 7) exhibits typical structured anthracene absorption at 320-400 nm. Heating of the solution to 70 °C and cooling to the rt induces disappearance of the anthracene absorption band with a concomitant increase of the spectral baseline, suggesting aggregation of the molecules and formation of nanoparticles that cause light scattering. Addition of an aqueous solution of ct DNA to the solution of **1** resulted in a hypochromic change of the absorption band at 320-400 nm, suggesting binding of **1** to the DNA. However, increase of the temperature to 70 °C induced absorbance increase, suggesting dissociation of the complex of DNA and **1** (Fig 4).

Compounds that bind to DNA cause stabilization of the duplex and increase of the DNA melting temperature (ΔT_m). Although the absorption spectra (Fig. 4) indicated DNA/**1** complex formation, **1** did not stabilize ds-DNA against thermal denaturation. Possible explanation is related to the observed aggregation of **1**. Nevertheless, the irradiation of DNA in the presence of **1** at rt is anticipated to cause alkylation of DNA, whereby resulting DNA adducts could show higher thermal stability than non-treated DNA. However, determination of T_m for DNA in the absence of **1**, and in the presence of **1** after 5 min of irradiation at 350 nm, gave the same $T_m = 79.0 \pm 0.3$ °C. This lack of measurable stabilization could be due to low efficiency of adduct formation or because of instability of nucleobase-adducts **9** (Fig. 2). Thus, thermal denaturation experiment could not give the unambiguous conclusion with respect to the ability of QM **2** to alkylate DNA.

CD spectroscopy as a highly sensitive method to assess conformational changes in the secondary structure of polynucleotides was applied to investigate the mode of binding of **1** to ct-DNA. It was anticipated that binding of achiral **1** to the helically chiral DNA should give rise to induced CD spectrum of **1**, aside expected changes in CD spectrum of DNA (220 – 300 nm).⁵⁰ However, no change was observed between the CD spectrum of free ct DNA, and DNA in the presence of **1** ($c = 1 \times 10^{-5}$ M, ct-DNA, $c = 4 \times 10^{-5}$ M). The finding indicates that **1** does not form one dominant complex within well defined DNA binding site (e.g. intercalation or minor groove binding), but

rather aggregates nonspecifically along both grooves of polynucleotide double helix due to the hydrophobic effect and aromatic interactions between molecules of **1** (latter supported by hypochromic changes in the UV-vis spectra, Fig. 4). Nevertheless, even such nonspecific aggregation brings **1** close to the number of potential alkylation sites along DNA. Thus, to check if the photochemical reaction could lead to the changes in CD spectra due to alkylation with QM **2**, the solution **1** in the presence of ct-DNA was irradiated (5 min, 8 lamps at 350 nm). However, no change in the CD spectrum could be observed, either due to lack of measurable reactivity of QM **2** with DNA, or because the photochemical product does not induce significant changes of the DNA helical chirality.

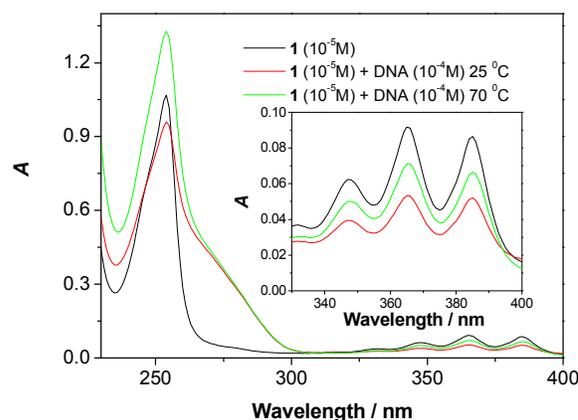


Fig. 4 Absorption spectrum of **1** ($c = 1 \times 10^{-5}$ M) in aqueous cacodylate buffer (50 mM, pH = 7) in the presence of ct-DNA ($c = 1 \times 10^{-4}$ M) at different temperatures; Inset: enlarged part of the spectra between 330 and 400 nm.

