

DNA damage and induction of apoptosis in pancreatic cancer cells by a new dinuclear bis(triazacyclonane) copper complex

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Abstract

The dinuclear copper(II) complex $[\text{Cu}_2\{\text{bcmp}(-\text{H})\}(\mu\text{-OH})](\text{NO}_3)_2 \cdot \text{H}_2\text{O}$ (**1**, bcmp = 2,6-bis(1,4,7-triazacyclonon-1-ylmethyl)-4-methylphenol) has been synthesized and characterized by electrospray ionization mass spectrometry, potentiometric titration and cyclic voltammetry. The X-ray structure of the analogous perchlorate salt $[\text{Cu}_2\{\text{bcmp}(-\text{H})\}(\mu\text{-OH})](\text{ClO}_4)_2 \cdot 2.5\text{H}_2\text{O}$ (**2**) was determined. Cytotoxicity studies showed very promising activity of **1** against various pancreatic tumor cell lines with IC_{50} values comparable or even lower than those of cisplatin. The Cu complex displayed low toxicity against a human non-tumor cell line (HEK 293) demonstrating selectivity for cancer cells. **1** converts supercoiled pUC19 plasmid DNA into the nicked form at micromolar concentrations in the absence of added reductants. A detailed kinetic study on the hydrolysis of the DNA model bis(2,4-dinitrophenyl) phosphate (BDNPP) has been performed. **1** hydrolyses BDNPP with a second order rate constant of 0.047 M s^{-1} at pH 8 and $40 \text{ }^\circ\text{C}$. Finally, single cell electrophoresis (comet assay) and fluorescence microscopy analysis showed that **1** interacts with cellular DNA and induces apoptotic cell death of Capan-1 pancreatic cancer cells. Western blotting analysis indicated that the Cu complex activates the p53 dependent pathway of apoptosis.

Keywords: copper, cytotoxicity, DNA damage, metallodrugs

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1. Introduction

Copper is an essential element for most aerobic organisms. Employed as a structural and catalytic cofactor it is involved in many biological pathways [1-3]. In the search for new anticancer metallodrugs that can overcome the limitations of routinely used Pt drugs such as severe side effects, toxicity and inherent and acquired resistance, there is growing interest in biometals like Cu, as they are biocompatible and less toxic than non-endogenous heavy metals. Furthermore, Cu is an interesting candidate for the treatment of cancers because of its bioavailability and the observation of increased Cu levels in cancer tissue [4].

The importance of Cu in anticancer therapy is demonstrated by the ever increasing group of cytotoxic Cu complexes described in the literature, with the number of papers reported in 2012 being eight times greater than in 2008 [5,6]. Several synthetic Cu complexes have been found to be active both *in vitro* and *in vivo* [4].

Cytotoxic Cu(II) complexes often show DNA cleavage activity [4,7,8], suggesting a connection between these two properties. The mechanism of Cu complex-induced DNA strand breaks often involves the formation of reactive oxygen species (ROS) through the Fenton reaction following the reduction of Cu(II) to Cu(I) by biological reducing agents such as ascorbic acid. Metal-catalysed hydrolysis of the phosphodiester backbone of DNA, however, is also an attractive strategy for damaging tumor cell DNA irreversibly, because the mechanism does not require the addition of external agents. Furthermore, hydrolytic metallonucleases may find additional applications in molecular biology as substitutes for restriction enzymes or footprinting agents or as biomimetic systems helping to elucidate the mechanism of action of natural nucleases [9]. While natural enzymes have an active site with one or more metal ions surrounded by amino acids that are essential for the catalytic processes, in artificial nucleases the protein matrix is replaced by simple organic ligands.

In this paper we report the synthesis and full characterization of a novel dicopper(II) complex with a binucleating ligand containing a phenol scaffold and two triaza crown binding sites. The metallonuclease properties of the Cu complex were investigated in isolated DNA and the effects of DNA cleavage on cancer cell growth were assessed in Capan-1 pancreatic cancer cells.

2. Experimental

2.1. Materials and instrumentation

$\text{Cu}(\text{NO}_3)_2 \cdot 2.5\text{H}_2\text{O}$ was purchased from TCI Europe and used as supplied. All the other chemicals and solvents were of analytical or spectroscopic grade, obtained from commercial sources and used without further purification. Deuterated solvents were obtained from Apollo Scientific. MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) and cisplatin were obtained from Sigma Chemical Co, St. Louis, USA. Bis(2,4-dinitrophenyl) phosphate (BDNPP) was prepared according to the literature [10]. 2,6-bis(1,4,7-triazacyclonon-1-ylmethyl)-4-methylphenol was synthesised by a modified literature procedure [11] as recently described by us [12].

