ORIGINAL ARTICLE



Synthesis, Structural, DNA Binding and Cleavage Studies of Cu(II) Complexes Containing Benzothiazole Cored Schiff Bases

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Received: 15 June 2016/Accepted: 15 August 2016/Published online: 27 August 2016 © Springer Science+Business Media New York 2016

Abstract Novel benzothiazole Schiff bases L^1 [1-((4,6difluorobenzo[d]thiazol-2-ylimino)methyl) naphthalen-2-ol], L^{2} [3-((4,6-difluorobenzo[d]thiazol-2-ylimino) methyl)benzene-1,2-diol], L^{3} [2-((4,6difluorobenzo[d]thiazol-2-ylimino)methyl)-5methoxyphenol], L⁴ [2-((4,6-difluorobenzo[d]thiazol-2vlimino)methyl)-4-chlorophenol] and their binary Cu(II) complexes were synthesized. The structures of all the compounds have been discussed on the basis of elemental analysis, FT-IR, NMR, UV-Visible, ESI-Mass, TGA, ESR, SEM, powder XRD and magnetic moments. Based on the analytical and spectral data a square planar geometry has been assigned to all complexes in which the Schiff bases act as monobasic bidentate ligands, coordinating through the azomethine nitrogen and phenolic oxygen atom. DNA binding ability of these complexes was studied on CT-DNA by using UV-Vis absorption, fluorescence and viscometry. DNA cleavage ability of the complexes was examined on pBR322 DNA by using gel electrophoresis method. All the DNA binding studies reveal that they are good intercalators. The bioefficacy of the ligands and their complexes was examined against the growth of bacteria and fungi in vitro to evaluate their antimicrobial potential. The screening data revealed that the complexes showed more antimicrobial activity than the corresponding free ligands.

Electronic supplementary material The online version of this article (doi:10.1007/s10895-016-1911-3) contains supplementary material, which is available to authorized users.

Shivaraj shivaraj_sunny@yahoo.co.in Keywords Schiff base \cdot Cu(II) complex \cdot DNA binding \cdot DNA cleavage \cdot Antimicrobial activity

Introduction

DNA is a significant cellular receptor, many chemicals bring to bear their antitumor effects by binding to DNA and by this means change the replication of DNA and inhibit the growth of the cancer cells, which is the basis of preparing new and more efficient antitumor drugs [1-3]. During the last decade several transition metal complexes have been used as tools for understanding DNA structure, as agents for mediation of DNA cleavage or as chemotherapeutic agents. Benzothiazole nucleus possesses broad range of biological properties such as antitumour [4, 5], cytotoxic [6], antiglutamate/antiparkinson [7], inhibition of enzymes [8]. Moreover, Schiff bases derived from 2-aminobenzothiazole are of great importance due to their wide applications as antimicrobial, anti-inflammatory, antidegenerative and anti-HIV agents [9-13]. There is an increasing prominence and evergreen curiosity in the field of metal containing drugs. Medicinal inorganic chemistry covering a wide applications of metals in therapeutics and diagnostics. The biological behaviour of Cu(II) complexes has been subjected to intense investigation for DNA binding and cleavage activities [14] for novel chemotherapeutics and highly sensitive diagnostic agents [15]. Metal complexes can bind to DNA via covalent or non-covalent type of binding (intercalating, groove or electrostatic). Copper has a long history of medicinal application, the anticancer and antiparasitic properties of Cu(II) complexes have increased the interest of scientists in their thorough studies [16, 17]. Synthesis, characterization, biological activity and DNA interaction of Cu(II) complexes of isoxazole Schiff bases were reported earlier from our laboratory [18, 19].

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The aim of the present study is to synthesize the desired Cu(II) complexes from benzothiazole cored Schiff bases and to study the DNA binding and cleavage efficiency. Further, to investigate their effect on pathogenic strains of bacteria and fungi.

Experimental

Materials

All the chemicals and solvents used were purchased from Sigma-Aldrich, Merck and Hi media Ltd. All the solvents such as methanol, chloroform, petroleum ether, acetone and water were purified by standard procedures. The pBR322 DNA and CT-DNA were purchased from Genei Biolab, Bangalore, India and stored at 4 °C. All the buffer solutions used in the DNA interaction studies were prepared from doubly distilled water.

Instrumentation

The elemental analysis of Schiff bases and their Cu(II) complexes was carried out by using Perkin Elmer 240C (USA) elemental analyzer. ¹H-NMR and ¹³C-NMR spectra of the Schiff bases were recorded on Bruker 400 MHz NMR instrument using TMS as internal reference. FT-IR spectra of the compounds were recorded in the range 4000–400 cm^{-1} on Perkin-Elmer Infrared model 337 using KBr pellets. Electronic spectra were recorded on Shimadzu UV-VIS 1601 spectrophotometer. ESI mass spectra of all compounds were recorded on VG AUTOSPEC mass spectrometer. Magnetic susceptibilities of Cu(II) complexes were measured on Gouy balance model 7550 using Hg[Co(SCN)₄] as standard. Melting temperatures of the ligands and their complexes were determined on Polmon instrument (Model No. MP-96). Copper content of the complexes was estimated by atomic absorption spectroscopy after decomposing the complexes with concentrated HNO₃ using GBC Avanta 1.0 AAS. The ESR spectra were recorded at liquid nitrogen temperature on JES-FA200 ESR spectrometer (JEOL-Japan). The thermo gravimetric analysis was carried out in a dynamic nitrogen atmosphere with a heating rate of 10 °C min⁻¹ using a Shimadzu TGA-50H in the temperature range of 27-1000 °C. Viscosity measurements were performed using Ostwald viscometer (Vensil). Powder X-ray diffraction (XRD) analysis was recorded on X'pert Pro diffractometer. The morphology of ligands and complexes was measured by SEM (scanning electron microscopy, Zeiss evo18).

