



Research Article

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Hetero-Binuclear Complexes of Lanthanum (III) Using Bridging *N,N'*-Bis(2-Pyridylmethyl)Oxamide and Terminal 1,10-Phenanthroline: Synthesis, Characterization and Biological Evaluation

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ABSTRACT

In the present study, hetero-binuclear La(III) complexes were newly synthesized using well known bridging ligand *N,N'*-bis(2-pyridylmethyl)oxamide (2PMO, **1**) and “complexes as ligands” [Cu(2PMO)] (**2**) and [Ni(2PMO)] (**3**). The newly synthesized mononuclear (**4**) as well as hetero-binuclear complexes (**5** and **6**) were characterized by analytical and spectroscopic techniques. The newly synthesized complexes were tested for their binding ability towards CT-DNA and chemical nuclease property towards SC pUC 19 DNA. The complexes were tested for antibacterial activities against Gram negative bacteria (*E. coli* and *K. pneumonia*) and Gram positive bacteria (*B. subtilis* and *S. aureus*), antioxidant activities by DPPH radical scavenging and ferrous-ion chelation methods and cytotoxicity against MCF-7 cell lines by MTT assay. The structure-activity relationship revealed that the newly synthesized hetero-binuclear complexes show potent DNA binding and chemical nuclease activities, antibacterial, antioxidant and cytotoxic activities.

Keywords: Antibacterial, antioxidant, antitumor, DNA interactions, oxamides, Lanthanum (III) complexes.

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INTRODUCTION

The cleavage of DNA by metal complexes is of pharmaceutical industries. Chemical nuclease activity by metal complexes usually takes place either by hydrolytic pathway where phosphodiester bond is hydrolyzed or

oxidative pathway which involves abstraction of proton from sugar-phosphate backbone or nitrogenous bases. Hydrolytic pathway results in the 3'-OH and 5'-OPO₃H fragments that can be relegated by natural enzymes whereas oxidative pathway results in the formation of non-sticky ends which cannot be relegated

by natural enzymes. [1-2] The oxidative pathway is catalyzed by the activation of metal complex in presence of oxidizing agent or reducing agent to generate diffusible free radicals which damage DNA. Thus, cleavage of DNA by oxidative pathway by metal complexes is highly beneficial in designing antitumor and antimicrobial agents. The chemical nuclease activity of the metal complexes depends on their interaction with DNA. The modes of non-covalent interactions include electrostatic effect, groove binding and intercalation of metal complexes with DNA or proteins are directed by the nature of the ligand, metal centers, geometry and coordination sphere of the metal complexes. [3-6] The efficiency of the metal complex in binding to the DNA is decided by the type of coordinated ligand. The presence of extended aromatic planar ring in the ligand moiety directs the metal complex in targeting DNA. [7] The oxidative stress induced by the free radicals that are generated in human body by various metabolic processes releases many free radicals such as hydroxyl radicals, superoxide radicals, singlet oxygen etc. The enzymes in the body to counteract these ions are limited and in turn these radicals are responsible for life threatening diseases such as carcinogenesis, inflammations etc. The compounds having the ability to suppress the activities of these free radicals are known as antioxidants. [8-9] These free radicals if produced in excess can cause oxidative damage of the DNA that causes cancerous effects. Thus, there is a need for drug molecules which possess diversified applications such as chemical nuclease activities, antimicrobial, antioxidant and antitumor activities are highly beneficial.

The coordination of suitable ligands also decides the biological properties of the metal complexes. Ligands having bioactive moieties such as $-C=O$, $-NH$, $-SH$, $-OH$ and aromatic planar rings can enhance the antimicrobial properties of the metal complexes. The coordinating ligands increase the lipophilicity of the complexes by sharing the positive charge on the metal ions by chelation thereby enhancing the permeability of the complexes into biological membranes. [10] These properties introduce metal complexes as potent antimicrobial agents.

The structure and geometry of the metal complexes decide the biological role of the metal complexes. In this regard, several investigations have been carried out in exploring the pharmaceutical values of lanthanum(III) complexes. La(III) in its stable +3 oxidation state mimics biological ions such as Ca^{2+} and K^+ and hence can be highly beneficial in synthesizing chemical nucleases. [11] La(III) complexes have been used as probes in immunoassays and as MRI agents, [12-14] anticancer agents, [15-19] binding agents towards DNA, RNA, proteins and as chemical nucleases, [20-24] antimicrobial agents [25-30] as well as antioxidant agents. [31-32] In this view, herein we report the synthesis and characterization of mononuclear as well as hetero-

binuclear complexes containing lanthanum(III), copper(II) or nickel(II) ion bridged by versatile ligand *N,N'*-bis(2-pyridylmethyl)oxamide and ancillary ligand 1,10-phenanthroline which is a classical DNA intercalator. We aimed to evaluate their biological properties such as DNA interactions, antibacterial, antioxidant and antitumor activities by standard procedures. The newly synthesized hetero-binuclear complexes were found to possess higher DNA binding strengths towards CT-DNA, chemical nuclease activity towards SC pUC19 DNA, antibacterial activity against Gram negative bacteria (*E. coli* and *K. pneumonia*) and Gram positive bacteria (*B. subtilis* and *S. aureus*), free radical scavenging and metal ion chelating abilities and antitumor activities against MCF-7 cell lines. Thus, they can be utilized as potent diagnostic as well as pharmaceutical tools.

MATERIALS AND METHODS

Materials

All the reagents and chemicals were obtained from Sigma Aldrich (USA) and Fluka (USA) and used as obtained. Solvents used for spectroscopic and electrochemical procedures were purified by standard procedure reported earlier. [33] Super coiled (SC) pUC 19 DNA (cesium chloride purified) was purchased from Bangalore Genei (India). Calf-thymus DNA (CT-DNA), Agarose (molecular biology grade), Ethidium bromide (EB) was obtained from Sigma Aldrich (USA). 2,2-diphenyl-1-picrylhydrazyl (DPPH), Ferrozine (Extra pure) were purchased from Himedia and used as received. Ferrous chloride and Tris (tris-hydroxymethyl(aminomethane)) were purchased from Merck and used as purchased. Tris-(hydroxymethyl) aminomethane-HCl (pH = 7.2) buffer solution was prepared using deionized, double distilled water.

Bacterial media was purchased from Himedia. The bacterial strains *Escherichia coli* (ATCC 25922), *Klebsiella pneumonia* (MTCC 109), *Staphylococcus aureus* (MTCC 3160) and *Bacillus subtilis* (MTCC 441) were procured from IMTECH, Chandigarh, India. Human breast cancer cells (MCF-7) were purchased from ATCC. Foetal bovine serum (FBS), penicillin (100 IU/mL), streptomycin (100 µg/mL), glutamine, RPMI-1640, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT), dimethyl sulfoxide (DMSO) chemicals required for cytotoxicity studies were purchased from Hi-media (Mumbai, India).

Methods

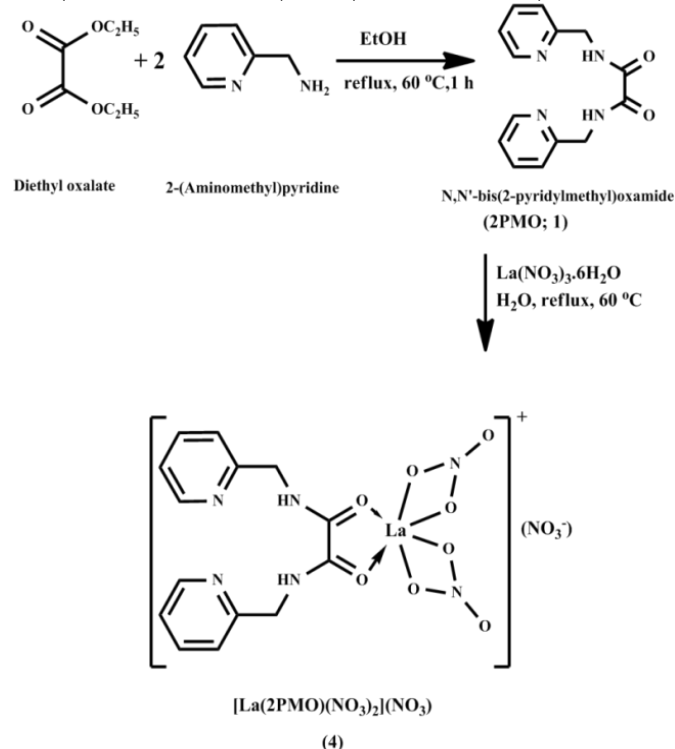
Syntheses

The ligand *N,N'*-bis(2-pyridylmethyl)oxamide (2PMO, 1), [34] $[Cu(2PMO)]$ (2) and $[Ni(2PMO)]$ (3) were synthesized by modifying literature procedure [35] for the comparison of their activities with those of newly synthesized mononuclear and binuclear-complexes. The new mononuclear $[La(2PMO)(NO_3)_2](NO_3)$ (4) and hetero-binuclear complexes 5-6 were synthesized by following reported methods with modifications. [36]