Non-covalent binding to proteins and photoreaction in protein

Anthracene derivative **1** is nonpolar, so it is plausible that it could bind to hydrophobic pockets of proteins. It has been demonstrated that anthracenes bound to bovine serum albumin (BSA) undergo photochemical reactions.⁵¹ Therefore, non-covalent binding of **1** to BSA was investigated by UV-vis, fluorescence and CD spectroscopy. After addition of an aqueous BSA solution to the solution of **1** ($c = 1 \times 10^{-5}$ M) in phosphate buffer (pH = 7, $c = 1$ mM), a hyperchromic change of the absorption bands at 250 and 320-400 nm was observed (Fig. 5 top). The hyperchromic change suggests that **1** binds to BSA upon which it deaggregates. After addition of BSA ($c = 3 \times 10^{-6}$ M) to the solution of **1** ($c = 1 \times 10^{-5}$ M) in phosphate buffer (pH = 7, $c = 1$ mM), a positive signal in the CD spectrum was observed at 250-260 nm, corresponding to the $S_0 \rightarrow S_2$ anthracene absorption (Fig 5. bottom). In the region 350-400 nm the CD signal was not observed due to much lower absorptivity. The induced CD spectrum strongly indicates non-covalent binding of achiral **1** to the protein. Similar induced CD spectra of anthracene derivatives in BSA with a positive signal at 250-260 nm have been reported.⁵¹

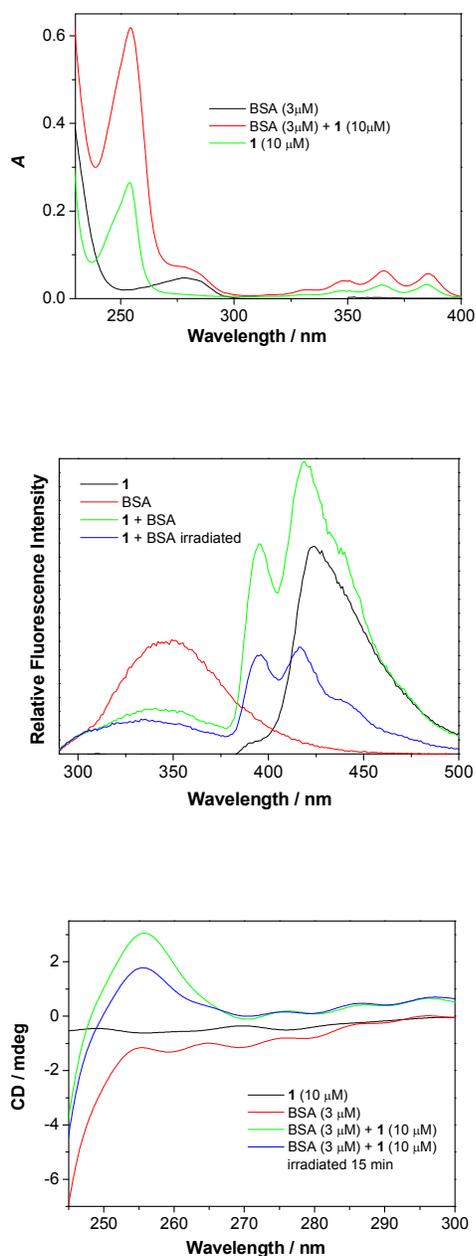


Fig. 5 Absorption (top), fluorescence (middle, $\lambda_{\text{ex}} = 280 \text{ nm}$), and CD spectra (bottom) of **1** ($c = 1 \times 10^{-5} \text{ M}$), BSA ($c = 3 \times 10^{-6} \text{ M}$) and their mixture in aqueous phosphate buffer (1 mM, pH = 7) at 25 °C before and after the irradiation (Luzchem, $8 \times 350 \text{ nm}$, 15 min).

Fluorescence excitation and emission spectra of **1** ($c = 1 \times 10^{-5} \text{ M}$) in aqueous phosphate or cacodylate buffer indicate aggregation of molecules (see the supporting info.). In contrast to the vibronically well resolved emission spectrum of **1** in CH_3CN ,³⁶ in aqueous buffered solution the spectrum (regardless of the excitation wavelength, 270, 280, 345 or 365 nm) has a shoulder at 395 nm, most probably corresponding to the non-aggregated molecules, and much stronger band with a maximum at 430 nm, corresponding to the emission from the aggregated molecules

(Fig. 5 middle and supporting info.). In a fluorescence spectrum of **1** and BSA mixture, fluorescence from both species can be detected. However, fluorescence from the protein is quenched compared to the neat BSA solution of the same concentration, whereas the anthracene emission is stronger and more resolved. The observation can be explained by non-covalent binding of **1** to BSA, whereupon the anthracene molecules deaggregate. Moreover, formation of the BSA/**1** complex leads to the quenching of fluorescence from tryptophanes in protein by energy transfer to **1**. Consequently, fluorescence from **1** is stronger in the complex than in the aqueous solution not containing BSA.

