

Antimicrobial, Cytotoxicity and Molecular Docking Study of New Quinoline Schiff Base and its Metal Complexes

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A new quinoline Schiff base ligand was synthesized by the reaction of 2-hydroxy-7-methylquinolin-3-carbaldehyde and 4-methylbenzene sulfonohydrazide. Synthesized Schiff base further utilized for the formation of stable metal complexes with Cu(II), Ni(II), Co(II) and Cd(II) metal salts and characterized by different spectroscopic techniques i.e., ¹H NMR, ¹³C NMR, FT-IR, UV-visible, ESR, MASS and TGA. The low molar conductance values indicate that synthesized metal(II) complexes were non-electrolytes. The magnetic moment value indicates that Cu(II), Ni(II) and Co(II) complexes were paramagnetic. Further, these compounds were screened for inhibition activity against four bacterial strains, three fungal strains and cytotoxicity against the A-549 and MCF-7 cell lines by using the MTT method. Among the synthesized complexes, metal complexes exhibited excellent anticancer activity against the human lung cancer cell line (A-549). Ligand and its Cd(II) complex showed good antibacterial activity. Furthermore, molecular docking study shows the significant binding affinity of metal complexes with tubulin protein. Hence, present study proposed that all the synthesized Schiff base metal(II) complexes have excellent biological activity and could be act as potential anticancer agents.

Keywords: Schiff base, Sulfonohydrazide, Quinoline, MIC, Cytotoxicity.

INTRODUCTION

Schiff bases are an interesting class of compounds, which attracts considerable attention of researchers. This is because of their diversity in its property, structural variability and their easy preparation [1,2]. They play an important role in the formation of the chelate compounds [3]. Schiff base having electrons reaches functional groups such as -OH, -SH and -NH₂ at adjacent positions to the azomethine group help to develop coordination with metal ions, which form stable complexes [4-12].

Metal complexes derived from Schiff bases are an interesting area of research. Such complexes have been widely used as biological [13-20], analytical [21,22] and catalyst [23-25] field. From the study, it was observed that the coordination of Schiff base with metal ions increase the biological activity of Schiff base [26,27].

Among the heterocyclic compounds, quinoline and its derivatives were found to be a significant class in the biological

field [28]. Several derivatives of quinoline are found to be effective antibacterial [17], antimicrobial [29], fungicides [17], antiviral [30], anti-inflammatory [31, 32] and antitumor activities [33]. Simultaneously, metal complexes derived from quinoline Schiff bases have extensive applications in different areas such as, catalyst in various types of reactions [34,35], dyes in solar cells [36], corrosion inhibitor [37], antioxidant [2], cytotoxic [28], DNA cleavage [38], anticancer [39], etc.

To find better antimicrobial and anticancer drug, we have designed and synthesized novel quinoline Schiff base and its metal(II) complexes. Synthesized compounds were confirmed by different analytical techniques and studied for its antibacterial, antifungal and cytotoxicity activities.

EXPERIMENTAL

All the required chemicals were purchased from Sigma-Aldrich Chemical Co., (USA), Molychem Chemical Supplier (Mumbai, India) and used as such for further synthesis. Fourier