General Procedure for the Synthesis of Schiff Bases

Hot methanolic solution of 4,6-difluorobenzo[d]thiazol-2amine (10 mmol) was magnetically stirred in a round bottom flask followed by drop wise addition of 2-hydroxy naphthalene-1-carbaldehyde / 2,3-dihydroxy benzaldehyde / 2-hydroxy-4-methoxy benzaldehyde / 5-chloro-2-hydroxy benzaldehyde (10 mmol). The reaction mixture was then refluxed with stirring at 60–80 °C for 2 h. After cooling, the yellow solid precipitate was collected by filtration. The resultant product was washed and recrystallized with methanol. The Schiff bases were dried in vacuum desiccators over anhydrous CaCl₂. The purity of the ligands was checked by TLC.

Ligand L^1 : Yield 78 %. **M. P**: 175 °C. **Mol Wt**: 340. *Anal.* Calc. (C₁₈H₁₀F₂N₂OS) (%): C, 63.52; H, 2.96; N, 8.23. Found: C, 63.58; H, 3.02; N, 8.26. **IR (KBr)** (cm⁻¹): $v_{(OH)}$ 3468; $v_{(CH=N)}$ 1623; $v_{(C-O)}$ 1219. **UV (CHCl₃)** $\lambda_{max/nm(cm^{-1})}$: 385 (25974); 261 (38314). ¹H NMR **(400 MHz, CDCl₃)**: δ = 14.29 (s, 1H), 10.18 (s, 1H), 8.30 (d, J = 8.5 Hz, 1H), 7.95 (d, J = 9.0 Hz, 1H), 7.78 (d, J = 8.0 Hz, 1H), 7.65–7.60 (m, 1H), 7.45–7.41 (m, 1H), 7.37–7.34 (m, 1H), 7.12 (d, J = 9.0 Hz, 1H), 7.05–6.99 (m, 1H), Fig. S1. ¹³C NMR (100 MHz, CDCl₃): δ = 176.1, 160.5, 159.8, 158.4, 157.9, 151.0, 139.4, 138.2, 136.8, 133.6, 129.1, 127.3, 125.0, 123.7, 117.3, 108.1, 104.2, 100.2. MS (ESI): m/z = 341 [M + H]⁺.

Ligand L^2 : Yield 75 %. **M. P**: 168 °C. **Mol Wt**: 306. *Anal.* Calc. (C₁₄H₈F₂N₂O₂S) (%): C, 58.26; H, 2.44; N, 7.50. Found: C, 59.02; H, 2.50; N, 7.53. **IR (KBr)** (cm⁻¹): $v_{(OH)}$ 3468; $v_{(CH=N)}$ 1633; $v_{(C-O)}$ 1247. **UV (CHCl₃)** λ_{max} ,nm(cm⁻¹): 387 (25839); 322 (31055); 263 (38022). ¹H **NMR (400 MHz, CDCl₃)**: δ = 10.23 (s, 1H), 7.74 (s, 2H), 7.51–7.42 (m, 1H), 7.17–7.09 (m, 3H), 6.84–6.79 (m, 1H). ¹³C **NMR (100 MHz, CDCl₃)**: δ = 192.8, 166.6, 157.2, 155.1, 152.0, 149.7, 146.1, 137.5, 133.3, 122.5, 121.3, 119.5, 103.7, 101.0, Fig. S2. **MS (ESI)**: m/z = 307 [M + H]⁺.

Ligand L^3 : Yield 76 %. **M. P**: 172 °C. **Mol Wt**: 320. *Anal.* Calc. (C₁₅H₁₀F₂N₂O₂S) (%): C, 56.24; H, 3.15; N, 8.73. Found: C, 56.29; H, 3.01; N, 8.71. **IR (KBr)** (cm⁻¹): $v_{(OH)}$ 3317; $v_{(CH=N)}$ 1614; $v_{(C-O)}$ 1247. **UV (CHCl₃)** $\lambda_{max/nm(cm^{-1})}$: 383 (26109); 263 (38022). ¹H NMR **(400 MHz, CDCl₃)**: δ = 9.28 (s, 1H), 7.86 (s, 1H), 7.84 (s, 1H), 7.51–7.41 (m, 3H), 7.15–7.09 (m, 2H), 3.86 (s, 3H). ¹³C **NMR (100 MHz, CDCl₃)**: δ = 170.6, 166.1, 162.9, 157.6, 154.9, 152.3, 149.5, 137.3, 133.6, 113.0, 108.1, 104.4, 102.4, 100.9, 55.6. **MS (ESI)**: m/z = 321 [M + H]⁺.