Irradiation of **1** (Luzchem, 350 nm, 15 min) in the presence of BSA resulted in a bleaching of the typical anthracene absorption band (see the SI) and the decrease of fluorescence, in agreement with the photoreaction of **1** giving anthracene adducts. Furthermore, induced CD spectrum in the region 250-260 nm was weaker after the irradiation, indicating that anthracene bound in the protein underwent photoreaction. The photoreaction inside the protein could be due to the photoreaction with BSA or due to photohydration giving H_2O -adduct **3**. Attempt to detect MS of the BSA-anthracene photo-adducts by MALDI failed. However, no unambiguous conclusion could be made due to too high molecular weight of BSA and a small m/z difference between the potential adduct and the BSA M^+ for which the MALDI analysis is not sensitive enough.

DNA alkylation experiment

To check the DNA-alkylation ability, we performed irradiations of supercoiled plasmid DNA in the presence of compound **1** and **7** (psoralen), followed by alkaline DNA gel electrophoresis. Clear differences in the various plasmid forms' migration performances can be seen after the irradiation of plasmid DNA in the presence of **7**, due to its well-known ability to intercalate and, on exposure to ultraviolet radiation, to form covalent interstrand cross-links (ICL) (Fig. 6).

Interstrand cross-linking (XL) activities of psoralen were evident as X-bands of circular (X-C), open circular (X-OC) and linear X-L forms). However, no X-bands and no difference in migratory ability compared to control DNA were observed after irradiation of **1**. Such result is not surprising since DNA alkylation products of **1** were demonstrated to decompose very easily, and besides, alkaline agarose gel electrophoresis allows distinguishing between cross-linked and uncross-linked species, but is probably not sensitive enough to visualize mono-alkylation,⁵² which we expected to occur after the irradiation of **1**.

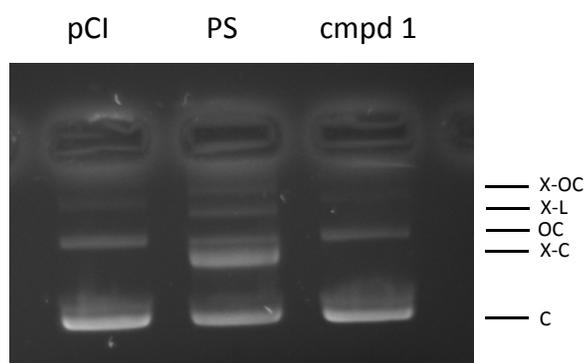


Fig. 6 Alkaline gel electrophoresis of DNA in the presence of psoralen and **1** (lane 1: irradiated pCI 0.8 μ g, lane 2: irradiated pCI + psoralen (PS), 5 μ M, lane 3: irradiated pCI + **1** 100 μ M) after irradiation (5 min at 350 nm). Non-reacted circular and open-circular plasmid forms (C and OC) and their cross-linked species (X-C and X-OC) along with cross-linked linear (X-L) are indicated at the right side of the gel image.

10 Cell cycle perturbation

To shed more light on the antiproliferative mechanism(s) of **1**, we assessed its influence on the cell cycle of HCT 116 cells at the IC₅₀ concentration obtained for both non-irradiated (20 μ M; data not shown), irradiated (2 μ M), along with 0.2 μ M concentration after 48 h and 72 h of incubation with or without 2 \times 5 min, or 3 \times 5 min of irradiation, respectively. Interestingly, without irradiation compound **1** negligibly influences the cell cycle, mostly causing a slight delay in the S phase-progression (DNA synthesis), and a slight induction of subG1 cells (Fig. 7b). These results do not point to DNA intercalative mode of action (usually strong G2/M arrest) and are in accordance with the previously described study of non-covalent DNA binding. The irradiation of cells for 2 \times 5 min incubated with 2 μ M concentration of **1**, caused a strong reduction in the number of cells in the S phase and induced both G1 and G2/M phase arrest, and significantly increased the number of cells in subG1, pointing to the activation of apoptosis (Fig. 7a). The additional 5 min of irradiation (72 h of incubation) led to massive activation of apoptosis (Fig. 7b). While these results clearly demonstrate a strong influence on the cell cycle and induction of cell death, the qualitative outcome do not undoubtedly point to DNA damage caused by DNA alkylation as a trigger. Namely, it has been previously demonstrated in the same tumor cell model (HCT 116 cells) that different alkylating agents, which induce either monoalkylation of DNA or DNA cross-links induced strong G2/M arrest.⁵³ Similar observations were also found in other tumor cell models.⁵⁴