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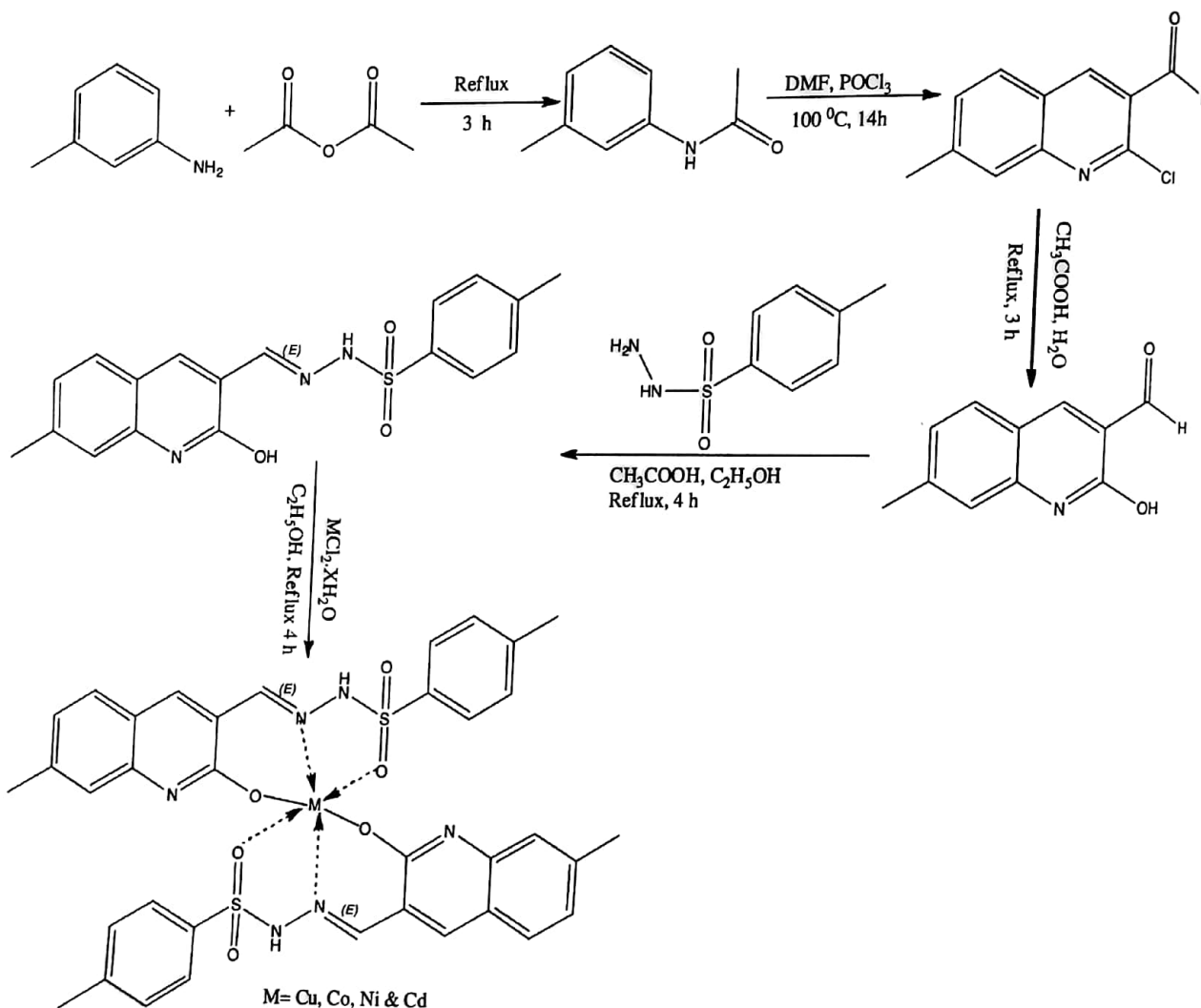


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52 transform infrared spectroscopy (FTIR) spectra were recorded
 53 on a Nicolet iS10, thermos Scientific, USA spectrophotometer
 54 using KBr pellets in the range of 4000-400 cm^{-1} . Proton nuclear
 55 magnetic resonance (^1H and ^{13}C NMR) spectra of Schiff base
 56 were measured in $\text{DMSO-}d_6$ solvent on a Bruker Avance 400
 57 MHz and 100MHz spectrometers, respectively. Electronic
 58 (UV-Visible) spectra were recorded using Carry 100 UV-visible
 59 spectrophotometer. Electron spins resonance (ESR) spectra
 60 of Cu(II) complex were performed on the JES-FA200 ESR
 61 Spectrometer. Thermogravimetric analysis (TGA) of metal
 62 complexes was performed on Mettler-Toledo instrument at the
 63 heating range of 20 $^\circ\text{C}/\text{min}$ with a temperature range of 25 to
 64 1000 $^\circ\text{C}$. Electrospray ionization mass spectra (ESIMS) were
 65 recorded on a Waters Micromass Q-T of Micro with atmos-
 66 pheric pressure chemical ionization (APCI) sources. Elemental
 67 analyses were performed on a FLASH EA 1112 series instru-
 68 ment. The magnetic moments were measured by the Gouy
 69 method at 25 $^\circ\text{C}$ using the MK1 Johnson Matthey model. Molar
 70 conductance was measured on DDS-11C type conductivity
 71 Bridge in DMSO solution at a concentration of 10^{-3} M.