Synthesis of $[La(2PMO)(NO_3)_2](NO_3)$ (4)

Diethyl oxalate (0.408 mL, 0.1 M) and 2-(aminomethyl)pyridine (0.432 mL, 0.2 M) were dissolved in 20 mL ethanol. The solution was refluxed at 60 °C for 1 h. Aqueous solution of $\text{La}(\text{NO}_3)_3 \cdot 6\text{H}_2\text{O}$ (0.433 g, 0.1 M) was added to the above reaction mixture and refluxing was continued for 2 h. The clear solution obtained was kept for slow evaporation, yielded white solid. The solid obtained was washed with warm ethanol: water mixture (1:1 v/v) and dried in a desiccator. Yield: 75%.

Anal. Calcd. (%) for $\text{C}_{14}\text{H}_{14}\text{LaN}_7\text{O}_{11}$ (**4**): C, 28.25, H, 2.37, N, 16.47. Found, C, 28.23, H, 2.35, N, 16.44. $^1\text{H-NMR}$ ($\text{DMSO}-d_6$, 400 MHz, ppm): δ = 4.265 – 4.485 (m, 2H), 7.261 – 7.302 (q, 2H), 7.762 – 7.766 (td, 1H), 9.268 – 9.313 (dt, 1H). FT-IR (KBr disc, cm^{-1}): ν = 3298 (N-H), 1660 (C=O), 1518 (C=N), 1480 and 1319 (NO_3^- , coordinated), 1384 (NO_3^- , ionic), 404 (La-O). λ_{max} (DMF/Tris-HCl) 234.95 nm (ϵ = 12,633 $\text{M}^{-1}\text{cm}^{-1}$), 260.34 nm (ϵ = 11,720 $\text{M}^{-1}\text{cm}^{-1}$). Λ_{M} (DMF, $\text{S cm}^2\text{M}^{-1}$): 117.05.



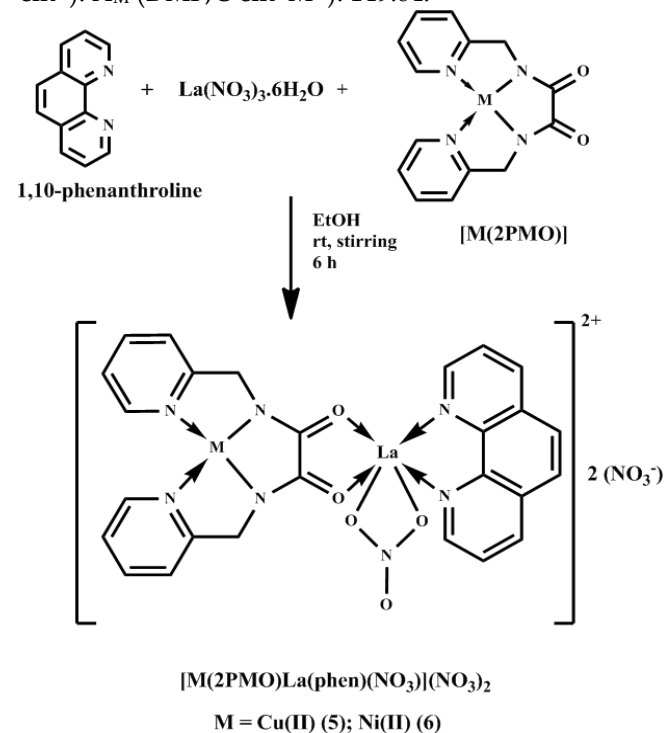
Scheme 1: Proposed synthetic route of $[\text{La}(\text{2PMO})(\text{NO}_3)_2](\text{NO}_3)$ (**4**).

The concept of “complexes as ligands” was used to synthesize binuclear complexes. $[\text{Cu}(\text{2PMO})]$ and $[\text{Ni}(\text{2PMO})]$ were used as “complex as ligands”. A solution of 1,10-phenanthroline (0.2 mM, 0.040 g, 10 mL) in ethanol was stirred constantly. To this an aqueous solution of $\text{La}(\text{NO}_3)_3 \cdot 6\text{H}_2\text{O}$ (0.2 mM, 120 mg, 15 mL) was added dropwise with continued stirring. Then aqueous solutions of $[\text{Cu}(\text{2PMO})]$ (0.2 mM, 0.066 g, 10 mL for **5**) and $[\text{Ni}(\text{2PMO})]$ (0.2 mM, 0.060 g, 10 mL for **6**) were added respectively in dropwise and stirring was continued for further 6 h. The resulting solutions were filtered. Dark green solid (**5**) and brown solid (**6**) were obtained on slow evaporation of the respective solutions. The solids were washed with

ethanol: water (1:1 v/v) mixture and dried in a desiccator. Yield: 68% (**5**); 76% (**6**) (**Scheme 2**).

Anal. Calcd. (%) for $\text{C}_{26}\text{H}_{20}\text{CuLaN}_9\text{O}_{11}$ (**5**): C, 37.31, H, 2.41, N, 15.06. Found, C, 37.29, H 2.39, N, 15.02. $^1\text{H-NMR}$ ($\text{DMSO}-d_6$, 400 MHz, ppm): δ = 4.209 – 4.488 (d, 3H), 7.295 – 7.788 (m, 4H), 8.260 – 8.637 (m, 2H), 9.307 – 9.457 (d, 1H). FT-IR (KBr disc, cm^{-1}): 1664 (C=O), 1520 (C=N), 1484 and 1318 (NO_3^- , coordinated), 1389 (NO_3^- , ionic), 402 (La-O), 494 (Cu-N). λ_{max} (DMF/Tris-HCl): 230.30 nm (ϵ = 13,366 $\text{M}^{-1}\text{cm}^{-1}$), 260.08 nm (ϵ = 7,766 $\text{M}^{-1}\text{cm}^{-1}$). Λ_{M} (DMF, $\text{S cm}^2\text{M}^{-1}$): 145.65.

Anal. Calcd. (%) for $\text{C}_{26}\text{H}_{20}\text{NiLaN}_9\text{O}_{11}$ (**6**): C, 37.53, H, 2.42, N, 15.15. Found, C, 37.51, H 2.40, N, 15.12. $^1\text{H-NMR}$ ($\text{DMSO}-d_6$, 400 MHz, ppm): δ = 4.199 – 4.434 (d, 3H), 7.262 – 7.872 (m, 4H), 8.416 – 8.626 (m, 2H), 9.287 – 9.298 (d, 1H). FT-IR (KBr disc, cm^{-1}): 1666 (C=O), 1519 (C=N), 1489 and 1312 (NO_3^- , coordinated), 1382 (NO_3^- , ionic), 401 (La-O), 491 (Cu-N). λ_{max} (DMF/Tris-HCl): 230.58 nm (ϵ = 13,367 $\text{M}^{-1}\text{cm}^{-1}$), 260.08 nm (ϵ = 10,333 $\text{M}^{-1}\text{cm}^{-1}$). Λ_{M} (DMF, $\text{S cm}^2\text{M}^{-1}$): 149.64.



Scheme 2. Proposed synthetic route of $[\text{M}(\text{2PMO})\text{La}(\text{phen})(\text{NO}_3)](\text{NO}_3)_2$. M = Cu (**5**); Ni (**6**).