The pD values of D_2O solutions were adjusted using concentrated DNO_3 and NaOD . UV/Vis measurements were carried out on a Varian Cary 50 scan spectrophotometer coupled to a Grant thermostatted water circulation bath. Elemental analyses (carbon, hydrogen and nitrogen) were performed with a PerkinElmer 2400 series II analyzer. Electrospray ionization (ESI) mass spectra were recorded with a Waters LCT Premier XE Spectrometer. Electrochemical measurements were performed in 50 mM phosphate buffer (pH 7) at room temperature using a CH Instruments 620 series potentiostat, coupled to an electrochemical cell containing an Ag/AgCl reference electrode (Bioanalytical Systems), a platinum foil counter electrode (Goodfellow) and a 3 mm glassy carbon working electrode (Goodfellow). Potentials are quoted versus the Ag/AgCl reference electrode.

2.2. Potentiometric titration

An aqueous solution of **1** (1 mM, 50 mL) was titrated with a standardized 0.1 M NaOH solution at 25 °C. The ionic strength was maintained at 0.1 M with KNO_3 . pH values were measured with a Jenway 3510 pH meter fitted with a Reflex Sensor Ltd. EC-1910-11 glass electrode. The program HYPERQUAD was used to calculate the deprotonation constant from the titration data [13].

2.3. UV kinetic measurements

The hydrolysis rates of BDNPP were measured by monitoring the increase in the visible absorbance at 400 nm due to the release of 2,4-dinitrophenolate. Rate constants were obtained by the initial rate method (< 5% conversion). Concentrations of 2,4-dinitrophenolate were calculated from the extinction coefficients ($12100 \text{ M}^{-1}\text{cm}^{-1}$ for 2,4-dinitrophenolate). Concentrations were corrected for the degree of ionization of 2,4-dinitrophenol at the respective pH value using pK_a (2,4-dinitrophenol) = 4.0 [14]. In a typical experiment 15 μL of a freshly prepared BDNPP stock solution (5 mM in DMSO) was added to a solution of the metal complex (1.5 mL, 0.25 - 5 mM) at 40 °C. The metal complex solutions were buffered with 50 mM PIPBS (pH 3.5-5), MES (pH 5-6.7), HEPES (pH 6.8-8.5) and CHES (pH 8.5-11.0). The ionic strength was maintained at 0.1 M with KNO_3 . Cleavage rates were corrected for the spontaneous hydrolysis of the substrate [10]. The kinetic pK_a value was determined by fitting the data to

$$k_{\text{obs}} = (k_{\text{H}_2\text{O}}[\text{H}^+] + k_{\text{OH}}K_a)/(K_a + [\text{H}^+]).$$

To determine the solvent deuterium isotope effect, analogous kinetic experiments were performed in 99.9% D_2O . The correction $\text{pD} = \text{pH}_{\text{measured}} + 0.4$ was applied to the pH meter readings [15,16]. Kinetic runs were run in duplicate to give a reproducibility of $\pm 15\%$.

2.4. X-ray analysis

Crystal data for compound **2** were collected at room temperature on an Agilent (formerly OxfordDiffraction) Xcalibur CCD diffractometer using graphite-monochromated Mo- $\text{K}\alpha$ radiation ($\lambda = 0.71069 \text{ \AA}$) [17]. The structure was solved by direct methods and subsequent Fourier syntheses and refined by full-matrix least squares on F^2 using SHELXS-97, SHELXL-97 [18-20] and Oscale [21]. The scattering factors were those given in the SHELXL program. Hydrogen atoms except those for the water molecules of crystallization were generated geometrically and refined as riding atoms with isotropic displacement factors equivalent to 1.2 times those of the atom to which they were attached (1.5 for methyl groups). The asymmetric unit of **2** that crystallized in the triclinic space group P1 contains two crystallographically independent complex cations. A MISSYM check [22, 23] was carried out to rule out any missed symmetry. The Flack's parameter [24] converged to $x = 0.65(2)$ indicating racemic twinning, which was taken into account during the final cycles of least

squares refinement using the twin law -1 0 0 0 -1 0 0 0 -1 2. Graphics were produced with ORTEX [25]. Crystallographic data and details of refinement are reported in Table S1.

2.5. DNA plasmid cleavage

pUC19 DNA cleavage activity of compound **1** was evaluated by agarose gel electrophoresis. Plasmid pUC19 DNA (1 µg/µL) was incubated with increasing concentration (range 5 - 50 µM) of **1** in Tris buffer (50 mM Tris, 18 mM NaCl, pH 8.2) at 37 °C for 3 h in the dark. The reaction was quenched by the addition of 3 µL of loading buffer (0.25% bromophenol blue and 30% glycerol) and samples were loaded onto a 1% agarose gel in TBE buffer (89 mM Tris, 89 mM boric acid, 2 mM EDTA, pH 8.2). The gels were subjected to electrophoresis for 4 h at 50 V, followed by staining with 0.5 µg/mL ethidium bromide overnight. Gel bands were visualized using a UV transilluminator and photographed using an Olympus digital camera.

2.6. Experiments with cultured human cells

Compound **1** was dissolved in DMSO just before the experiment and a calculated amount of drug solution was added to the cell growth medium to a final solvent concentration of 0.5%, which had no detectable effect on cell killing. Cisplatin was dissolved in 0.9% NaCl solution.