Ligand L^4 : Yield 70 %. **M. P**: 169 °C. **Mol Wt**: 324. *Anal.* Calc. (C₁₄H₇F₂N₂ClOS) (%): C, 51.78; H, 2.17; N, 8.63. Found: C, 51.59; H, 2.09; N, 8.81. **IR (KBr)** (cm⁻¹): $v_{(OH)}$ 3471; $v_{(CH=N)}$ 1647; $v_{(C-O)}$ 1257. **UV (CHCl₃)** $\lambda_{max/nm(cm^{-1})}$: 333 (30030); 264 (37878). ¹H NMR **(400 MHz, CDCl₃**): δ = 12.06 (s, 1H), 9.27 (s, 1H), 7.50 (d, J = 2.7 Hz, 1H), 7.45–7.41 (m, 1H), 7.39–7.36 (m, 1H), 7.06– 7.01 (m, 2H), Fig. S3. ¹³C NMR (100 MHz, CDCl₃): δ = 174.8, 161.9, 130.5, 159.1, 157.2, 139.8, 135.6, 132.4, 130.9, 125.7, 120.9, 118.3, 104.5, 102.2. **MS (ESI)**: m/z = 325 [M + H]⁺.

Synthesis of Binary Cu(II) Complexes

The preparation of all Cu(II) complexes were described in Scheme 1.

The complexes $[Cu(L^1)_2]$, $[Cu(L^2)_2]$, $[Cu(L^3)_2] \& [Cu(L^4)_2]$ were prepared by using the following general procedure. Hot methanolic solution (10 mL) of copper acetate monohydrate (0.5 mmol) was added drop wise to a hot methanolic solution (10 mL) of the corresponding Schiff bases (1 mmol). After complete addition, the mixture was refluxed with stirring at 60–80 °C for 3 h. The solid precipitate was separated out upon cooling. The precipitate was filtered under suction and washed with methanol and petroleum ether, finally kept for drying in vacuum desiccators over anhydrous CaCl₂.

Scheme 1 Synthesis of Schiff bases and their Cu(II) complexes

[*Cu*(*L*¹)₂]: Yield 70 %. **M. P**: 240 °C. **Mol Wt**: 742. *Anal.* Calc. (C₃₆H₁₈CuF₄N₄O₂S₂) (%): C, 58.26; H, 2.44; N, 7.50; Cu, 8.56. Found: C, 58.22; H, 2.50; N, 7.55; Cu, 8.51. **IR** (**KBr**)(cm⁻¹): $v_{(C=N)}$ 1612, $v_{(C-O)}$ 1238, $v_{(M-O)}$ 589, $v_{(M-N)}$ 429. **ESR**: g_{||} = 2.2778, g_⊥ = 2.0594, G = 4.6767. **UV-Vis (DMSO)** $\lambda_{max}/nm(cm^{-1})$: 257 (38610), 339 (23498), 381 (26246), 525 (19047). μ_{eff} (BM):1.83. **MS (ESI)**: 765 [M + Na]⁺.

[$Cu(L^2)_2$]: Yield 72 %. **M. P**: 235 °C. **Mol Wt**: 673. Anal. Calc. ($C_{28}H_{14}CuF_4N_4O_4S_2$) (%): C, 49.89; H, 2.09; N, 8.13; Cu, 9.43. Found: C, 49.95; H, 2.15; N, 8.19; Cu, 9.48. **IR** (**KBr**)(cm⁻¹): $v_{(C=N)}$ 1622, $v_{(C-O)}$ 1257, $v_{(M-O)}$ 587, $v_{(M-N)}$ 441. **ESR**: $g_{\parallel} = 2.3792$, $g_{\perp} = 2.0700$, G = 5.4171. **UV-Vis (DMSO)** λ_{max} /nm(cm⁻¹): 260 (38461), 342 (29239), 385 (25974), 520 (19230). μ_{eff} (BM):1.84. **MS (ESI)**: 691 [M + NH₄]⁺.



 $[Cu(L^2)_2], [Cu(L^3)_2], [Cu(L^4)_2]$

 $[Cu(L^1)_2]$

[$Cu(L^3)_2$]: Yield 69 %. **M. P**: 230 °C. **Mol Wt**: 701. Anal. Calc. ($C_{30}H_{18}CuF_4N_4O_4S_2$) (%): C, 52.37; H, 2.20; N, 8.73; Cu, 9.05. Found: C, 52.43; H, 2.26; N, 8.81; Cu, 9.10. **IR** (**KBr**)(cm⁻¹): $v_{(C=N)}$ 1609, $v_{(C-O)}$ 1256, $v_{(M-O)}$ 550, $v_{(M-N)}$ 429. **ESR**: $g_{\parallel} = 2.3689$, $g_{\perp} = 2.0658$, G = 5.6063. **UV-Vis** (**DMSO**) $\lambda_{max}/nm(cm^{-1})$: 259 (38610), 314 (31847), 411 (24330). μ_{eff} (BM):1.74. **MS (ESI)**: 624 [M + Na]⁺.