General Methods

The elemental analyses of carbon, hydrogen and nitrogen of the newly synthesized La(III) mononuclear (**4**) and binuclear complexes (**5** and **6**) were carried out using a Thermo Finnigan FLASH EA 1112 CHNS analyzer. ^1H NMR spectra of the complexes were recorded on 400 MHz liquid state Bruker NMR spectrometer obtained in $\text{DMSO}-d_6$ solution at approximately 25°C using tetramethyl silane (TMS) as internal standard. The IR spectra of the ligand and complexes were recorded using KBr pellets over the region 4000–400 cm^{-1} on a Shimadzu FT-IR spectrophotometer. UV-Visible absorption spectra of

the ligand and complexes were recorded on SpectraMax Plus 384 spectrophotometer. 10^{-3} M solutions of the ligand and complexes were prepared in DMF solvent and recorded in the wavelength ranging from 200 to 800 nm. The time-dependent stability of the complexes **4-6** in solution state were measured upto 6 h at pH = 7.2, 25 °C in DMF/Tris-HCl buffer by UV-Visible absorption spectroscopy using SpectraMax Plus 384 spectrophotometer. Full scan mass spectra (ESI-MS) of the complexes were recorded in positive ion mode on Shimadzu-LCMS equipped with APCI and ESI probes. The corresponding *m/z* values were recorded in the expected range. Conductivity measurements were carried out using 10^{-3} M solutions in DMF on Control Dynamics (India) conductivity meter with a dip-type cell, calibrated with a (0.1 M) solution of AnalaR potassium chloride. The molar conductivity values were used to determine the electrolytic nature of the complexes. Electrochemical experiments were performed on a CHI 600E Electrochemical analyzer controlled by PC. The cyclic voltammograms of 10^{-3} M solutions of ligand and binuclear complexes in DMF solvent were recorded in the potential range of -0.4 V to 0.7 V at the scan rate of 25 – 200 mV s⁻¹ at 25°C using 0.1 M tetra butyl ammonium perchlorate (TBAP) as supporting electrolyte. The three-electrode configuration was made up of glassy carbon as the working electrode, standard calomel electrode (SCE) as reference electrode and platinum wire as auxiliary electrode. Prior to each measurement, the working electrode and auxiliary electrodes were cleaned and polished with solid alumina of 1.0 μ, 0.5 μ and 0.05 μ respectively. The ligand and complex solutions were degassed with N₂ atmosphere for 10 min prior to the electrochemical measurements.

DNA Interaction Studies

The ability of the newly synthesized complexes **4-6** in comparison with mononuclear complexes **2-3** and the ligand 2PMO (**1**) to interact with nucleic acid (DNA) is studied by their binding interactions with CT-DNA and cleaving ability towards SC pUC 19 DNA.

DNA binding experiments

UV-Visible absorption titrations

UV-Visible absorption titration experiment was carried out to study the interaction of La(III) complexes with CT-DNA. Absorbance was measured at 260 and 280 nm for CT-DNA in 5mM Tris-HCl buffer (pH = 7.2) gave the ratio of 1.9:1, indicating the DNA was free of protein. The intensity of the absorbed band at 260 nm with a known ϵ value ($6600 \text{ M}^{-1}\text{cm}^{-1}$) was used to calculate the concentration of CT-DNA. [37] The absorption titrations were carried out at constant concentration of the complexes (30 μM) whereas concentration of CT-DNA was varied. Stock solutions (10^{-3} M) of complexes **2-6** were prepared by dissolving the complexes in DMF and diluted further to 30 μM using 5mM Tris-HCl buffer. The absorbance of DNA while measuring the spectra was eliminated by adding equal amounts of DNA to both complex and reference

solutions. The complex solution was allowed to get equilibrated for 5 mins after the addition of known aliquots of CT-DNA solution to allow the complex to bind to the DNA. [31] The changes in the absorption intensities were monitored at regular intervals and used to calculate the intrinsic equilibrium binding constant (K_b) values by McGhee-von Hippel (MvH) method using the following equation. [38]

$$(\epsilon_a - \epsilon_b) / (\epsilon_b - \epsilon_f) = (b - (b^2 - 2K_b C_t [\text{DNA}]/s)^{1/2}) / 2K_b C_t$$

Where b is $(1 + K_b C_t + K_b [\text{DNA}]/2s)$, ϵ_a is the extinction coefficient observed for the charge transfer absorption band at a given concentration of DNA, ϵ_f is the extinction coefficient observed for the absorption band of the complex in the absence of DNA, ϵ_b is the extinction coefficient of the complex in fully bound form with DNA, K_b is the intrinsic equilibrium constant, C_t is the total concentration of the metal complex in solution, $[\text{DNA}]$ is the concentration of DNA in nucleotides and s is the binding site size in base pairs. The non-linear least square analyses were carried out using the software Origin Lab, version 6.1. [39]

Electrochemical titrations - Cyclic voltammetry

The mode of interaction of the ligand 2PMO (**1**) and the complexes **2-6** with CT-DNA were further confirmed by carrying out electrochemical experiments. [40] Cyclic voltammetric measurements were recorded using glassy carbon as working electrode, saturated calomel as reference electrode and platinum wire as auxiliary electrode. All the measurements were recorded using Tris-HCl (5 mM)/NaCl (50 mM) buffer (pH = 7.2) as supporting electrolyte at 25 °C. The reaction solutions were degassed in N₂ atmosphere for 10 mins before carrying out the experiment. The redox behavior of the 30 μM 2PMO (**1**) and the complexes **2-6** were studied in the absence and presence of CT-DNA (90 μM) at the scan rate of 25 mV s⁻¹ to 200 mV s⁻¹ in the potential range -0.4 to 0.7 V. The cyclic voltammetric experiment was also carried out to study the binding nature of 2PMO (**1**) and the complexes **2-6** at constant concentration (30 μM) with increasing concentration of CT-DNA at the scan rate of 200 mVs⁻¹ in the potential range of -0.4 to 0.7 V. The shift in the peak potential values indicates the mode of interaction of the ligand and complexes with CT-DNA. The formal potentials of the ligand and complexes in the free form ($E_f^{0'}$) and fully bound form ($E_b^{0'}$) were used to calculate the binding constant ratios using the following formula:

$$E_b^{0'} - E_f^{0'} = 0.0591 \log [K_R/K_O]$$

Where K_R/K_O is the ratio of binding constants of the complexes in the reduced form to the oxidized form.

Viscosity measurements

Viscometric titrations were carried out using Micro-Ubbelohde Viscometer which was thermo-stated at 37°C in a constant temperature bath. The concentration of CT-DNA was kept constant (150 μM, NP) while concentration of complex solutions were increased at regular aliquots. The flow times were measured with automated times and each sample was measured three

times for accuracy and average time flow was calculated for each complex solution. A graph of $(\eta/\eta_0)^{1/3}$ vs. $[\text{complex}]/[\text{DNA}]$ was plotted where η is the viscosity of CT-DNA in the presence of complex solution and η_0 is the viscosity of CT-DNA alone. Viscosity values were calculated using observed flow times of DNA containing solutions (t) corrected to that of buffer alone (t_0).^[41]

$$\eta = (t - t_0) / t_0$$

DNA cleavage studies

Chemical nuclease activity – Pilot experiment

The agarose gel Electrophoresis method was used to determine the ability of the 2PMO (**1**) and the complexes **2-6** to cleave super coiled (SC) pUC 19 DNA to the nick circular (NC) form. The gel electrophoresis was carried out using 16-toothed combed gel trays of 210-150 mm. A constant voltage of 60 V at 10 mA was supplied using an Electrophoresis Power Supply Bangalore-Genei (India) system during electrophoresis. The cleavage experiment was carried out using SC pUC19 DNA (33.3 μM , 2 μg , NP) in 50mM Tris-HCl/50 mM NaCl (pH = 7.2). The concentration of the ligand and complexes were corresponding to the final volume of the reaction mixture in the gel-well. The band intensities were used to measure the extent of cleavage of SC pUC 19 DNA using UVITEC Gel Documentation System. Due corrections were made for the low level of nicked circular (NC) form present in the original super coiled (SC) DNA sample and for the low affinity of EB binding to SC compared to NC and linear forms of DNA.^[42]