2.7. Cell cultures

Human breast (MCF-7), colon (HCT-15), and pancreatic (BxPC3, MIA Paca-2, AsPC-1, Capan-1, PANC-1) carcinoma cell lines along with melanoma (A375) cells and nontransformed embryonic kidney (HEK293) were obtained from American Type Culture Collection (ATCC, Rockville, MD). The human ovarian 2008 cancer cell line was kindly provided by Prof. G. Marverti (Dept. of Biomedical Science of Modena University, Italy). Human cervical carcinoma (A431) cells were kindly provided by Prof. F. Zunino (Division of Experimental Oncology B, Istituto Nazionale dei Tumori, Milan, Italy). Cell lines were maintained in the logarithmic phase at 37 °C in a 5% carbon dioxide atmosphere using the following culture media containing 10% fetal calf serum (Euroclone, Milan, Italy), antibiotics (50 units/mL penicillin and 50 µg/mL streptomycin) and 2 mM L-glutamine: (i) RPMI-1640

medium (Euroclone) for 2008, MCF-7, HCT-15, A431, AsPC-1 and BxPC3 cells; (ii) F-12 HAM'S (Sigma Chemical Co.) for A549 cells; (iii) DMEM (Sigma Chemical Co.) for MIA Paca-2, PANC-1, A375 and HEK293 cells; IMDM (Sigma Chemical Co.) for Capan-1 cells.

2.8. *MTT assay*

The growth inhibitory effect toward tumor cells was evaluated by means of the MTT assay [26]. Briefly, $(3 - 8) \times 10^3$ cells/well, dependent upon the growth characteristics of the cell line, were seeded in 96-well microplates in growth medium (100 μ L). After 24 h, the medium was removed and replaced with fresh media containing the compound to be studied at the appropriate concentration. Triplicate cultures were established for each treatment. After 72 h, each well was treated with 10 μ L of a 5 mg/mL MTT saline solution, and following 5 h of incubation, 100 μ L of a SDS solution in 0.01 M HCl were added. After an overnight incubation, cell growth inhibition was detected by measuring the absorbance of each well at 570 nm using a Bio-Rad 680 microplate reader. The mean absorbance for each drug dose was expressed as a percentage of the control untreated well absorbance and plotted vs drug concentration. IC_{50} values, the drug concentrations that reduce the mean absorbance at 570 nm to 50% of those in the untreated control wells, were calculated by a four parameter logistic (4-PL) model.

2.9. *Comet assay*

Single-cell gel electrophoresis for detection of DNA damage was performed using the comet assay reagent kit purchased from Trevigen (Trevigen Inc., Gaithersburg, MD, US) according to the manufacturer's instructions. Briefly, Capan-1 (2×10^5) cells were seeded in 25 cm^2 flasks in growth medium (6 mL). After 24 h, cells were incubated for different exposure times (6 and 12 h) with increasing concentrations of the tested compound. Subsequently, cells were washed twice with cold phosphate buffered saline (PBS), harvested, centrifuged, and resuspended at 1×10^5 cell/mL in 1% low melting point agarose (LMPA, Trevigen). Then 50 μ L of the cells-LMPA mixture was layered onto frosted microscope slides which were pre-coated with 1% normal agarose. After the agar had been allowed to set at 4 $^{\circ}C$, the slides were immersed in lysis buffer (100 mM Na_2EDTA , 2.5 M NaCl, 10 mM Tris pH 10.0, and 1% Triton X-100) for 1 h at 4 $^{\circ}C$. The slides were then incubated in an alkaline electrophoresis solution (1 mM EDTA, 300 mM NaOH, pH > 13) at 4 $^{\circ}C$ for 40 min,

followed by electrophoresis (1 V/cm) at 4 °C for 30 min. The slides were washed three times with neutralization buffer before immersion in absolute ethanol for 20 min. and air-dried at room temperature. The DNA was stained with SYBR Green (1 µg/mL) for 5 min. at 4 °C. A total of 50 comets per slide, randomly captured at a constant depth of the gel, were examined at 40-fold magnification in a fluorescence microscope (Olympus BX41, Milano, Italy; excitation, 495 nm; emission, 521 nm) connected through a black and white camera to a computer-based image analysis system. Comets were randomly captured at a constant depth of the gel, avoiding the edges of the gel, occasional dead cells, and superimposed comets. DNA damage was measured as tail length (distance of DNA migration from the middle of the body of the nuclear core) using Cell-F software (Olympus).

2.10. Hoechst 33258 staining

Capan-1 cells were seeded into 8-well tissue-culture slides (BD Falcon, Bedford, MA, USA) at 5×10^4 cells/well (0.8 cm²). After 24 h, cells were washed twice with PBS and following 48 and 72 h of treatment with IC₅₀ doses of the tested compound, cells were stained for 5 min with 10 µg/mL of Hoechst (2'-(4-hydroxyphenyl)-5-(4-methyl-1-piperaziny)-2,5'-bi-1H-benzimidazole trihydrochloride hydrate, Sigma–Aldrich) in PBS before being examined by fluorescence microscopy (Olympus).