Influence of compounds on protein synthesis

It was previously demonstrated that the alkylation of DNA suppressed transcription and translation in cells and in *in vitro* translation systems.^{49,55,56} If tumor cells were transfected with a prealkylated plasmid coding for a reporter protein GFP with *N*-methylquinolinium quinone methide, a significant suppression of GFP protein expression was detected. This inhibition was correlated with the amount of DNA adducts formed, i.e. the

concentration of alkylating agent.⁵⁴ Moreover, if a prealkylated DNA was also used as a template for coupled transcription and translation, a significant inhibition of reporter protein expression occurs.^{49,53} Therefore, to further examine whether the enhancement of tumor cells' growth inhibition is a consequence of DNA alkylation/damage by photogenerated QMs, produced upon irradiation of **1**, we assessed the expression of Enhanced Green Fluorescence protein (EGFP) in a cell free environment using a high-capacity T7 promoter driven *in vitro* translation system. Samples of DNA encoding the EGFP gene were incubated with compound **1** (100 μ M) and psoralen and subsequently irradiated at 350 nm for 5 min or kept in dark. Afterwards, the samples were subjected to coupled transcription and translation to find a correlation between the effect of DNA damage and the amount of transcribed and subsequently translated gene product. As expected, the irradiation of psoralen almost completely inhibited the EGFP expression. Similar results were obtained previously with the cross-linking agent cisplatin.⁴⁹ The irradiation of compound **1** led to a certain inhibition of EGFP expression, but compared to control, irradiated DNA, this inhibition was not statistically significant, pointing to its lower alkylating potential (Figure 7). However, in our previous study we tested the ability of another alkylating agent, chlorambucyl to inhibit protein translation in the same experimental system, and observed significantly lower inhibitory potency (55% of inhibition) of protein translation compared to cisplatin, or psoralen (94% of inhibition).⁴⁹ Therefore, again we could observe certain inhibition, which correlates with lower DNA alkylation ability of **1**.

Conclusions

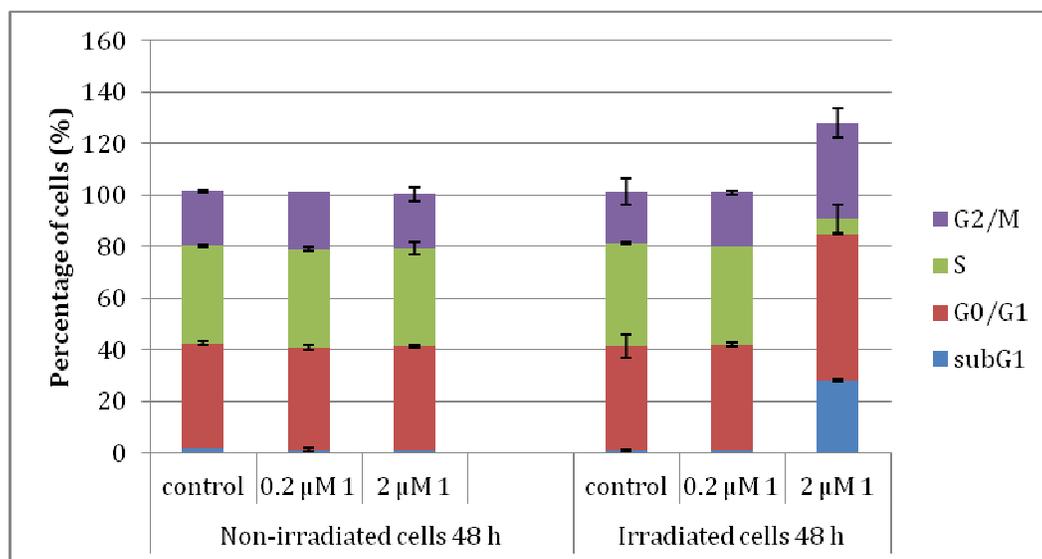
All investigated anthracenes moderately inhibit the proliferation of tumor cells. However, the irradiation (350 nm) of tumor cells incubated only with **1** considerably enhances the antiproliferative effect, by a drastic reduction of the percentage of cells in S phase, and an associated accumulation of cells in G1 and G2/M phase. This growth inhibition effect was dependent on the irradiance, whereby longer irradiation time caused massive tumor cells death. The enhancement of the antiproliferative effect was rationalized as being due to the formation of QM **2** from **1** by excited state proton transfer, which then acts as an electrophilic species capable of reacting with biological molecules. Even though the exact mechanism of the enhanced antiproliferative activity was not determined, we showed that **1** binds to DNA by nonspecific manner causing negligible photoinduced alkylation, at variance to evident impediment of protein synthesis. Anthracene **1** binds also to BSA, and probably to many other proteins. Photoreaction in the presence of proteins can in principle lead to alkylation, and most likely it is the major reason of photoinduced damages of the cellular proteins. This indication to major role of protein-impairment is an important finding with impact to the further development of anticancer agents based on QMs since to now all QM agents were sought to have DNA as the only target.