72 **Synthesis of quinoline Schiff base ligand (HL):** 2-Chloro-
 73 7-methylquinoline-3-carbaldehyde was synthesized using

74 Vilsmeier-Haack reaction as reported method [40]. The formed
 75 2-chloro-7-methylquinoline-3-carbaldehyde was further used
 76 for the formation of 2-hydroxy-7-methylquinoline-3-carbal-
 77 dehyde. 2-Chloro-7-methylquinoline-3-carbaldehyde (20 mmol)
 78 and 2 mL H_2O was dissolved in 4 mL acetic acid and refluxed
 79 for 4 h. The progress of the reaction was checked by thin-layer
 80 chromatography (TLC). The obtained product, 2-hydroxy-7-
 81 methylquinoline-3-carbaldehyde was washed with distilled
 82 water and recrystallized in absolute ethanol. The recrystallized
 83 2-hydroxy-7-methylquinoline-3-carbaldehyde was further used
 84 for the synthesis of the final Schiff base ligand. For the formation
 85 of the final ligand, a mixture of 2-hydroxy-7-methylquinoline-
 86 3-carbaldehyde (1 mmol), 4-methylbenzenesulfonylhydrazide
 87 (1 mmol) and 5-10 drops acetic acid in 15 mL ethanol was
 88 placed in a round bottom flask. The mixture was refluxed at
 89 75 $^\circ\text{C}$ for 5 h. The progress of the reaction was checked by
 90 TLC. The reaction mixture was quenched with crushed ice and
 91 extracted with ethyl acetate. The organic extracts were washed
 92 with brine solution and dried over anhydrous sodium sulphate.
 93 The solvent was evaporated under reduced pressure to obtain
 94 the corresponding crude compound, which was purified with
 95 ethanol (Scheme-I). ^1H NMR: (100 MHz, $\text{DMSO-}d_6$, δ ppm):



Scheme-I: Synthesis of ligand and its metal complexes

96 11.57 (s, 1H, NH), 11.18 (s, 1H, OH), 8.12 (s, 1H, Ar-H), 8.07
 97 (s, 1H, Ar-H), 7.73-7.71 (d, 2H, Ar-H, $J = 7.5$ Hz), 6.89-6.87 (d,
 98 2H, Ar-H, $J = 7.5$ Hz) 7.56 (s, 1H, Ar-C=CH), 7.38-7.36 (d,
 99 1H, Ar-H, $J = 8$ Hz), 7.22-7.20 (d, 1H, Ar-H, $J = 8$ Hz) and
 100 2.30 (s, 6H, -CH₃); ¹³C NMR (100 MHz, DMSO-*d*₆, δ ppm):
 101 161.84, 143.48, 142.43, 141.93, 139.29, 136.45, 135.19,
 102 129.45, 128.45, 127.45, 124.18, 123.90, 117.21, 115.43, 21.90
 103 and 21.51.

104 **Synthesis of metal(II) complexes:** A hot ethanolic
 105 solutions of metal(II) chloride (5 mL, 0.0015 mol) were added
 106 to 30 mL hot ethanolic ligand solution (0.0030 mol) in 250
 107 mL round bottom flask. The reaction mixture was stirred for
 108 30 min and few drops of 5% NaOH solution were added to
 109 maintain basic condition of the reaction. Further, the reaction
 110 mixture was refluxed for 4 h to complete the formation of the
 111 metal(II) complexes. The formed coloured metal(II) complexes
 112 were washed with distilled water followed by ethanol.