Chemical nuclease activity – Mechanistic studies

Mechanistic studies were carried out to study the involvement of free radicals in the oxidative cleavage of SC pUC 19 DNA by complexes. Agarose gel electrophoresis was carried out in the presence of radical quenchers such as DMSO which inhibit hydroxyl radicals and sodium azide (NaN_3) which inhibits singlet oxygen. Methyl green, a major groove binder was used to study the site of binding of hetero binuclear La(III) complexes **5** and **6** (10 μM) during oxidative cleavage.^[43]

Antibacterial activities

The antibacterial properties of the ligand 2PMO (**1**) and its complexes **2-6** were determined by following standard protocol of Agar disc diffusion method.^[44-45] The test organisms *Escherichia coli*, *Klebsiella pneumonia*, *Bacillus subtilis* and *Staphylococcus aureus* were maintained on nutrient agar slants overnight. The bacterial cultures were then centrifuged at 8000 rpm for 10 min. The bacterial cells were suspended in saline to make a suspension of 10^5 CFU/ mL and used for assay. The bacterial culture was then spread over the nutrient agar plate using sterile swab and the petri plates were allowed to dry. The test samples 2PMO (**1**) and its complexes (**2-6**) were dissolved in DMSO (1 mg mL⁻¹). The discs were prepared from Whatman filter paper and had been impregnated with DMSO solutions of 2PMO (**1**) and its complexes (**2-6**) respectively were

placed on agar surface. Each test plate consists of one standard antibiotic tetracycline (30 μg mL⁻¹) and the other test samples (50 μg mL⁻¹ and 100 μg mL⁻¹). The discs were placed equidistant to one another. The plates were then incubated at 37°C for 24 h. After incubation, the plates were checked for zone of inhibition. The inhibition zones were measured and compared with the standard, tetracycline. DMSO was used as negative control.

Antioxidant activities

The antioxidant activities of synthesized 2PMO (**1**) and its complexes (**2-6**) were tested by DPPH free radical scavenging method and ferrous ion chelating activity.

DPPH free radical scavenging activity

The most widely used method to check the antioxidant activity is the DPPH radical scavenging activity method. DPPH being a stable nitrogen based free radical gives violet colour in its oxidized form, changes to yellow colour on reduction either by electron transfer or hydrogen transfer. The compounds which facilitate this reaction are considered to be radical scavengers or antioxidants. DPPH has an absorbance of 517 nm in its radical form that disappears after accepting hydrogen radical or electron from the antioxidant and becomes a stable diamagnetic substance.^[46-47] The DPPH radical scavenging ability of the synthesized ligand 2PMO (**1**) and its complexes (**2-6**) were determined by modifying the method reported by Brand-Williams *et al.*^[57] 1 mg mL⁻¹ stock solutions were prepared by dissolving 2PMO (**1**) and its complexes (**2-6**) in DMF. Test solutions (10-70 μg mL⁻¹) were taken in 96-well plate and the volumes were made up to 200 μL using 5 mM Tris-HCl buffer (pH = 7.4). 200 μL of positive controls were prepared using Butylated hydroxy toluene (BHT) and Ascorbic acid (AA) in a manner exactly same as test samples. DPPH in ethanol (500 μM) was added to all the wells. The 96-well plate was then kept under dark for 30 min incubation. Blank solution contained same experimental solutions except that of test samples. Absorbance was then measured at 517 nm against the blank. The decrease in absorbance measures the radical scavenging ability of the complexes. Radical scavenging ability was expressed as %I i.e., the ability of complexes to scavenge DPPH radicals.

Ferrous ion chelating assay

Ferrozine has the ability to chelate Fe^{2+} ions resulting in red colour. When other chelating agents are present, the ferrozine- Fe^{2+} complex formation is disrupted and the red colour will be diminished. The decrease in the red colour formation is a measure of chelating ability of the chelating agents present and the decreased intensity is measured at 562 nm by UV-Visible spectrophotometer. The ability of the 2PMO (**1**) and its complexes (**2-6**) to chelate Fe^{2+} ions were determined by the reported methods with slight modifications.^[48-49] 1 mg mL⁻¹ stock solutions were prepared by dissolving 2PMO (**1**) and its complexes (**2-6**) in DMF. 10-70 μL from the stock solutions of the tested complexes were

added to a 96-well microplate and the volumes were made upto 150 μ L using distilled water. To these wells, 2 mM FeCl_2 solution (20 μ L) was added. The reaction was initiated by adding 5 mM Ferrozine (30 μ L) to each well and shaken vigorously. The microplate was then incubated for 10 min at room temperature. Once the reaction mixture attained equilibrium, absorbance of each well was measured at 562 nm against the blank (reaction mixture without test samples). EDTA- Na_2 was used as standard. The reaction mixture without test sample is used as blank. The ferrous ion chelating ability (% inhibition of formation of ferrozine- Fe^{2+} complex) was calculated using the formula:

$$\text{Ferrous ion chelating ability (\%)} = [1 - A_s/A_0] \times 100$$

Where A_s is the absorbance in the presence of test sample and A_0 is the absorbance of blank. A graph using %I i.e., % inhibition of ferrozine- Fe^{2+} complex formation vs. concentration of sample were plotted. Using the graph, EC_{50} , the efficient concentration at which 50% of the metal chelation was occurred was calculated.

Cytotoxicity activities – MTT assay

The cytotoxic effects of 2PMO (**1**) and its La(III) complexes (**5** and **6**) were tested against human breast cancer cells (MCF-7) by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay as per the reported procedure with slight modifications. [50-51] The assay of the test samples 2PMO (**1**) and its complexes (**5** and **6**) was carried out in the concentration range 10-320 $\mu\text{g mL}^{-1}$. MTT assay is a colorimetric assay based on the fact that 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) dye is a pale yellow substrate taken up by live cell mitochondria which is then reduced by mitochondrial oxidoreductase enzymes into an insoluble purple color formazan product. The cellular oxidoreductase enzymes may reflect the number of viable cells under given conditions. Briefly, MCF-7 cells (3×10^3 cells/well) in RPMI-1640 medium (supplemented with 10% FBS, 100 U/mL penicillin, 100 $\mu\text{g mL}^{-1}$ streptomycin and 2 mM glutamine) with a final volume of 200 μ L were seeded into 96-wells culture plate and incubated overnight at 37°C with the supply of 5% CO_2 . The stock solutions of 2PMO (**1**) and its La(III) complexes (**5** and **6**) were prepared in DMSO of molecular biology grade and diluted to required concentrations using RPMI-1640 medium. The cells were treated with or without test samples of known concentrations and were incubated at 37°C with the supply of 5% CO_2 for 48 h. After incubation, the cells were washed with phosphate buffer saline (PBS). At the end of the treatment period, 20 μ L MTT (5 mg mL^{-1}) was added to each well and incubated for 4 h at 37°C with the supply of 5% CO_2 . After incubation, the blue formazan products formed in the cells were dissolved in DMSO (100 μ L) and measured spectrophotometrically at 540 nm. The percentage cell viability of the complex untreated and

treated cells were calculated and represented graphically.

Table 1: Physicochemical data of complexes 4-6.

	16	17	18
M. W. ^a	595.21	836.94	832.09
M. P. ^b (°C)	204	> 300	> 300
IR ^c , cm^{-1} [$\nu(\text{C}=\text{O})$]	1660	1660	1666
λ^d , nm (ϵ , $\text{M}^{-1} \text{cm}^{-1}$)	234.95 (12,633)	230.30 (13,366)	230.58 (13,367)
Λ_m^e , $\text{S cm}^2 \text{M}^{-1}$	117.05	145.65	149.64
$E_{1/2}^f$ (V)	-0.1233	-0.1122	-0.1073

^a Molecular weight.

^b Melting point

^c IN KBr phase

^d UV-Visible absorption band in DMF.

^e Molar conductivity values in DMF.

^f The formal potentials are vs. SCE. $E_{1/2}$ is the average of cathodic and anodic peak potentials.