Cite this: DOI: 10.1039/c0xx00000x

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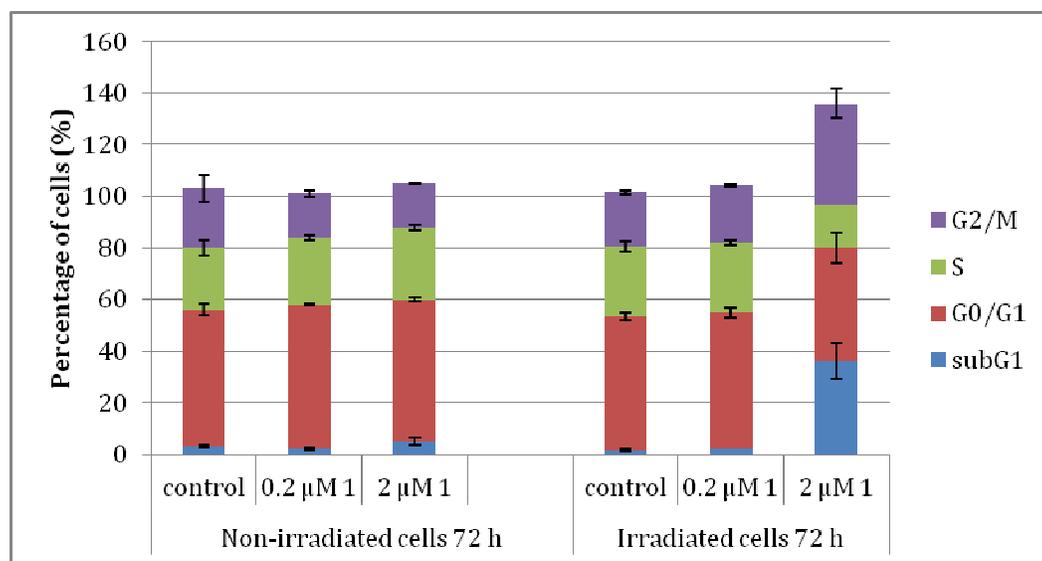


Fig. 7 The effects of compound **1** at 0.2 and 2 μM concentrations on the cell cycle distribution of HCT 116 cells after 24 and 48 h treatments (see text for details). The histograms represent the percentages of cells ± sd in the respective cell cycle phase (G1, S and G2/M), along with the percentage of cells in the subG1 (dead/apoptotic cells) obtained by flow cytometry.