 $[Cu(L^4)_2]$: Yield 74 %. **M. P**: 247 °C. **Mol Wt**: 708. *Anal.* Calc. (C₂₈H₁₂CuF₄N₄O₂S₂Cl)(%): C, 47.30; H, 1.70; N, 7.88; Cu, 9.97. Found: C, 47.33; H, 1.66; N, 7.81; Cu, 9.91. **IR (KBr)**(cm⁻¹): $v_{(C=N)}$ 1635, $v_{(C-O)}$ 1262, $v_{(M-O)}$ 559, $v_{(M-N)}$ 462. **ESR**: $g_{\parallel} = 2.3719$, $g_{\perp} = 2.0613$, G = 6.0668. **UV-Vis (DMSO)** λ_{max} /nm(cm⁻¹): 259 (38610), 339 (29498), 383 (26109), 517 (19342). μ_{eff} (BM):1.77. **MS (ESI)**: 726 [M + NH₄]⁺.

DNA Binding Studies

UV–Vis Spectroscopic Studies

The UV-Visible absorption spectroscopic method is used to study the DNA binding nature of metal complexes. The absorption titrations were carried out in Tris–HCl buffer (5 mM Tris– HCl/50 mM NaCl, pH 7.2) at 25 °C by keeping the Cu(II) complex concentration constant (10 μ M) and varying the concentration of the CT-DNA from 0 to 10 μ M. The solution of CT-DNA in the buffer gave a ratio of UV absorbance of 1.8: 1.9 at 260 nm and 280 nm, which indicates the CT-DNA was sufficiently free of protein [20]. The CT-DNA concentration per nucleotide was measured by employing an extinction coefficient of 6600 M⁻¹ cm⁻¹ at 260 nm [21]. To eliminate the absorbance of CT-DNA itself, equal increments of CT-DNA were added to both the sample solution and the reference solution.

Fluorescence Study

By the fluorescence spectral method, the relative binding of the complexes to CT-DNA was studied with an EB-bound CT-DNA solution in Tris-HCl buffer (pH 7.2) at 25 °C were excited at 350 nm by varying concentrations of complexes from 0 to 60 μ M. The emission intensity showed reduction upon addition of the complex. The effect of the metal complexes on the emission intensity of EB-bound CT-DNA was used to determine the binding properties of the metal complexes. The binding constants were calculated from the slopes of the lines in the plot of the fluorescence intensity versus complex concentration. The relative binding of the complexes to CT-DNA was determined by Stern–Volmer eq. $I_0/I = 1 + K_{SV} r$.

Where I_0 and I are the fluorescence intensities in the absence and presence of complexes respectively, K_{SV} is a linear Stern–Volmer constant, and r is the concentration of complex to that of DNA.

Viscosity Measurements

Ostwald capillary viscometer was used to study the viscosities of CT-DNA at temperature 30 ± 0.1 °C in a thermostatic water bath. CT- DNA concentration (100 µM) was kept constant and complex concentrations (0–100 µM) were varied in Tris–HCl buffer (pH 7.2) solution. The flow time of the solution was measured by a digital stop-watch. An average flow time was taken after three times measurement for the same complex solution to maintain the accuracy. A plot was made between $(\eta/\eta_o)^{1/3}$ and the ratio of the concentration of the Cu(II) complexes to CT-DNA, where η is the viscosity of CT-DNA alone. Viscosity values were calculated from the experimental flow time of CT-DNA containing solutions corrected from the flow time of buffer alone.

DNA Cleavage Experiments

DNA cleavage experiments were performed with pBR322 DNA at pH 7.2 in Tris–HCl buffer solution by agarose-gel electrophoresis method by varying concentration of Cu(II) complexes. Oxidative DNA cleavage was monitored by treating pBR322 DNA with H₂O₂ and photolytic DNA cleavage in presence of UV light. The loading dye 0.25 % bromophenol blue is added to the samples and made to a total volume of 16 μ L. The samples were incubated at 37 °C for 2 h. The samples were loaded in the agarose gel and electrophoresis was carried out at 75 V for 1 h.

Biological Evaluation

The ligands and their Cu(II)complexes were screened for their in vitro anti bacterial activity against gram negative *Escherichia coli (E. coli), Pseudomonas putida (P. putida)* and *Klebsiella pneumoniae (K. pneumoniae)* and gram positive *Bacillus subtilis (B. subtilis)* and *Staphylococcus aureus (S. aureus),* Ampicillin was used as standard drug. The antifungal activity was screened against *Aspergillus niger (A. niger)* and *Candida albicans (C. albicans)* by using Ketoconazole as standard drug. Nutrient agar was used as the medium for antimicrobial activity [22]. The sterilized medium was inoculated with the suspension of the microorganism and then poured into a petridish. The test compounds (1 mg/mL in DMSO solvent) were impregnated on the paper disc and then placed on the solidified medium. The plates were kept for incubation at 37 °C for 24 h.

Results and Discussion

All Cu(II) complexes are coloured, stable at room temperature and non-hygroscopic and melt at high temperatures. All the

Table 1 Some important IR absorption frequencies (cm^{-1}) of Schiffbases and their Cu(II) complexes

Compound	$\upsilon_{\text{(O-H)}}$	$\upsilon_{(CH=N)}$	$\upsilon_{(C\text{-}O)}$	$\upsilon_{(M\text{-}O)}$	v _(M-N)
L^1	3468	1623	1219	-	-
$[Cu(L^1)_2]$	-	1612	1238	589	429
L ²	3468,	1633	1247	-	-
$[Cu(L^2)_2]$	-	1622	1257	587	441
L ³	3317	1614	1247	-	-
$[Cu(L^3)_2]$	-	1609	1256	550	429
L^4	3471	1642	1257	-	-
$[Cu(L^4)_2]$	-	1635	1262	559	462

complexes are soluble in DMF and DMSO but insoluble in water, alcohol and chloroform. The analytical data obtained for all complexes are in good agreement with the theoretical values and the metal to ligand ratio is 1:2.