RESULTS AND DISCUSSION

Syntheses

One mononuclear complex (**4**) and two binuclear complexes (**5-6**) are newly synthesized following literature method using *N,N'*-bis(2-pyridylmethyl)oxamide (2PMO) as ligand (**1**), $[\text{Cu}(\text{2PMO})]$ (**5**) and $[\text{Ni}(\text{2PMO})]$ (**6**) “complexes as ligands” concept under given conditions as in **Schemes 1-2**. All the newly synthesized complexes are characterized by $^1\text{H-NMR}$, FT-IR spectroscopy, UV-Visible absorption spectroscopy, ESI-Mass spectrometry, molar conductivity measurement, electrochemical measurements and elemental analyses. The complexes are soluble in methanol, dichloromethane, DMF, DMSO and Tris-HCl buffer (pH = 7.2) and are highly stable in DMF/Tris-HCl buffer solutions. All the results are correlated and are in the expected range. Molar measurements reveal that newly synthesized complex **4** is 1:1 electrolyte whereas complexes **5** and **6** are 1:2 electrolytes in DMF solvent.

General Methods

$^1\text{H-NMR}$ spectra reveal the presence of $-\text{CH}_2$ protons in the region 4.19 – 4.48 ppm in the all the newly synthesized complexes **4-6**. Complex **4** has chemical shift values around 7.76 – 7.77 ppm which correspond to $-\text{NH}$ of amide protons. The aromatic protons appear around 7.26 – 7.87 ppm and 9.25 – 9.31 ppm which correspond to pyridyl group in complexes **4-6**. Complexes **5** and **6** have additional aromatic protons around 8.26 – 8.63 ppm, correspond to 1,10-phenanthroline. [7]

IR spectra reveals the presence of characteristic carbonyl ($-\text{C}=\text{O}$) peak of the oxamide ligand (2PMO) in all the complexes **4-6**. The oxamide $-\text{C}=\text{O}$ appears at 1666 cm^{-1} , 1660 cm^{-1} and 1666 cm^{-1} in complexes **4**, **5** and **6** respectively. The ligand carbonyl peak at 1649 cm^{-1} has been shifted to higher frequencies in complexes **4-6**, reveal the coordination of the 2PMO ligand through $-\text{C}=\text{O}$. When oxamide coordinates through $-\text{C}=\text{O}$, the $-\text{N}-\text{C}-\text{O}$ vibrations have been shifted to higher frequencies as in protonated species. This confirms the coordination of oxamide through $-\text{C}=\text{O}$.

C=O bond. The -NH group is deprotonated and coordinated to d-metal ion and hence the secondary -NH peak is absent in IR spectra of complexes **5** and **6**. The secondary -NH peak appears at 3298 cm⁻¹ in complex. The presence of coordinated NO₃⁻ peaks appear at 1480 and 1319 cm⁻¹, 1484 and 1318 cm⁻¹ and 1489 and 1312 cm⁻¹ in complexes **4**, **5** and **6** respectively are due to the ν_4 and ν_1 vibrations of the nitrate group of C_{2v} symmetry of the coordinated nitrate. The ionic nitrate appears at 1384 cm⁻¹ in complex **4**, 1389 cm⁻¹ in complex **5** and 1382 cm⁻¹ in complex **6**. [52-53] The frequencies observed around 404 cm⁻¹, 402 cm⁻¹ and 401 cm⁻¹ in complexes **4**, **5** and **6** respectively are due to $\nu(\text{La-O})$ [36] and the stretching bands $\nu(\text{Cu-N})$ at 494 cm⁻¹ in complex **5** and $\nu(\text{Ni-N})$ 491 cm⁻¹ in complex **6** confirms the coordination of oxamide through amide nitrogen with M(II) ion.

UV-Visible absorption spectra reveal the $n \rightarrow \pi^*$ transitions of the oxamide -C=O at 234.95 nm, 232.58 nm and 233.30 nm in complexes **4**, **5** and **6** respectively. The $\pi \rightarrow \pi^*$ transitions absorb at 262.8 nm, 260.08 nm and 260.08 nm in complexes **4**, **5** and **6** respectively. The $n \rightarrow \pi^*$ transitions in complexes have been red shifted when compared with the ligand 2PMO (229.53) and the $\pi \rightarrow \pi^*$ transitions show hyperchromism on complexation.

ESI-Mass spectra confirm the formation of ionic complexes. The *m/z* value that appeared for complex **4** at 583.05 in negative scan mode corresponds to [M+OH+CH₃OH-(NO₃)⁻] where M = [La(2PMO)(NO₃)₂]⁺ ion. Complex **5** has the *m/z* value at 425.11 in positive ion mode. This value corresponds to [M+Na+CH₃OH-2(NO₃)⁻] where M = [Cu(2PMO)La(phen)(NO₃)₂]²⁺ ion. Complex **6** has *m/z* value at 391.20 which is corresponding to [M+K-2(NO₃)⁻] where M = [Ni(2PMO)La(phen)(NO₃)₂]²⁺ ion.

The electrochemical responses of newly synthesized complexes in DMF were determined by using 0.1 M TBAP as supporting electrolyte at the scan rate of 200 mVs⁻¹. Complexes **4-6** have shown the change in reduction and oxidation peak potentials which is an indicative of the redox behavior of the complexes. [30] The formal potential values of new complexes **4-6** are given in Table 1.

DNA binding experiments

UV-Visible absorption titrations

Electronic absorption titrations were carried out to study the DNA binding propensities of the complexes **2-6** to the calf-thymus (CT) DNA in 5 mM Tris-HCl/50 mM NaCl buffer (pH = 7.2). The changes in the absorption intensities of the spectral bands of the complexes as a function of increased concentration of CT-DNA were determined (Fig. 1). The intrinsic binding constants (*K_b*) of the complexes to the CT-DNA were determined by monitoring changes in the spectral band with increasing CT-DNA concentration, keeping complex concentration constant (30 μM). The binding constant values were given in Table 2.

Table 2: UV-Visible absorption titration data of 30 μM complexes **2-6** in the presence of CT-DNA.

Complex	Empirical formula	<i>K_b</i> (M ⁻¹)	[s]
2	[Cu(2PMO)]	$4.69 \times 10^4 \pm 0.76$	0.21 ± 0.0059
3	[Ni(2PMO)]	$3.12 \times 10^4 \pm 0.9$	0.20 ± 0.0053
4	[La(2PMO)(NO ₃) ₂](NO ₃)	$8.53 \times 10^4 \pm 0.2$	0.23 ± 0.0046
5	[Cu(2PMO)La(phen)(NO ₃) ₂](NO ₃)	$6.35 \times 10^5 \pm 0.1$	0.24 ± 0.0033
6	[Ni(2PMO)La(phen)(NO ₃) ₂](NO ₃)	$4.93 \times 10^5 \pm 0.4$	0.22 ± 0.0067

K_b= intrinsic binding constant. [s]= site size in base pairs.