2.11. Western blot analysis

About 10⁶ Capan-1 cells were treated for 24 and 48 h with IC₅₀ doses of the tested compound. Afterwards, cells were harvested, lysed in RIPA buffer (1% NP40, 0.5% sodium deoxycholate, 0.1% SDS), and centrifuged at 13,000 x g for 15 min. at 4 °C. β-actin was used as a loading control. An equal amount of proteins for each sample was electrophoresed on a 12% SDS-PAGE and blotted to a nitrocellulose membrane. The membrane was incubated for 1 h in PBS-Tween20 (0.05%) containing 5% nonfat milk and then at 37 °C for 1 h with primary antibodies. The membranes were stained with the corresponding peroxidase-conjugated secondary antibodies for 1 h at room temperature and detected by ECL according to the manufacturer's protocol (GE).

2.12. Synthesis of the Cu complexes

[Cu₂{bcmp(-H)}(μ-OH)](NO₃)₂·H₂O (**1**) was prepared by mixing Cu(NO₃)₂·2.5H₂O (380 mg, 2.06 mmol), dissolved in 15 mL of water and a solution of bcmp (402 mg, 1.03 mmol) in 5 mL methanol. After adjusting the pH value to 6.8 with NaOH the green solution was evaporated using a rotavapor. The residue was dissolved in 15 mL methanol and filtered. Slow diffusion of diethyl ether into the alcoholic solution afforded green crystals of formula [Cu₂{bcmp(-H)}(μ-OH)](NO₃)₂·H₂O. Yield: 480 mg (0.71 mmol, 69%). Anal. Calcd. (%) for C₂₁H₄₀N₈O₉Cu₂: C, 37.33; H, 5.96; N, 16.58; found: C, 37.15, H, 5.31; N, 16.88%. ESI MS (positive ion mode): m/z: 596.11 [M-NO₃-H₂O]⁺. Crystals of [Cu₂{bcmp(-H)}(μ-OH)](ClO₄)₂·2.5H₂O (**2**) suitable for X-ray analysis were obtained by slow evaporation (5 weeks) of an aqueous solution of 30 mg of **1** in the presence of 10.8 mg NaClO₄. **Caution!** *Although no problems were encountered in this work, transition metal perchlorate complexes are potentially explosive and should be handled with proper precaution.*

3. Results and discussion

3.1. Synthesis and characterisation of [Cu₂{bcmp(-H)}(μ-OH)](NO₃)₂·H₂O (**1**) and [Cu₂{bcmp(-H)}(μ-OH)](ClO₄)₂·2.5H₂O (**2**)

In this study, the dinucleating ligand 2,6-bis(1,4,7-triazacyclonon-1-ylmethyl)-4-methylphenol (bcmp) with two 9-membered aza-crown binding sites was chosen because of the well-established high affinity of Cu²⁺ ions for 1,4,7-triazacyclonane (tacn). The logK value for mononuclear Cu(tacn)²⁺ is 15.5, suggesting that a highly stable bcmp complex is formed under physiological conditions [27]. Furthermore, Cu complexes of tacn ligands are generally good catalysts for phosphate ester hydrolysis [28, 29]. The aromatic spacer provides a rigid scaffold for the bimetallic site, while the bridging phenoxy group shields the electrostatic repulsion between the metal ions and holds the two Cu²⁺ ions in close proximity. We have previously reported a modified synthetic procedure for bcmp [12].

The dinuclear Cu(II) complex [Cu₂{bcmp(-H)}(μ-OH)](NO₃)₂·H₂O (**1**) was prepared by reacting the ligand bcmp with two equivalents of Cu(NO₃)₂ in a methanol/water mixture at pH 6.8 (Chart 1). The complex has been characterized by elemental analyses and ESI mass

spectrometry. The latter shows a signal centered at $m/z = 596.1$ which can be assigned to the isotopomers of $[\text{Cu}_2\{\text{bcmp}(-\text{H})\}(\mu\text{-OH})(\text{NO}_3)]^+$ (Fig. S1). $[\text{Cu}_2\{\text{bcmp}(-\text{H})\}(\mu\text{-OH})(\text{ClO}_4)_2 \cdot 2.5\text{H}_2\text{O}$ (**2**) was obtained by recrystallizing complex **1** from water in the presence of NaClO_4 . X-ray suitable crystals of **2** were grown by slow evaporation of an aqueous solution at r.t. and the structure is described below.