Spectral Characterization

FT-IR Spectra

The IR spectra of free Schiff base ligands were compared with the spectra of their corresponding Cu(II) complexes to understand the coordinating mode of the Schiff base ligands to Cu(II) centre in the complexes. A representative IR spectra of L^3 and $[Cu(L^3)_2]$ are given in Figs. S4 and S5. The Schiff base ligands

Fig. 1 SEM and EDX graphs of L^1 and $[Cu(L^1)_2]$



showed a strong band in the region $1614-1642 \text{ cm}^{-1}$ which corresponds to the azomethine (C = N) group. Further, this band is shifted to a lower frequency region 1609–1635 cm⁻¹ indicating co-ordination of azomethine nitrogen to Cu(II) ion [23–25]. A strong intense band observed at 1219–1257 cm⁻¹ in Schiff bases due to phenolic C-O stretching is shifted to higher frequency by 1238–1262 cm⁻¹ indicating Schiff bases are coordinated through the phenolic oxygen to the metal atom. A broad band appeared in the range of 3317-3471 cm⁻¹ attributed to vOH in the Schiff bases, which is disappeared in complexes indicating deprotonation of the phenolic proton prior to coordination. The co-ordination of the azomethine nitrogen and phenolic oxygen are further supported by the appearance of two new non ligand bands at 520–589 and 429–462 cm^{-1} due to vM-O and vM-N respectively [26, 27]. The IR absorption frequencies are given in Table 1.

Electronic Spectra and Magnetic Moments

The magnetic moment values (μ_{eff}) of all Cu(II) complexes are in the range of 1.74–1.84 BM which reveals the presence of a single unpaired electron [28, 29]. The Schiff base ligands exhibited characteristic high energy absorption bands in the range of 260–263 nm and 314–344 nm which are attributed to intraligand π - π * transitions. Other low energy bands observed at 381–387 nm are due to n- π * transition of non bonding electrons of azomethine nitrogen of Schiff base ligands [30, 31]. The Cu(II) complexes show a broad band in the lower frequency region 411–525 nm corresponds to the d-d



SEM photograph of L¹



EDX graph of L¹



SEM photograph of [Cu(L¹)₂]



EDX graph of [Cu(L¹)₂]



Fig. 2 Powder XRD patterns of L^1 and $[Cu(L^1)_2]$

transition due to ${}^{2}B_{1g} \rightarrow {}^{2}E_{g}$ transition. A representative electronic spectra of L¹ and [Cu(L¹)₂] are given in Fig. S6. Based on electronic spectral data and magnetic moment values a square planar geometry is assigned to all Cu (II) complexes [32].

SEM

The surface morphology of ligands and their metal complexes was investigated by scanning electron micrograph analysis. All the compounds were analyzed by EDX analysis to find the elements present on surface. SEM photographs and EDX graphs of L¹, L², [Cu(L¹)₂], [Cu(L²)₂] are given in Fig. 1 and Fig. S7. The SEM analysis of ligands L¹, L², L³ and L⁴ showed irregular cubes, rod shaped, irregular tube and elongated flake like structures respectively and the complexes [Cu(L¹)₂], [Cu(L²)₂], [Cu(L³)₂] & [Cu(L⁴)₂] resemble twisted fibre, granular, grass like, irregular particle shaped structures

Fig. 3 ESR spectrum of $[Cu(L^2)_2]$

respectively. The EDX analysis of ligands contains C, H, N, O, S, F and Cu(II) complexes contain C, H, N, O, S, F & Cu elements. The appearance of no other elemental peaks in the spectra assures the purity of the compounds.

Powder XRD

The powder XRD patterns of all the compounds were recorded at $2\theta = 10-80^{\circ}$ range. The powder XRD patterns of L¹, L³, [Cu(L¹)₂], [Cu(L³)₂] are shown in Fig. 2 and Fig. S8, display sharp crystalline peaks for the ligands indicating their crystalline nature, whereas the complexes do not exhibit well defined sharp peaks possibly due to their amorphous nature. The crystallite sizes are calculated from XRD peak by using the Debye-Scherrer's equation (D = $0.9 \lambda / \beta \cos\theta$). The crystallite sizes calculated for ligands are 26 nm (L¹), 17 nm (L²), 26 nm (L³), 14 nm (L⁴) and for complexes are 17 nm [Cu(L¹)₂], 23 nm [Cu(L²)₂], 8 nm [Cu(L³)₂], 17 nm [Cu(L⁴)₂].