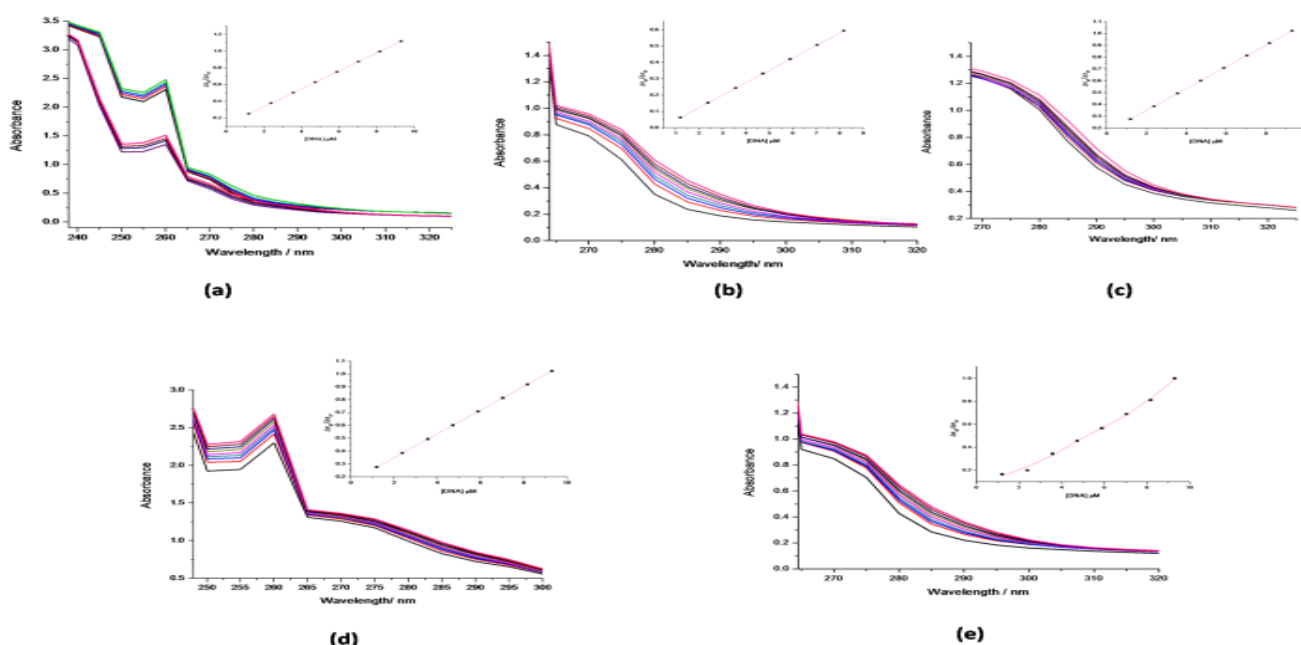


Fig. 1: UV-Visible absorption titration spectra (a-e) of complexes **2-6** (30 μM) respectively in the presence of increased amount of CT-DNA in 5 mM Tris-HCl/ 50 mM NaCl buffer (pH=7.2). The inset plots show $\Delta\epsilon_{at}/\Delta\epsilon_{bf}$ vs. [DNA]/ μM that used to calculate intrinsic binding constant *K_b* M⁻¹.

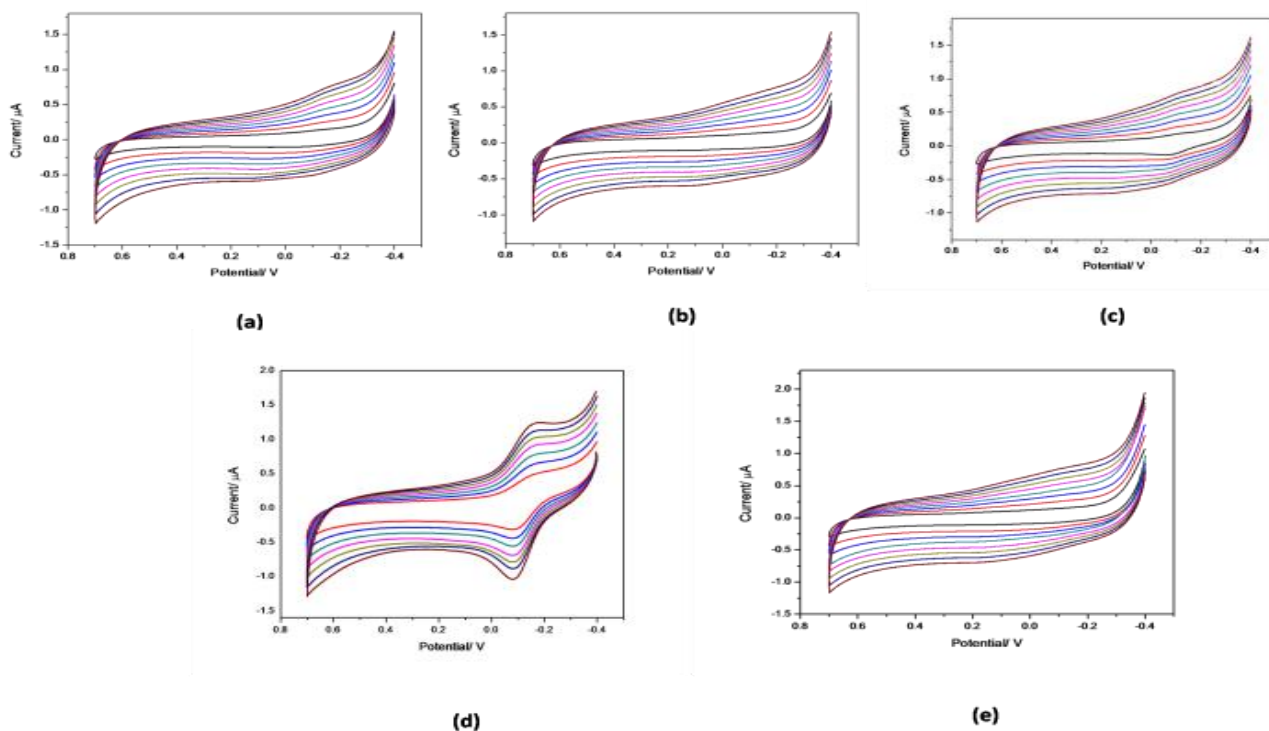


Fig. 2: Cyclic voltammograms of 2-6 (a-e) (30 μM) in the presence of CT-DNA (100 μM) in 5 mM Tris-HCl buffer (pH = 7.2) using 50 mM NaCl as supporting electrolyte in the potential range -0.4 to 0.7 V at the variable scan rate 25 – 200 mVs^{-1} .

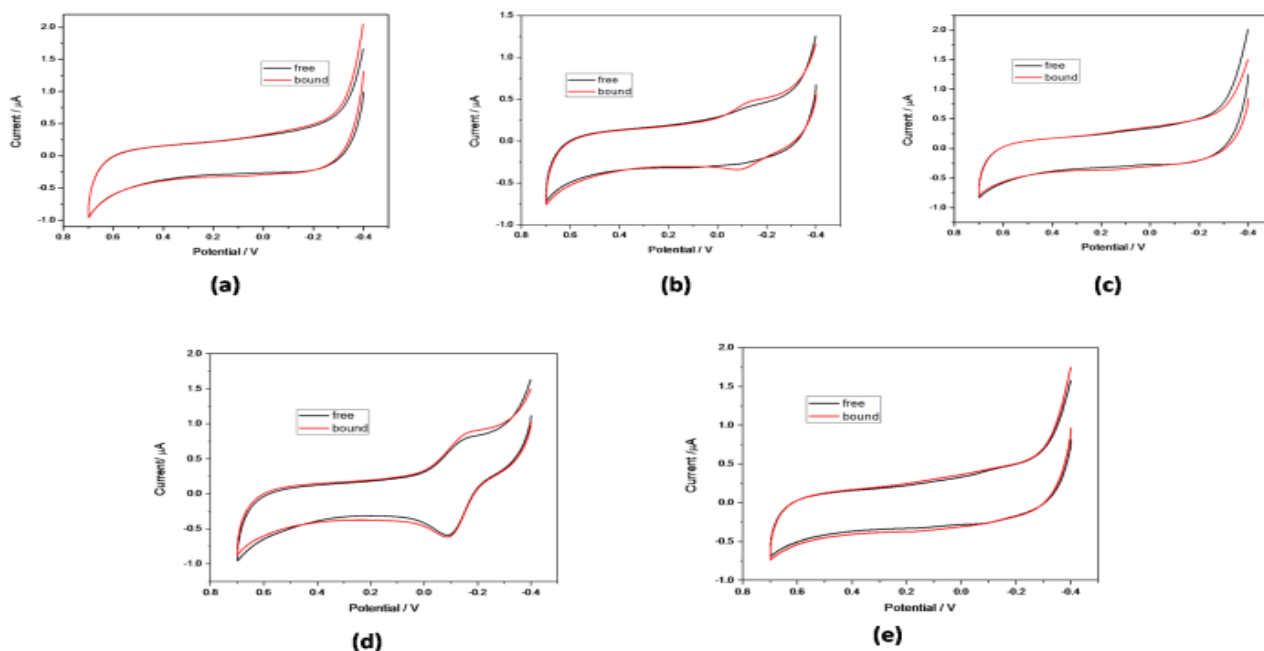


Fig. 3: Cyclic voltammograms of 2-6 (a-e) in free complex and complex fully bound with CT-DNA in 50 mM Tris-HCl buffer (pH = 7.2) using 50 mM NaCl as supporting electrolyte in the potential range -0.4 to 0.7 V at the scan rate 200 mVs^{-1} at room temperature.