113 Biological study

114 **Antibacterial study:** The broth dilution method was used
 115 to measure the minimum inhibitory concentration (MIC) of
 116 prepared compounds [41]. Dimethylsulfoxide (DMSO) was
 117 used as a solvent for diluent and it has no biological effect on
 118 selected bacterial strain [42]. In this study, two Gram-negative
 119 bacteria viz., *Escherichia coli* (MTCC 443) and *Pseudomonas*
 120 *aeruginosa* (MTCC 1688) and two Gram-positive bacteria
 121 viz., *Staphylococcus aureus* (MTCC 96) and *Staphylococcus*
 122 *pyogenus* (MTCC 442) were tested against the synthesized
 123 Schiff base ligand and its metal(II) complexes. Chloram-
 124 phenicol and ampicillin were used as the standard drugs for
 125 reference. Serial dilutions of Schiff base ligand and its metal
 126 (II) complexes were prepared for the primary and secondary
 127 screening. The control plate with no prepared compounds and
 128 drug was subculture spreading evenly over a plate suitable for
 129 the growth of selected bacterial pathogens and kept overnight
 130 at 37 °C in incubator. The MIC of the control bacterial strain
 131 was assessed to check the efficacy of the reference drug concen-
 132 trations. The lowest concentration was recorded as the MIC.
 133 The amount of growth from the control plate before incubation
 134 was compared. Synthesized compounds were diluted to 2000
 135 $\mu\text{g/mL}$ concentration as a stock solution. In primary screening,
 136 125, 250 and 500 $\mu\text{g/mL}$ concentrations of synthesized comp-
 137 ounds were taken. The synthesized compounds found active
 138 in primary screening were further tested in the second set of
 139 dilutions against all the selected pathogens. The particles found
 140 active in primary screening were diluted similarly to 100, 50,
 141 25, 12.5, 6.250, 3.125 and 1.5625 $\mu\text{g/mL}$ concentrations. The
 142 MIC was considered for the dilution showing at least 99%
 143 inhibition.

144 **Antifungal study:** The antifungal activity of the synthe-
 145 sized compounds was studied with three fungal strains viz.,
 146 *Aspergillus clavatus* (MTCC 1323), *Candida albicans* (MTCC
 147 227) and *Aspergillus niger* (MTCC 282) using agar dilution
 148 protocol [41]. To determine MIC, a stock solution of synthe-
 149 sized compounds was prepared in DMSO and then incorp-
 150 orated in a specified quantity of sterile molten dextrose agar
 151 for antifungal screening. The inoculate was prepared by taking

a stock to about 100 mL of nutrient broth in 250 mL sterilized
 and clean conical flasks. The conical flasks were incubated at
 27 °C for 24 h before the experiment. The plates were kept
 under aseptic conditions to allow the diffusion of the solution
 properly into potato-dextrose agar medium. Then, the plates
 were incubated at 25 °C for 48 h. The highest dilution dis-
 playing at least 99% inhibition zone was taken as MIC with
 nystatin and griseofulvin as a standard reference drugs. The
 triplicate analysis was performed to minimize errors.

Cytotoxicity: The cells were seeded at a density of approx-
 imately 5×10^3 cells well in a 96-well flat bottom microtitre
 plate and maintained at 37 °C overnight in 95% humidity and
 5% CO₂. Different concentrations (50, 40, 30, 20, 10, 5 μM)
 of samples were treated and the cells were incubated for the
 next 48 h. The cells in well were washed twice with phosphate
 buffer saline (PBS) and 20 μL of MTT [3-(4,5-dimethylthiazol-
 2-yl)-2,5 diphenyltetrazolium bromide] staining solution (5
 mg mL⁻¹ in phosphate buffer saline) was added to each well
 and the plate was incubated at 37 °C. After 4 h, 100 μL DMSO
 was added to each well to dissolve the formant crystals and
 the absorbance was recorded at 570 nm using a microplate
 reader.