The potentiometric titration of **1** revealed a single deprotonation step with a pK_a value of 4.69 ± 0.09 (Fig. S2). The pK_a value for the deprotonation of the first of the two terminally bound water ligands in mononuclear $[\text{Cu}(\text{tacn})(\text{H}_2\text{O})_2]^{2+}$ is 7.3 [29]. Thus, the low pK_a value found for **1** is in line with a strong binding of the resulting hydroxide to two Cu ions and can be assigned to the formation of the $\mu\text{-OH}$ species. As expected, the pK_a value is lower than that of the analogous Zn complex ($\text{pK}_a = 5.37$) [12]. The electrochemical properties of the dicopper complex were investigated using cyclic voltammetry (CV). Fig. S3 shows the cyclic voltammogram of the complex in phosphate buffer displaying oxidation and reduction peaks at -0.04 V and -0.62 V (vs. Ag/AgCl), respectively. The redox process is irreversible which suggests a fast decomposition of the $\text{Cu}^{\text{I}}\text{-Cu}^{\text{I}}$ species. This has also been observed for analogous binuclear Cu(II) complexes where redox peaks have been assigned to a Cu centered two-electron reduction process $\text{Cu}^{\text{II}}_2 \rightarrow \text{Cu}^{\text{I}}_2$ [30, 31].

The X-ray structure of **2** is shown in Fig. 1, selected bond lengths and angles are listed in Table 1. The asymmetric unit of the Cu complex contains two crystallographically independent complex cations, four perchlorate anions and five water molecules of crystallization. The two complex cations have very similar geometric parameters and exhibit the typical features of five-coordinate Cu complexes of triazacyclononane and its derivatives [32-34]. Each Cu center adopts a distorted square pyramidal coordination geometry with the bridgehead nitrogen, one of the secondary nitrogens, the bridging phenolate oxygen and the bridging hydroxo group defining the basal plane, while the apical site is occupied by the other secondary nitrogen. The four Cu ions lie 0.205(6) (Cu(1)), 0.197(5) (Cu(2)), 0.180(6) (Cu(3)), and 0.198(6) Å (Cu(4)) above their respective basal planes. The τ values which indicate the degree of trigonal distortion from an ideal square pyramidal coordination geometry range from 0.137 to 0.252. According to the definition by Addison ($\tau = (\alpha - \beta)/60$, where α and β are the two largest bond angles) an ideal square pyramid has a τ value of 0, while τ is 1 for a trigonal bipyramid [35]. As expected for d^9 Cu ions with square pyramidal coordination geometry, the Cu-N bond distances are shorter for equatorial nitrogens (2.001 to 2.017 Å) than for the apical nitrogen (2.198 to 2.257 Å) [36-37]. The apical nitrogens of the

Cu ions are anti to each other. The Cu₂O₂ units are planar (maximum deviation from the best weighted plane 0.005 Å in cation 1 and 0.002 Å in cation 2). The Cu₂O₂ plane and the bridging phenyl ring form a dihedral angle of 29.5(8) and 29.0(5) ° in cation 1 and 2, respectively.

3.2. Biological studies

3.2.1. Cytotoxicity

The antiproliferative effects of **1** were assessed on a panel of six human tumor cell lines including examples of human breast (MCF-7), cervical (A431), colorectal (HCT-15), pancreatic (BxPC3) and ovarian (2008) cancers as well as melanoma. For comparison purposes, the cytotoxicity of cisplatin, the most widely used metal-based anticancer drug, was evaluated under the same experimental conditions. IC₅₀ values, calculated from the dose-survival curves obtained after 72 h of drug treatment from the MTT test, are shown in Table S2. While the cytotoxic activity was considerably lower than that of cisplatin against MCF-7, A431, HCT-15, A549 and 2008 tumor cell lines, the dimetallic Cu(II) complex retained a significant in vitro antitumor activity against pancreatic cancer cells. Tested on a panel of five different pancreatic cancer cell lines (Table 2), compound **1** showed a cell killing activity that was similar or even better than that exerted by cisplatin. Interestingly, the greater efficacy was detected in metastatic site derived pancreatic Capan-1 and AsPC-1 cancer cell, on the other hand, the cytotoxic potency was about two times lower than that of cisplatin against p53 null Mia PaCa-2 cells.

Complex **1** was also tested against a human non-tumor cell line, namely human embryonic kidney HEK293 cells (human non-cancerous cells in rapid proliferation). Notably, against non cancerous cells, complex **1** elicited a cytotoxic activity approximately 5-fold lower than that recorded after cisplatin treatment (Table 2). The selectivity index value, (i.e. the average IC₅₀ towards normal HEK293 cells divided by the average IC₅₀ towards all malignant cells included in the panel) calculated for **1** was approximately 1.5-fold higher than that obtained with the reference metallodrug, thus suggesting a noticeable preferential cytotoxicity of **1** versus neoplastic cells.

3.2.2. DNA damage

The ability of **1** to cause DNA damage in Capan-1 cells was investigated. After treatment with increasing concentrations of **1** for 6 and 12 h, evidence for cleavage of cellular DNA was obtained by alkaline single cell gel electrophoresis (Fig. 2). Cells treated with **1** displayed a statistically significant increase in electrophoretic migration of the DNA fragments, evidenced by well-formed comets. As evident from the comet tail lengths (length of DNA migration), which are directly related to the DNA fragment size, complex **1** caused a marked time- and dose-dependent increase in comet tail lengths confirming its ability to cleave cellular DNA.