Table 2 ESR data ofCu(II) complexes

Complex	g	g_{\perp}	G
$[Cu(L^{1})_{2}]$	2.2778	2.0594	4.6767
$[Cu(L^{2})_{2}]$	2.2600	2.0650	4.2695
$[Cu(L^3)_2]$	2.2508	2.0608	4.3067
$[Cu(L^4)_2]$	2.2398	2.0613	4.0250

ESR Spectra

The liquid state X band ESR spectra of all the Cu(II) complexes were recorded at liquid nitrogen temperature (77 K). The ESR spectra of Cu(II) complexes provide the information about the extent of the delocalization of unpaired electron. The ESR spectra of $[Cu(L^1)_2]$, $[Cu(L^2)_2]$ are shown in Fig. S9 and Fig. 3. The g_{\parallel} , g_{\perp} and G values are given in Table 2. The trend observed $g_{\parallel} > g_{\perp} > g_e(2.0023)$ is a typical of Cu(II) d⁹ system with axial symmetry and the unpaired electron is localized in $d_{x}^{2} g^{-2}$ orbital with ${}^{2}B_{1g}$ ground state of Cu(II) square planar complex [33, 34]. The g_{||} values are found to be less than 2.3 suggesting the covalent nature of metal-ligand bond. The G values are more than 4 indicating negligible exchange interaction between the Cu(II) centres [35].

ESI Mass Spectral Studies

0.30

The ligands and their Cu(II) complexes have been studied for their mass spectral analysis. The ESI mass spectra of ligands exhibit $[M + H]^+$ peaks at m/z 341(L¹), 307(L²), 321(L³) and 325(L⁴). The ESI mass spectra of complexes exhibit peaks at m/z 765 $[M + Na]^+$ for $[Cu(L^1)_2]$, 691 $[M + NH_4]^+$ for $[Cu(L^2)_2]$, 624 $[M + Na]^+$ for $[Cu(L^3)_2]$ and 726 [M +NH₄]⁺ for $[Cu(L^4)_2]$. The m/z fragments of complexes confirm the stoichiometry of the complexes as $[Cu(L)_2]$ type. The representative mass spectra of L¹, L², $[Cu(L^1)_2]$ and $[Cu(L^2)_2]$ are given in Figs. S10-S13.



 $[Cu(L^3),]$ 0.25 0.20 Absorbance 0.15 0.10 0.05 0.00 250 300 350 450 400 500 Wavelength(nm) 0.30 $[Cu(L^4),]$ 8.00E-00 0.25 0.20 Absorbance 0.15 0.10 0.05 0.00 250 300 350 400 450 Wavelength(nm)

Fig. 4 UV-Vis absorption spectra of complexes $[Cu(L^1)_2]$, $[Cu(L^2)_2]$, $[Cu(L^3)_2]$ and $[Cu(L^4)_2]$ in the absence (dashed line) and presence (solid line) of increasing amounts of CT-DNA in Tris HCl buffer (pH 7.2) at 25 °C. Conditions: $[Complex] = 10 \ \mu M$, [DNA] = 0-

10 $\mu M.$ Arrow (\downarrow) shows the hypochromic and red shift upon increasing CT-DNA concentration. Inset: linear plot for the calculation of the intrinsic DNA binding constant, K_b



1.8-

1.5

1.3-1.2





200

150

 $\left[\operatorname{Cu}(\operatorname{L}^{3})_{2}\right]$

Fig. 5 Fluorescence emission spectra of CT-DNA-EB system in the absence and presence of increasing concentration of complexes $[Cu(L^1)_2]$, $[Cu(L^2)_2]$, $[Cu(L^3)_2]$ and $[Cu(L^4)_2]$ in Tris HCl buffer (pH 7.2) at 25 °C. Conditions: [Complex] = $0-60 \mu$ M,

Fig. 6 Effect of increasing amounts of EB, [Cu(L1)2], $[Cu(L^2)_2]$, $[Cu(L^3)_2]$ and $[Cu(L^4)_2]$ on the relative viscosity of CT-DNA at 30 \pm 0.1 °C

[DNA] = 125 μ M, [EB] = 12.5 μ M. Arrow (\downarrow) shows the emission intensity decreases upon increasing concentration of the complexes. Inset: I₀/I versus r





Fig. 7 Oxidative cleavage of supercoiled pBR322 DNA($0.2 \ \mu g$, 33.3 μ M) at 37 °C in 5 mM Tris. HCl/5 mM NaCl buffer by the [Cu(L¹)₂] and [Cu(L²)₂]. **a** Lane 1, DNA control; Lane 2, DNA + H₂O₂ (1 mM); Lane 3–8, DNA + H₂O₂ (1 mM) + [Cu(L¹)₂] (20, 30, 40, 50, 60 and 70 μ M) respectively. **b** Lane 1, DNA control; Lane 2, DNA + H₂O₂ (1 mM); Lane 3–8, DNA + H₂O₂ (1 mM); Lane 3–8, DNA + H₂O₂ (1 mM) + [Cu(L²)₂] (20, 30, 40, 50, 60 and 70 μ M) respectively



Thermal Analysis

The thermal decomposition of Cu(II) complexes was studied using thermo gravimetric technique. The experiment was performed at temperature 27 °C to 1000 °C at heating rate of 10 °C min⁻¹ under nitrogen atmosphere. The TGA curves of complexes [Cu(L¹)₂], [Cu(L²)₂] are shown in Fig. S14. The TGA curves of the complexes show a two step decomposition process. All the complexes are stable up to 200 °C confirming that there is no coordinated water molecules present in the complex molecule. The first weight loss occurred in the range of 209–336 °C corresponding to the partial loss of ligand moiety. The second mass loss is appeared at 375–789 °C corresponding to the complete departure of the complexes and the loss of their ligand portion. The residue was the copper oxide (CuO).