Molecular docking study: To investigate the binding
 mode of various drug-metal complexes (Cd, Co, Cu and Ni)
 with β -tubulin receptor, molecular docking was performed
 using Auto Dock software [43]. The microtubules are essential
 in cell division [44]. The inhibition of microtubules structure
 leads to disturb its dynamics that's leads to cell apoptosis and
 death [45]. Hence, we used β -tubulin as target receptor for the
 molecular docking study, to understand the binding mode of
 various metal-drug complexes with β -tubulin. The crystal
 structure of tubulin (1JFF.pdb) was retrieved from the protein
 database. The three dimensional atomic coordinates of the
 metal complexes (Cd, Co, Cu and Ni) were built Discovery
 Studio Visualizer [46]. The grid box of $80 \times 80 \times 80$ was built
 around the paclitaxel binding pocket with grid spacing 0.375 Å.
 Herein, we performed a local docking protocol, to explore the
 binding mode of metal-drug complexes using AutoDock. Here,
 Lamarckian Genetic Algorithm was used for molecular docking
 and output conformations were further clustered using an all-
 atom RMSD with a cut-off of 4 Å. The lowest binding energy
 conformation were further utilized for bonding and non-
 bonding interactions analysis and visualization using (DeLano,
 2002) and Discovery Studio visualizer [46] and PyMol [47],
 respectively.

RESULTS AND DISCUSSION

Elemental analysis: From the elemental analysis data
 (Table-1), it was confirmed that the synthesized Schiff base
 ligand and its metal(II) complexes are completely formed. All
 the prepared compounds were subjected to molar conductance
 in DMSO solvent at the concentration of 10^{-3} M. The molar
 conductance of the metal complexes was found to be in the
 range of 47-68 $\Omega^{-1} \text{cm}^2 \text{mol}^{-1}$. From the obtained values (Table-
 1), it was proved that all the synthesized metal(II) complexes
 were non-electrolyte with evidence for the absence of water
 molecules in the coordination sphere [62].

TABLE-I
PHYSICAL AND ANALYTICAL DATA OF QUINOLINE SCHIFF BASE LIGAND AND ITS METAL(II) COMPLEXES

Compounds	Yield (%)	m.p. (°C)	Λ_m (cm ² Ω ⁻¹ mol ⁻¹)	μ_{eff} (B.M.)	m.w.	Elemental analysis (%): Found (calcd.)			
						C	N	H	S
C ₁₈ H ₁₇ N ₃ O ₃ S	81	220-222	-	-	355.10	59.98 (60.83)	11.52 (11.82)	4.56 (4.82)	8.65 (9.02)
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Cu]	73	>300	58	1.78	772.35	55.85 (55.98)	10.86 (10.88)	4.14 (4.18)	8.41 (8.30)
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Ni]	74	>300	47	3.56	766.50	56.19 (56.34)	10.57 (10.96)	4.13 (4.20)	8.29 (8.36)
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Co]	65	>300	63	4.81	765.74	56.24 (56.32)	10.69 (10.95)	4.10 (4.20)	8.17 (8.35)
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Cd]	78	>300	68	-	823.22	52.73 (52.65)	10.30 (9.85)	3.66 (3.93)	7.55 (7.81)