3.2.3. Mechanism of cell death induction

Cell death induction in Capan-1 pancreatic cancer cells treated for 48 and 72 h with IC_{50} doses of complex **1** was also studied by monitoring the morphological changes of Hoechst 33258-stained cells *via* fluorescent microscopy analysis (Figure 3A). Compared to control cells, **1**-treated cells presented brightly stained nuclei and morphological features typical of apoptosis, such as chromatin condensation and fragmentation, thus confirming the induction of an apoptotic cell death.

It has been shown that in response to treatment with a variety of DNA-damaging agents an induction of tumor suppressor p53 occurs. The activation of p53 which plays a pro-apoptotic role by inducing and downregulating many fundamental cell death effectors, ultimately leads to apoptotic cell death [38]. By taking into consideration that the cytotoxic potency of **1** is about two times lower against p53 null pancreatic Mia PaCa-2 cells than towards pancreatic p53 wild-type cells (BxPC3, AsPC-1, Capan-1 and PANC-1), we studied the effect of **1** on p53. Notably, complex **1** provoked a concentration-dependent increase in p53 activity (Fig. 3B). Taken together, cytotoxicity and p53 results indicate that this complex can activate apoptotic cell death pathways by a mechanism that is p53-dependent. Thus, cancer cell sensitivity to the studied Cu complex strongly depends on the p53 status.

3.3. DNA cleavage and phosphate diester hydrolysis

To gain insight into the molecular interactions of complex **1** with DNA, we performed *in vitro* experiments on supercoiled (SC) plasmid pUC19 DNA. The cleavage ability of the Cu complex was measured by monitoring the conversion of SC-DNA (form I) into the nicked-circular form (NC, form II) using gel electrophoresis. SC pUC19 DNA (1 $\mu\text{g/mL}$) was incubated with increasing complex concentrations (range 5 - 50 μM) for 3 h. Fig. 4 shows representative gel electrophoresis patterns resulting from incubation of SC-DNA with complex **1**. Treatment with complex **1** resulted in a concentration-dependent pUC19 cleavage. Notably, at 25 μM concentration (lane 3), complex **1** was able to convert SC-DNA into the NC form and at 50 μM concentration (lane 4) complete nicking of DNA was observed as evidenced by the absence of the form I. It is worthy of note that no external reducing agent was added. This does not necessarily suggest a hydrolytic cleavage mechanism, as several self-activating oxidative Cu metallonucleases have been described [39-45]. In fact, Cu complexes that actually cleave DNA by a hydrolytic mechanism are less numerous [46-48]. In order to distinguish between an oxidative and hydrolytic mechanism, ROS scavengers specific for singlet oxygen (DABCO), hydroxyl radical (DMSO) or hydrogen peroxide (DMTU), have been added. The nuclease activity of complex **1** (Figure S4) was not influenced by the addition of these scavengers, thus suggesting that DNA cleavage is hydrolytic. Notwithstanding, we thought it of interest to explore the potential of **1** to mediate phosphate ester hydrolysis using BDNPP (bis(2,4-dinitrophenyl) phosphate) which is a widely used model for the phosphodiester linkages in DNA.

1-mediated BDNPP hydrolysis was monitored by following the increase in the visible absorbance at 400 nm because of the release of 2,4-dinitrophenolate. The dependence of the reaction rate of BDNPP cleavage on the pH value in the presence of **1** was studied over the pH range 3.5 - 12.0. The plot of the pseudo-first-order rate constants *vs.* pH revealed a sigmoidal curve (Fig. 5). Fitting the data to

$$k_{\text{obs}} = (k_{\text{H}_2\text{O}}[\text{H}^+] + k_{\text{OH}}K_{\text{a}})/(K_{\text{a}} + [\text{H}^+])$$

gave $k_{\text{H}_2\text{O}} = 5.75 \times 10^{-6} \text{ s}^{-1}$, $k_{\text{OH}} = 6.0 \times 10^{-4} \text{ s}^{-1}$ and a kinetic $\text{p}K_{\text{a}}$ value of pH 6.97. The kinetic $\text{p}K_{\text{a}}$ value is 2.3 units higher than the thermodynamic $\text{p}K_{\text{a}}$ value for the bridging water in $[\text{Cu}_2\{\text{bcmp}(-\text{H})\}(\mu\text{-H}_2\text{O})]^{3+}$ determined by potentiometric titration ($\text{p}K_{\text{a}} = 4.69$). This might be explained by a weakening of one of the $\mu\text{-OH-Cu}$ bonds on substrate binding (Chart 2). The hydroxide in the catalytically active species can act as a nucleophile or a general base.