From the Fig. 1 it is observed that, on incremental additions of CT-DNA to the complex solutions, the absorption intensity decreases. The significant hypochromism and slight red shift are the characteristic spectral changes in the intercalative mode. This may be due to the coupling of π^* -orbital of the intercalated complex with that of π -orbital of the base pairs of DNA results in decreasing the π - π^* transition energy and hence bathochromic shift and hypochromism. [54-55] The binding strengths of the complexes follow the order: **5** > **6** > **4** > **2** > **3**. Among all the complexes

[Cu(2PMO)La(phen)(NO₃)](NO₃)₂ (**5**) has shown the highest binding constant values.

Electrochemical titrations – Cyclic voltammetry

The mode of binding and the nature of complexes in binding towards CT-DNA are further confirmed by electrochemical titrations. The cyclic voltammetry is carried out in two ways. Firstly, the concentration of complexes **2-6** (30 μM) and the concentration of CT-DNA (100 μM) both are kept constant while varying the scan rate from 25 to 200 mV s^{-1} . Secondly, at the fixed scan rate 200 mV s^{-1} , electrochemical titrations were

carried out by keeping concentration of complex constant (30 μM) and the DNA was added in regular aliquots. The formal peak potential values in the absence (E_f^0) and presence of CT-DNA (E_b^0) were determined and changes in the voltammetric current were observed. The E_f^0 and E_b^0 values are given in the **Table 3**. The changes in the redox behavior of the complexes are shown in **Figs. 2 and 3**.

From the **Table 3** it is observed that, the peak potential values are shifted to positive potential and the voltammetric current decreases on addition of complexes to CT-DNA solution. The reason may be attributed to the fact that, once the complexes are bound to DNA through intercalation, the molecules diffuse more slowly into the electrolyte solution and hence voltammetric current decreases. The E_f^0 and E_b^0 values are used to calculate the ratio of binding constants of complexes in reduced state to the oxidized state and the results reveal that, all the complexes bind to CT-DNA in the reduced state.

Viscosity Method

The binding mode of the complexes to CT-DNA was further confirmed by viscosity method. Viscosity method is known to be the least ambiguous technique to determine the binding interaction of complexes in the absence of crystallographic data of the complexes. It is well known fact that, when crystallographic data is unavailable, viscosity method is the least ambiguous hydrodynamic method that is used to determine the binding mode of metal complexes with DNA. A classical intercalator intercalates between the base pairs of the DNA, thus increases the length of DNA helix and hence viscosity of DNA increases whereas partial or non-classical interaction of complex results in bending or kinking of DNA, thereby decreasing the length of the DNA and in turn decreases the viscosity of DNA. [41, 46, 55] The concentration dependent viscosity method reveals that, as the concentration of the complexes increases, viscosity of the DNA increases. The viscosity is highest for binuclear complexes (5 and 6) than their mononuclear analogues (2-4). This is due to the fact that the planar aromatic ring of 1,10-phenanthroline stabilizes the DNA-complex interaction through intercalation and hence the viscosity is increased. The change in the viscosity of CT-DNA in presence of complexes has been shown in **Fig. 4**.

DNA cleavage studies

The ability of the complexes 2-6 in comparison with the ligand 2PMO (1) to cleave super coiled (SC) pUC19 DNA in the absence and presence of oxidizing agent H_2O_2 and reducing agent MPA was studied by agarose gel Electrophoresis in 50 mM Tris-HCl buffer/50 mM NaCl (pH = 7.2). Of all the complexes and ligands studied, at 10 μM , complexes 5 and 6 have shown the maximum cleavage (92% and 88% respectively) in presence of reducing agent MPA as shown in **Fig. 5**. The SC pUC 19 DNA cleavage data in hydrolytic and oxidative conditions are given in **Table 4**. The

mononuclear complexes 2-4 have shown poor chemical nuclease activity at the same concentration.

Table 3: Cyclic voltammetric data of complexes 2-6 in free form and fully bound with CT-DNA.

Complex	E_f (V)	I_c (μA)	E_b (V)	I_a (μA)	E_f^0 (V)	E_b^0 (V)	K_R/K_O
2	-0.244	-1.42	-0.204	-2.78	-0.122	-0.102	2.17
3	-0.101	-1.26	-0.084	-1.62	-0.050	0.042	1.43
4	0.0798	-2.08	0.1368	-3.19	0.0399	0.0684	3.02
5	-0.081	-1.57	-0.068	-3.55	-0.041	-0.034	1.28
6	-0.088	-4.65	-0.084	-4.82	-0.044	-0.042	1.08

E_f = Electrode potential of free complex, E_b = Electrode potential of complex bound to DNA.

Table 4: SC pUC 19 DNA cleavage data of 2PMO (1) and its complexes 2-6.

Lanes	Reaction Conditions	% SC	% NC
1	DNA + NaCl + Tris-HCl	99	01
2	DNA + NaCl + Tris-HCl + H_2O_2	98	02
3	DNA + NaCl + Tris-HCl + MPA	97	03
4	DNA + NaCl + Tris-HCl + 1	98	02
5	DNA + NaCl + Tris-HCl + 1 + H_2O_2	98	02
6	DNA + NaCl + Tris-HCl + 1 + MPA	97	03
7	DNA + NaCl + Tris-HCl + 2	94	06
8	DNA + NaCl + Tris-HCl + 2 + H_2O_2	92	08
9	DNA + NaCl + Tris-HCl + 2 + MPA	70	30
10	DNA + NaCl + Tris-HCl + 3	97	03
11	DNA + NaCl + Tris-HCl + 3 + H_2O_2	98	02
12	DNA + NaCl + Tris-HCl + 3 + MPA	80	20
13	DNA + NaCl + Tris-HCl + 4	97	03
14	DNA + NaCl + Tris-HCl + 4 + H_2O_2	94	06
15	DNA + NaCl + Tris-HCl + 4 + MPA	84	16
16	DNA + NaCl + Tris-HCl + 5	97	03
17	DNA + NaCl + Tris-HCl + 5 + H_2O_2	87	13
18	DNA + NaCl + Tris-HCl + 5 + MPA	08	92
19	DNA + NaCl + Tris-HCl + 6	94	06
20	DNA + NaCl + Tris-HCl + 6 + H_2O_2	89	11
21	DNA + NaCl + Tris-HCl + 6 + MPA	12	88

[Complex] = 10 μM ; [H_2O_2] = 200 μM ; [MPA] = 500 μM ; Tris-HCl/NaCl = 50 mM.

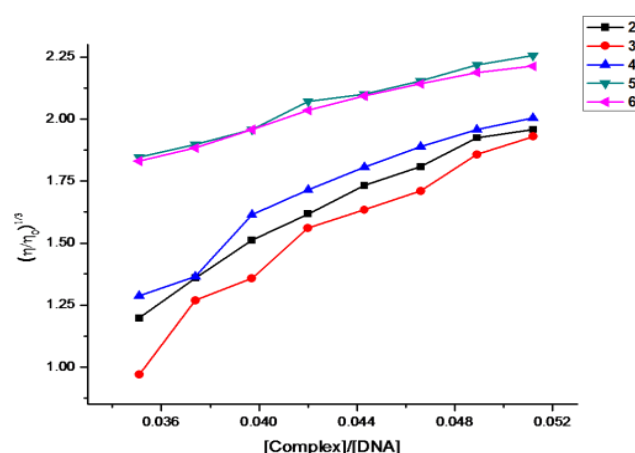


Fig. 4: A plot of $[\text{Complex}]/[\text{DNA}]$ vs. $(\eta/\eta_0)^{1/3}$ showing the change in viscosity of CT-DNA with increasing concentration of complexes 2-6 in the presence of 50 mM Tris-HCl/NaCl buffer (pH = 7.2) at 37°C.

The oxidative cleavage is mediated by the generation of diffusible free radicals by metal complexes. The involvement of highly reactive oxygen species such as hydroxyl radicals or singlet oxygen in cleaving SC pUC 19 DNA by complexes 5 and 6 were determined by the mechanistic studies carried out by agarose gel electrophoresis in presence of reducing agent MPA

using radical quenchers DMS and sodium azide (NaN_3) and major groove binder methyl green. No significant inhibition of cleavage is observed in presence of singlet oxygen quencher NaN_3 whereas cleavage activity is inhibited in presence of hydroxyl radical quencher DMSO and major groove binder MG. This confirms that the complexes 5 and 6 intercalate between the DNA base pairs in the major groove direction. The results are shown in Fig. 6 and the cleavage data is given in Table 5.