207 **FT-IR spectroscopy:** FT-IR spectra of all the prepared
208 Schiff base ligand and its metal(II) complexes were carried
209 out and clearly showed a difference with FT-IR peaks. The peak
210 observed at 1652 cm⁻¹ was due to the azomethine ν(C=N) stret-
211 ching, which shifted to a lower wave value (1634-1622 cm⁻¹)
212 in the complexes indicating the participation of azomethine
213 nitrogen in coordination with the metal ion (N-M) [48]. The
214 phenolic ν(O-M) stretching vibration band was observed at
215 1349 cm⁻¹ in the free ligand. In metal(II) complexes, this band
216 appeared at lower frequency 1036-1018 cm⁻¹ region, confirming
217 the participation of the phenolic group in complex formation
218 [49]. The vibration bands for the SO₂ group in the free ligand
219 molecule appeared at 1316 cm⁻¹ and 1188 cm⁻¹ ν_{asym}(SO₂) and
220 ν_{sym}(SO₂), respectively. In the metal(II) complexes, the asym/
221 symm. bands shifted to 1266-1222 and 1134-1112 cm⁻¹, respec-
222 tively, upon the coordination of the central metal ion [49-55].
223 The additional peaks observed in metal complexes in the range
224 of 460-419 cm⁻¹ were due to N-M bonding and 517-509 cm⁻¹
225 were due to O-M bonding [56,57]. The characteristic bands
226 of the stretching frequency are listed in Table-2.

227 **UV-Visible spectra and magnetic moment:** UV-Visible
228 spectra of the synthesized Schiff base ligand and its metal(II)
229 complexes were recorded in DMSO solvent (10⁻⁵ M). The
230 absorption band in ligand molecule appeared at 320 nm and
231 274 nm for π→π* and n→π* transitions, respectively. These
232 bands in metal(II) complexes were shifted at bathochromic
233 shift. For Ni(II), Cd(II), Cu(II) and Co(II) complexes, the
234 absorption bands appeared at 333 and 381, 343 and 389, 339
235 and 388, 352 and 390 nm, respectively. The absorption band
236 at a bathochromic shift in metal complexes confirmed that the
237 ligand moiety was coordinated with central metal ions [58].

238 The magnetic moment for Cu(II), Ni(II) and Co(II)
239 complexes was calculated using the Gouy balance at 25 °C.
240 The observed magnetic moment values for Cu(II), Ni(II) and
241 Cd(II) complexes were found to be 1.78, 3.56 and 4.81 B.M.
242 respectively. The magnetic moment obtained for Cu(II) complex
243 was approximately equal to spin-only value of one unpaired
244 electron 1.75 B.M. for octahedral geometry [59]. In case of
245 Ni(II) and Co(II) complexes, the observed magnetic moment

values were approximately equal to its reported octahedral
geometry [60,61].

246 **ESR spectra:** The X-band ESR spectra of copper(II)
247 complex were recorded at liquid nitrogen temperature in DMSO
248 solvent. Fig. 1 shown ESR spectra of Cu(II) complex. It provided
249 information about the environment of the central metal ion in
250 the complex. Covalency parameter α² was calculated to deter-
251 mine the bonding between central metal ions and surrounding
252 ligand. The following equation was used to calculate the coval-
253 ency parameter α²:
254
255

$$\alpha^2 \text{Cu(II)} = - (A_{||}/0.036) + (g_{||} - 2.002) + 3/7 (g_{\perp} - 2.002) + 0.04 (1) \quad 256$$



Fig. 1. ESR spectrum of Cu(II) complex

257 Hamiltonian parameter was used to calculate the ground
258 state of the Cu(II) complex. All the calculated values are given
259 in Table-3. The obtained g_{||} and g_⊥ values are greater than free
260 electron g values. The trend in g values is g_{||} > g_⊥ > 2.0023,
261 these values indicate that the unpaired electrons present in d_{xy}²
262 ground state, which are characteristics of octahedral geometry
263 [63]. The calculated G value (axial symmetry parameter) was
264 found to be > 4, which suggested that the interactions of Cu-Cu
265 ions are negligible [64]. The effective magnetic moment was
266 calculated using equation:

$$\mu_{eff}^2 = \frac{3}{4} (g_{av}^2) \quad 267$$

TABLE-2
KEY FT-IR FREQUENCY (cm⁻¹) OF QUINOLINE SCHIFF BASE LIGAND AND ITS METAL(II) COMPLEXES

Compounds	ν(C=N)	ν(C-O)	ν(N-M)	ν(O-M)	ν _{asym} (SO ₂)	ν _{sym} (SO ₂)
C ₁₈ H ₁₇ N ₃ O ₃ S	1652	1349	-	-	1316	1188
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Cu]	1622	1035	460	517	1246	1112
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Ni]	1632	1036	419	510	1223	1125
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Co]	1634	1018	459	516	1222	1134
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Cd]	1633	1035	452	509	1266	1125