To gain more insight into the reaction mechanism, the solvent deuterium isotope effect k_H/k_D was measured. The solvent deuterium isotope effect is the classical method for distinguishing between a nucleophilic and a general base mechanism. Generally, a k_H/k_D value ranging from 0.8 to 1.5 indicates a nucleophilic mechanism for phosphate ester hydrolysis, where no proton transfer is involved in the rate-determining step, while a k_H/k_D value greater than 2 supports a general base mechanism, where M-OH deprotonates an external nucleophile [49]. The average k_H/k_D value obtained at different pL values (L = H, D) for the hydrolysis of BDNPP in the presence of **1** is 0.98 (Table S3) suggesting a direct nucleophilic attack of the OH group to the phosphorous (Chart 2). This agrees with work reported by Neves and co-workers who showed that the hydrolysis of BDNPP by similar binuclear complexes of type $\text{Fe}^{\text{III}}(\mu\text{-OH})\text{M}^{\text{II}}$ and $\text{Fe}^{\text{III}}(\mu\text{-CH}_3\text{COO})_2\text{M}^{\text{II}}$ (M = Zn, Cu, Mn, Fe) proceeds *via* a mechanism that involves intramolecular attack of a terminally Fe^{III} -bound hydroxide [50-53]. Neves et al. was concluded that the $\mu\text{-OH}$ was a poorer nucleophile than a terminal OH. The dependence of the hydrolysis rate on the concentration of the Cu complex (pH 8.0, 40 °C, Fig. S5) shows a linear increase of the hydrolysis rate in the 0.5 - 2.5 mM concentration range for **1**. Higher concentrations could not be analysed, as the complex precipitated out of solution. A linear fit of the data gave an apparent second-order rate constant of 0.047 M s^{-1} . This agrees favorably with the apparent second order rate constant of 0.021 M s^{-1} reported for the closely related dinuclear $\text{Cu}^{\text{II}}(\mu\text{-OH})\text{Cu}^{\text{II}}$ complex with the Schiff base ligand 6-amino-6-methylperhydro-1,4-diazepine [54, 55]. However, DFT calculations for this complex suggested a mechanism that involves nucleophilic attack by the bridging hydroxide of the Schiff base complex.

4. Conclusions

$[\text{Cu}_2\{\text{bcmp}(-\text{H})\}(\mu\text{-OH})](\text{NO}_3)_2$ is a new dicopper(II) complex which shows interesting cytotoxic activity against pancreatic cancer cell lines and selectivity for cancer cells over non-cancer cells. The Cu complex can interact with cellular DNA and induce apoptotic cell death via the p53-dependent pathway. Furthermore, the complex promotes the hydrolysis of the DNA model BDNPP by a mechanism involving nucleophilic attack by Cu-OH and converts SC-DNA into the nicked form in the absence of external reductants. Taken together, the reported results herein confirm that the dinuclear bis(triazacyclonane) copper complex represents an attractive “self-activating” metallonuclease. Moreover, the remarkable

results obtained in terms of selectivity against Capan-1 pancreatic cancer cell call for further studies focused on characterizing the antitumor efficacy against this cancer type.

5. Abbreviations

bcmp	2,6-bis(1,4,7-triazacyclonon-1-ylmethyl)-4-methylphenol
BDNPP	bis(2,4-dinitrophenyl) phosphate
CHES	<i>N</i> -cyclohexyl-2-aminoethanesulfonic acid
DABCO	1,4-diazabicyclo[2.2.2]octane
DMSO	dimethylsulfoxide
DMTU	dimethylthiourea
EDTA	ethylenediaminetetraacetic acid
ESI	Electrospray ionization
HEPES	4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid
MES	2-(<i>N</i> -morpholino)ethanesulfonic acid
PBS	phosphate buffered saline
PIPBS	piperazine- <i>N,N</i> -bis(4-butanesulfonic acid)
TBE	tris/borate/EDTA

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Appendix A. Supplementary data

Crystallographic data for **2** have been deposited with the Cambridge Crystallographic Data Centre, CCDC no. 920701. Copies of the data may be obtained free of charge from The Director, CCDC, 12 Union Road, Cambridge CB2 1EZ, UK (fax:+44-1223-336033; e-mail: deposit@ccdc.cam.ac.uk or www: <http://www.ccdc.cam.ac.uk>). Supplementary data to this article can be found online at <http://dx.doi.org/.....> These data include cytotoxicity data of **1**, the ESI MS spectrum of **1**, potentiometric titration data of **1**, the cyclovoltammogram of **1**, gel electrophoresis patterns of radical scavenger experiments, the rate-concentration profile and k_H/k_D values for **1**-mediated BDNPP hydrolysis.

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Figure Legends

Chart 1. Synthetic route for the synthesis of the Cu complex **1**: i) BOC-ON, CHCl₃; ii) THF/CHCl₃; iii) TFA/CH₂Cl₂/Et₂O; iv) Cu(NO₃)₂.

Chart 2. Possible mechanism of action for the hydrolysis of BDNPP by complex **1**.

Fig. 1. View of the cation of [Cu₂{bcmp(-H)}(μ-OH)](ClO₄)₂·2.5H₂O (**2**) with the atom numbering scheme. For the sake of clarity, only one of the two crystallographically independent complex cations is shown. Thermal ellipsoids are drawn at the 40 % level.