DNA Binding Studies

UV-Vis Spectroscopic Studies

Electronic absorption spectroscopy is an effective technique to study the binding mode and extent of binding metal complexes with DNA [36, 37]. Absorption spectra of all the Cu(II) complexes were recorded in the absence and presence CT-DNA and shown in Fig. 4. The binding mode of the Cu(II) complexes to DNA through intercalation is characterized by the change in absorbance (hypochromism) and red shift in wavelength [38]. Hypochromism is observed due to the presence of aromatic chromophore which might facilitate the interaction of the complexes with the CT-DNA bases via non covalent π - π * interactions. When the complexes intercalate to the base pairs of CT-DNA, the π^* orbital of the intercalated ligand in the complexes can couple with π orbital of the DNA base pairs, and then decreasing the π - π * transition energies. On the other hand, the coupling π^* orbitals are partially filled by electrons, thus decreasing the transition probabilities [39, 40]. The absorption spectra of the Cu(II) complexes show a band at 407-379 nm. Upon incremental addition of CT-DNA, the band underwent hypochromism with a slight red shift. The K_b (intrinsic binding constant) is calculated from a plot of $[DNA]/(\varepsilon_a - \varepsilon_f)$ Vs [DNA] using the equation: [DNA]/ $(\varepsilon_a - \varepsilon_f) = [DNA]/(\varepsilon_b - \varepsilon_f) + 1/K_b(\varepsilon_b - \varepsilon_f)$, where [DNA] is the concentration of DNA in base pairs, ε_a is the apparent coefficient of A_{obsd}/[complex], ε_f and ε_b represent the extinction coefficients of the free and fully bound forms of the complex, respectively [41]. The binding constant K_b is calculated by the ratio of slope to the intercept. The binding constant K_b for $[Cu(L^{1})_{2}]$ is 8.5 ± 0.02 × 10⁶ M⁻¹, $[Cu(L^{2})_{2}]$ is $8.6 \pm 0.02 \times 10^5 \text{ M}^{-1}$, [Cu(L³)₂] is $2.86 \pm 0.02 \times 10^6 \text{ M}^{-1}$ and $[Cu(L^4)_2]$ is $1.91 \pm 0.02 \times 10^6 \text{ M}^{-1}$. The K_b values are

Fig. 8 Photoactivated cleavage of supercoiled pBR322 DNA $(0.2 \ \mu g, 33.3 \ \mu M)$ at 37 °C in 5 mM Tris. HCl/5 mM NaCl buffer by Cu(L¹)₂, Cu(L²)₂ using UV irradiation of wavelength 345 nm. **a** Lane 1, DNA control; Lane 2–7, DNA + [Cu(L¹)₂] (20, 30, 40, 50, 60, 70 and 80 \ \mu M) respectively. **b** Lane 1, DNA control; Lane 2–7, DNA + [Cu(L²)₂] (20, 30, 40, 50, 60, 70 and 80 \ \mu M) respectively



 Table 3
 The inhibition zone

 values of the ligands and their
 complexes at 1 mg/mL

 concentration
 concentration

Compound	Bacteriu	Bacterium (mm)					Fungi (mm)	
	Gram-negative bacteria			Gram-positive bacteria				
	E. coli	P. putida	K.pneumoniae	B. subtilis	S. aureus	A. niger	C. albicans	
L^1	11	10	11	8	10	12	10	
L ²	9	10	9	10	9	10	7	
L ³	11	9	10	9	11	11	9	
L^4	10	11	9	10	11	10	8	
$[Cu(L^1)_2]$	25	24	22	22	23	24	25	
$[Cu(L^2)_2]$	24	23	21	20	22	21	23	
$[Cu(L^3)_2]$	23	20	21	19	21	23	24	
$[Cu(L^4)_2]$	21	22	20	21	22	22	23	
Ampicillin	31	30	33	34	30	-	-	
Ketoconazole	-	-	-	-	-	30	32	

relatively less with the potential intercalators like ethidium bromide (EB) ($K_b = 7 \times 10^7 \text{ M}^{-1}$) [42].

Fluorescence Study

Fluorescence experiments were performed to gain support for the binding mode of the complexes with CT-DNA. EB is one of the most sensitive fluorescent probes which can bind with DNA. There is an increase in the intensity of fluorescence spectra of EB in presence of CT-DNA because of intercalation between the DNA basepairs [43, 44]. The quenching of EB bound to DNA is calculated with the addition of increasing concentration of metal complexes. The Fig. 5 is showing the decrease in the intensity of EB with increase in the concentration of Cu(II) complexes. The reduction in the emission intensity is due to displacement of EB bound DNA with Cu(II) complexes [45]. The K_{sv} values calculated for all the complexes are $6.4 \pm 0.02 \times 10^4$ M⁻¹ [Cu(L¹)₂], $\begin{array}{l} 3.3 \, \pm \, 0.02 \, \times \, 10^4 \, \, M^{-1} \, \, [Cu(L^2)_2], \, 1.6 \, \pm \, 0.02 \, \times \, 10^4 \, \, M^{-1} \\ [Cu(L^3)_2], \, 1.3 \, \pm \, 0.02 \, \times \, 10^4 \, \, M^{-1} \, [Cu(L^4)_2]. \end{array}$