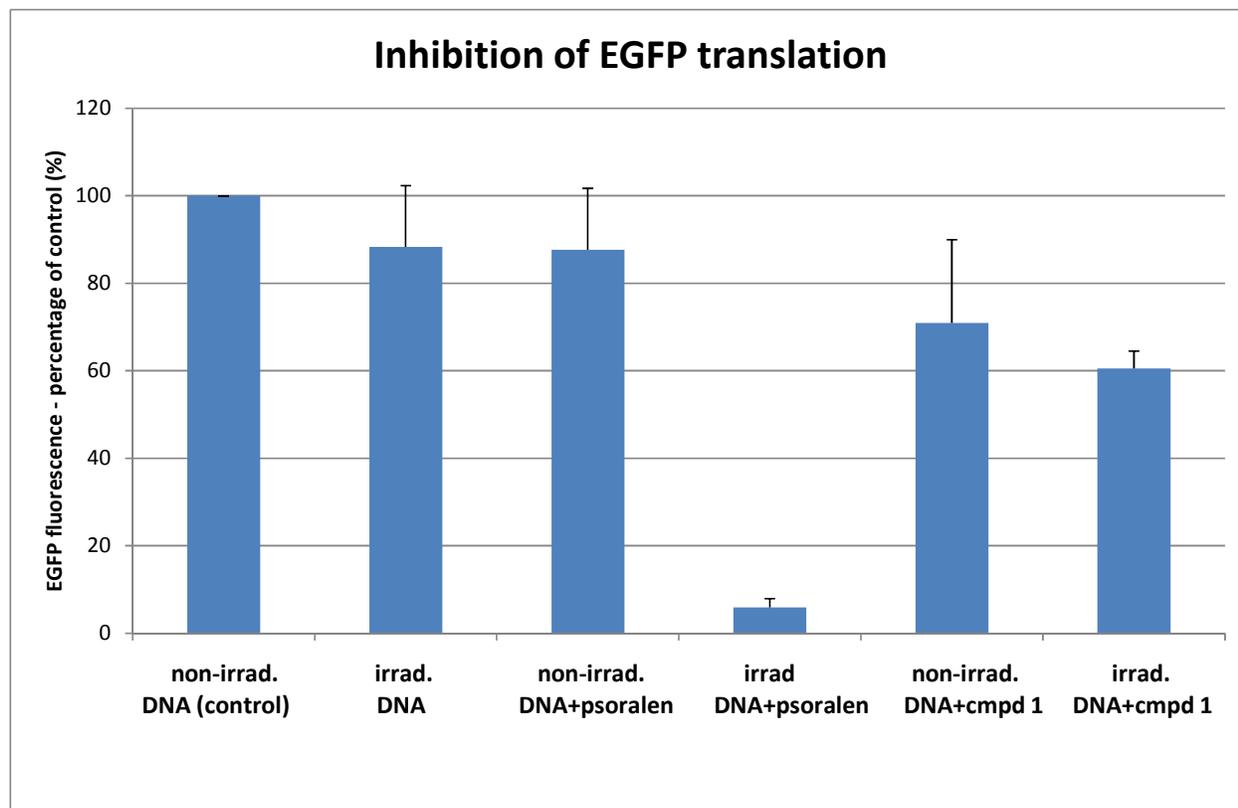


Fig. 6 Inhibition of EGFP protein translation. Plasmid containing EGFP gene under T7 promoter was incubated with psoralen (5 μ M) and with compound **1** (100 μ M) and was irradiated at 350 nm for 5 min (irradiated DNA), or kept in dark (non-irradiated DNA) at 24 $^{\circ}$ C. Non-irradiated DNA (positive control) and irradiated plasmid DNA without compounds were used as controls. Treated or untreated plasmid was then incubated with the S30 T7 High-Yield Protein Expression System, which enabled the EGFP protein translation/synthesis (see Experimental for details). The graph represents the percentages of EGFP fluorescence, corresponding to the amount of EGFP protein obtained during the *in vitro* translation procedure. The bars represent average values (\pm SEM) of three separate experiments performed in duplicates.

10 Acknowledgement

These materials are based on work financed by the Croatian Foundation for Science (HRZZ grants no. 02.05/25 and I-2364-2014), the Ministry of Science Education and Sports of the Republic of Croatia (grants No. 098-0982464-2514, 098-15 0982914-2918), the Natural Sciences and Engineering Research Council (NSERC) of Canada and the University of Victoria. The work was also supported by the FP7-REGPOT-2012-2013-1 project, Grant Agreement Number 316289 – InnoMol. The authors thank dr. S. Kazazić for recording MALDI MS spectra and dr. M. Matković for the help with CD spectrometer.

Notes and references

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[†] Electronic Supplementary Information (ESI) available: [HPLC chromatograms, NMR, MS, UV-vis, fluorescence and CD spectra]. See DOI: 10.1039/b000000x/

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Cite this: DOI: 10.1039/c0xx00000x

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