Fig. 2. Single cell gel electrophoresis. Capan-1 cells were treated for different incubation times (6 or 12 h) with increasing concentration of **1** and then processed for comet assay. The comet tail length was calculated from the center of the cell and measured in micrometers with CellF software. The error bars indicate the S.D.

Fig. 3. Apoptosis induction and p53 activation. **A.** Hoechst 33258 staining. Capan-1 cells were incubated for 48 and 72 h with IC₅₀ doses of complex **1** and stained with the fluorescent dye Hoechst 33258. **B.** Capan-1 cells were treated with increasing concentrations of **1** for 48 h. The amount of p53 was detected by Western blotting analysis as described in the Experimental Section.

Fig. 4. DNA cleavage. Agarose gel electrophoresis patterns of SC pUC19 DNA incubated with complex **1** in Tris buffer at 37 °C for 3 h. Lane 1: DNA control; lane 2: DNA + 5 μM **1**; lane 3: DNA + 25 μM **1**; lane 4: DNA + 50 μM **1**.

Fig. 5. Rate-pH profile for the cleavage of BDNPP (5×10^{-5} M) in the presence of **1** (1 mM) at 40 °C. [buffer] = 50 mM (buffer = PIPBS (pH 3.5 – 5), MES (pH 5 – 6.7), HEPES (pH 6.8 – 8.5), and CHES (pH 8.5 – 9.5); $I = 0.1$ M (KNO_3)).

Table 1. Selected bond lengths [Å], angles [°] and hydrogen bonding interactions in compound **2**

Cu(1)–N(1)	2.009(6)	Cu(3)–N(1')	2.035(6)
Cu(1)–N(2)	2.015(6)	Cu(3)–N(2')	2.223(7)
Cu(1)–N(3)	2.198(8)	Cu(3)–N(3')	2.013(6)
Cu(1)–O(1)	1.933(5)	Cu(3)–O(1')	1.934(5)
Cu(1)–O(2)	1.980(5)	Cu(3)–Cl(1')	1.944(5)
Cu(2)–N(4)	2.017(6)	Cu(4)–N(4')	2.044(6)
Cu(2)–N(5)	2.015(6)	Cu(4)–N(5')	2.001(6)
Cu(2)–N(6)	2.257(7)	Cu(4)–N(6')	2.232(6)
Cu(2)–O(1)	1.922(5)	Cu(4)–O(1')	1.941(5)
Cu(2)–O(2)	1.961(5)	Cu(4)–O(2')	1.946(5)
Cu(1)···Cu(2)	3.015(1)	Cu(3)···Cu(4)	2.981(1)
N(1)–Cu(1)–N(2)	86.5(3)	N(1')–Cu(3)–N(2')	84.0(3)
N(1)–Cu(1)–N(3)	85.4(3)	N(1')–Cu(3)–N(3')	86.9(3)
N(2)–Cu(1)–N(3)	84.3(3)	N(2')–Cu(3)–N(3')	83.3(3)
N(1)–Cu(1)–O(2)	161.1(2)	N(1')–Cu(3)–O(2')	161.4(2)
N(2)–Cu(1)–O(1)	173.1(3)	N(3')–Cu(3)–O(1')	176.5(3)
N(4)–Cu(2)–N(5)	86.5(2)	N(4')–Cu(4)–N(5')	86.7(2)
N(4)–Cu(2)–N(6)	83.6(2)	N(4')–Cu(4)–N(6')	83.2(2)
N(5)–Cu(2)–N(6)	82.0(2)	N(5')–Cu(4)–N(6')	82.8(2)
N(4)–Cu(2)–O(2)	166.7(2)	N(4')–Cu(4)–O(2')	162.2(2)
N(5)–Cu(2)–O(1)	176.4(3)	N(5')–Cu(4)–O(1')	172.9(3)
Cu(1)–O(1)–Cu(2)	102.9(2)	Cu(3)–O(1')–Cu(4)	100.6(2)
Cu(1)–O(2)–Cu(2)	99.8(2)	Cu(3)–O(2')–Cu(4)	100.0(2)

Table 2. IC₅₀ values (μM) of **1** and cisplatin for pancreatic cancer cell lines. Cells (3-8×10⁴·ml⁻¹) were treated for 72 h with increasing concentrations of tested compounds. Cell viability was evaluated by means of MTT test. IC₅₀ values were calculated by the dose-response curves by means of four parameter logistic model (p<0.05). SD = standard deviation.

Compound	IC₅₀ (μM) ± SD	
	1	cisplatin
MIA PaCa-2	30.43±3.15	16.76±1.89
Panc-1	21.19±4.25	8.54±3.54
Capan-1	11.67±6.52	13.17±0.97
BxPC3	18.77±2.24	11.13±2.36
AsPC-1	14.73±2.52	15.11±1.94
HEK293	89.76±4.65	17.62±2.87