Viscosity Measurements

The viscometric measurements were also carried out to further confirm the mode of interaction between the Cu(II) complexes and CT-DNA. A classical intercalation can cause an increase in the viscosity of DNA since it increases the separation between base pairs at intercalation site, hence an increase in overall length of the double helix [46, 47]. However a partial or non classical intercalation of the complex results in bending of the DNA helix, which decreases the effective length of DNA, with a concomitant decrease in its viscosity. The effects of all the Cu(II) complexes on the viscosity of DNA at 30 ± 0.1 °C are shown in Fig. 6. The relative specific viscosity of DNA is determined by increasing the concentration of the complex while the CT-DNA concentration was kept constant.



Fig. 9 a Anti bacterial activity, b Anti fungal activity of Schiff bases and their Cu(II) complexes

By increasing the complex concentration to DNA the viscosity of DNA increases gradually which leads to an intercalative mode of binding [48]. These results revealed that the binding or der of the complexes is found to be $[Cu(L^1)_2] > [Cu(L^2)_2] > [Cu(L^3)_2] > [Cu(L^4)_2]$. Among all Cu(II) complexes the increase in viscosity is more in the case of $[Cu(L^1)_2]$. This is due to lengthening of the DNA duplex by the extended planarity of bicyclic naphthalene moiety of $[Cu(L^1)_2]$ which increases the intercalation compare to remaining complexes with monocyclic phenyl ring moiety [49].

DNA Cleavage Experiments

The DNA cleavage ability of Cu(II) complexes is investigated by agarose gel electrophoresis method with super coiled pBR322 DNA at different complex concentrations in the presence of H₂O₂ and UV light. The ability of the complexes in DNA cleavage is estimated by the conversion of DNA from Form I to Form II and Form III. During electrophoresis the fastest migration is detected in the supercoiled form (Form I). While the scission occurs on one strand the supercoiled form relaxes to generate nicked form (Form II), if both strands are cleaved linear form (Form III) appears which migrates in between Form I and Form II [50]. The concentration effect has been studied for the complexes by increasing the concentrations from 20 to 70 µM. As shown in Figs. 7 and 8 with increase in the complex concentration the circular supercoiled DNA converted to nicked DNA. Oxidative DNA cleavage of all the complexes were carried out in the presence of H_2O_2 , there is no cleavage was observed in lane 1 and lane 2, while increasing the concentration of complexes (lane 3-8) form II is increased. The above results indicate that complexes can cleave the pBR322 DNA by oxidative cleavage via formation of hydroxyl radical. Photolytic DNA cleavage was performed in presence of UV irradiation at 345 nm, no DNA cleavage was observed in lane 1. From lane 2-8 DNA cleavage was observed by converting supercoiled DNA into nicked form, stating that cleavage capacity of the complexes is more at higher concentrations.

Biological Evaluation

The in vitro antimicrobial screening of all the ligands and their Cu(II) complexes were tested against the bacterial *species E. coli, P. Putida, K. pneumoniae, B. subtilis* and *S. aureus* and fungal species *A. niger*, and *C. albicans* by paper disc method. Ampicillin (antibacterial) and Ketoconazole (antifungal) were used as reference drugs. The antimicrobial activity stated that Cu(II) complexes show higher activity than corresponding free Schiff base ligands. The inhibition zone values of the compounds are summarized in Table 3. The inhibition zone

values of all the Cu(II) complexes revealed that the complexes exhibit moderate to strong antimicrobial activity when compared with the standard drug (Fig. 9). The increase in antimicrobial activity of the complexes can be explained on the basis of Overtone's concept [51] and Tweedy's Chelation theory [52]. According to Overtone's concept of cell permeability, the lipid membrane that surrounds the cell favors the passage of only the lipid soluble materials makes liposolubility as an important factor, which controls the antimicrobial activity. On chelation with metal ions the polarity of the metal ions reduced due to overlapping of ligand orbital and the partial sharing of its positive charge with the donor groups [53]. This process increases the lipophilicity of metal complexes [54].

Conclusion

Four novel Schiff bases and their binary Cu(II) complexes have been synthesized and characterized by analytical and spectral data. From the obtained data a square planar geometry has been assigned to all Cu(II) complexes. The UV-Vis absorption, fluorescence, and viscosity measurements revealed that the complexes bind with CT-DNA through an intercalation mode. From the DNA cleavage studies of Cu(II) complexes it is observed that the complexes effectively cleave supercoiled pBR322 DNA in the presence of H_2O_2 and also UV light. In vitro antimicrobial activity studies of Schiff base ligands and their Cu(II) complexes revealed that the activity increases upon coordination. Among all the complexes highest activity is observed for Cu(L¹)₂.

Acknowledgments We express our sincere thanks to the Head, Department of Chemistry for providing the necessary facilities. We are thankful to the Director, CFRD, Osmania University, Hyderabad, and the Director, IICT, Hyderabad, and the SAIF, IIT Bombay for providing spectral and analytical data. We are also thankful to DST-PURSE, DST-SERB and UGC-UPE (FAR) for providing financial assistance.

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