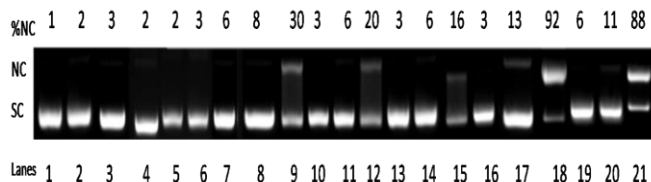


Fig. 5: Agarose gel electrophoretic diagram showing the cleavage of SC pUC 19 DNA by 2PMO (1) and its complexes 2-6 (10 μM) in presence of 50 mM Tris-HCl/ NaCl buffer (pH = 7.2), oxidizing agent H_2O_2 and reducing agent MPA. Lane 1, DNA alone; lane 2, DNA + H_2O_2 ; lane 3, DNA + MPA, lane 4, DNA + 1; lane 5, DNA + H_2O_2 + 1; lane 6, DNA + MPA + 1; lane 7, DNA + 2; lane 8, DNA + H_2O_2 + 2; lane 9, DNA + MPA + 2; lane 10, DNA + 3; lane 11, DNA + H_2O_2 + 3; lane 12, DNA + MPA + 3; lane 13, DNA + 4; lane 14, DNA + H_2O_2 + 4; lane 15, DNA + MPA + 4; lane 16, DNA + 5; lane 17, DNA + H_2O_2 + 5; lane 18, DNA + MPA + 5; lane 19, DNA + 6; lane 20, DNA + H_2O_2 + 6; lane 21, DNA + MPA + 6.

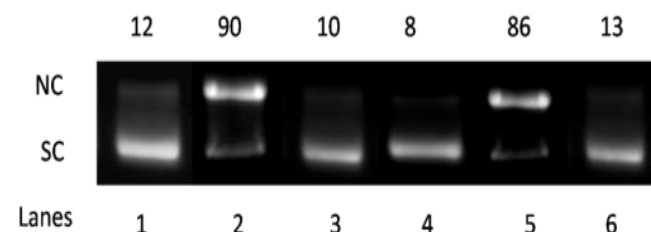


Fig. 6: Agarose gel electrophoretic diagram showing the cleavage of SC pUC 19 DNA by complexes 5 and 6 in the presence of additives like DMSO, NaN_3 and methyl green (MG) in 50 mM Tris-HCl/NaCl buffer (pH = 7.2). Lane 1, DNA + MPA + 5 + DMSO; lane 2, DNA + MPA + 5 + NaN_3 ; lane 3, DNA + MPA + 5 + MG; lane 4, DNA + MPA + 6 + DMSO; lane 5, DNA + MPA + 6 + NaN_3 ; lane 6, DNA + MPA + 6 + MG.

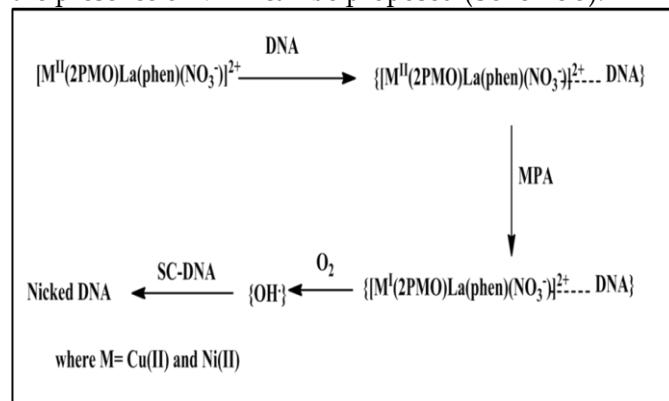
Table 5: SC pUC 19 DNA cleavage data of mechanistic studies of 5 and 6 in presence of DMSO, NaN_3 or MG at 10 μM .

Lane no.	Reaction Conditions	%SC	%NC
1	DNA + NaCl + Tris-HCl + MPA + 5 + DMSO	88	12
2	DNA + NaCl + Tris-HCl + MPA + 5 + NaN_3	10	90
3	DNA + NaCl + Tris-HCl + MPA + 5 + MG	90	10
4	DNA + NaCl + Tris-HCl + MPA + 6 + DMSO	92	8
5	DNA + NaCl + Tris-HCl + MPA + 6 + NaN_3	14	86
6	DNA + NaCl + Tris-HCl + MPA + 6 + MG	87	13

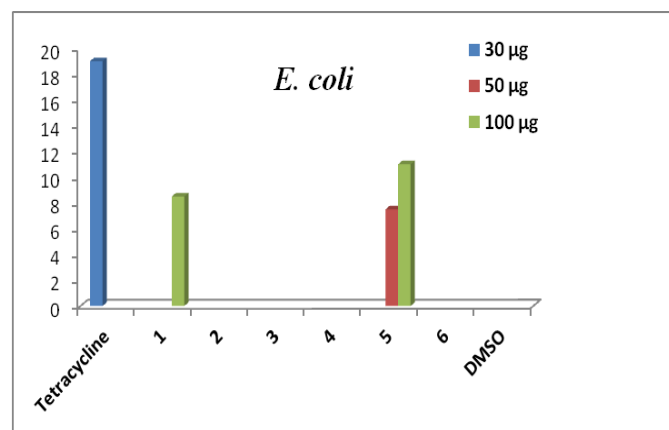
[MPA] = 500 μM , DMSO = 4 μL , [NaN_3] = 200 μM , Methyl green (MG) = 10 μM , [Complex] = 10 μM .

Thus it can be proposed that in presence of MPA, Cu(II) reduces to Cu(I) and Ni(II) to Ni(I) and in turn activate the molecular oxygen into hydroxyl radical which is responsible for the cleavage of SC pUC 19 DNA. [57]

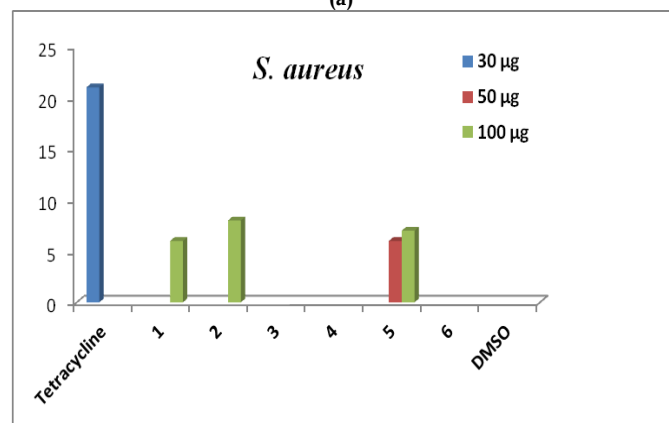
Based on the above observation, a mechanism for the cleavage of SC pUC19 DNA by complexes 5 and 6 in the presence of MPA can be proposed (Scheme 3).



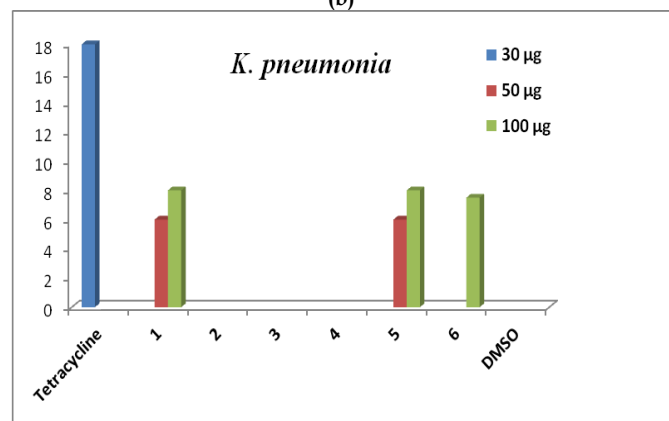
Scheme 3. Proposed mechanism for the cleavage of SC pUC 19 DNA by complexes 5 and 6 in the presence of MPA mediated by hydroxyl radicals.



(a)



(b)



(c)