TABLE 3
FT-IR SPECTRAL DATA OF Cu(II) COMPLEX

ν_1	ν_2	ν_3	ν_4	ν_5	ν_6
2.045	2.199	2.096	4.606	0.178	1.80

68 TGA: Table-4 summarized with decomposition stages,
69 temperature range, loss of weight (actual and calculated) and
70 assignments of the loss fragments. All the complexes start
71 decomposed near 200 °C temperature, which indicates an
72 absence of water moiety. The TG curve of the Co complex is
73 shown in Fig. 2. The thermogram of the Co²⁺ complex exhibited
74 three decomposition stages. In stage first, the 12.19% weight
75 loss (calculated 13.2%) was observed between the temperature
76 range 200-240 °C, corresponding to loss of C₈H₇ ligand moiety.
77 In the second stage, 28.97% weight loss was observed (calcd.
78 28.10%) in the temperature range 280-680 °C, due to loss of
79 C₁₁H₆N₃O₂ moiety of the coordinated ligand molecule. In step
80 third, 23.88% loss was observed (calculated 23.80%) in the
81 range of 720-980 °C with loss of C₈H₆N₂O₃ moiety. In the end,
82 21.91% residue remains present. The remaining residue contains
83 metal oxide along with non-decomposed organic moiety.

84 **Mass spectra:** To confirm the complete formation, all
85 the metal(II) complexes were subjected to ESI-MS. The ESI-
86 MS spectrum of Ni(II) complex (Fig. 3) shows a molecular
87 ion peak at M⁺ 766, which corresponds to its molecular weight.
88 Ligand, Cu(II), Co(II) and Cd(II) metal complexes show
89 molecular ion peak at M⁺ 356, 771, 765 and 823, respectively.
90 The obtained molecular ion peaks of prepared compounds are
91 exactly matched with its corresponding molecular weight. From

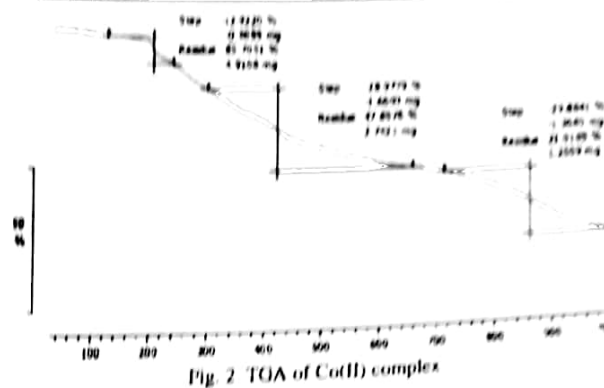


Fig. 2 TGA of Co(II) complex

study of ESI-MS spectra, it was confirmed that the synthesized
compounds are completely formed.

Antibacterial activity: The prepared compounds have
been screened for antibacterial activity with two Gram-positive
and two Gram-negative bacterial strains. Among the prepared
compounds, Schiff base ligand and its Cd(II) complex showed
excellent activity with all four bacterial strains. In case of
Cu(II), Co(II) and Ni(II) complexes, Cu(II) complex showed
excellent activity against *E. coli* and *P. aeruginosa* bacterial
strains (Fig. 4). The Co(II) complex has shown good to excellent
activity against *E. coli*, *S. aureus* and *S. pyogenes* bacterial strain.
Ni(II) complex was found to be weak active against the four
bacterial strains.

Antifungal activity: All the synthesized compounds were
screened against three fungal strain viz., *C. albicans*, *A. niger*
and *A. clavatus* at different concentrations ranging between

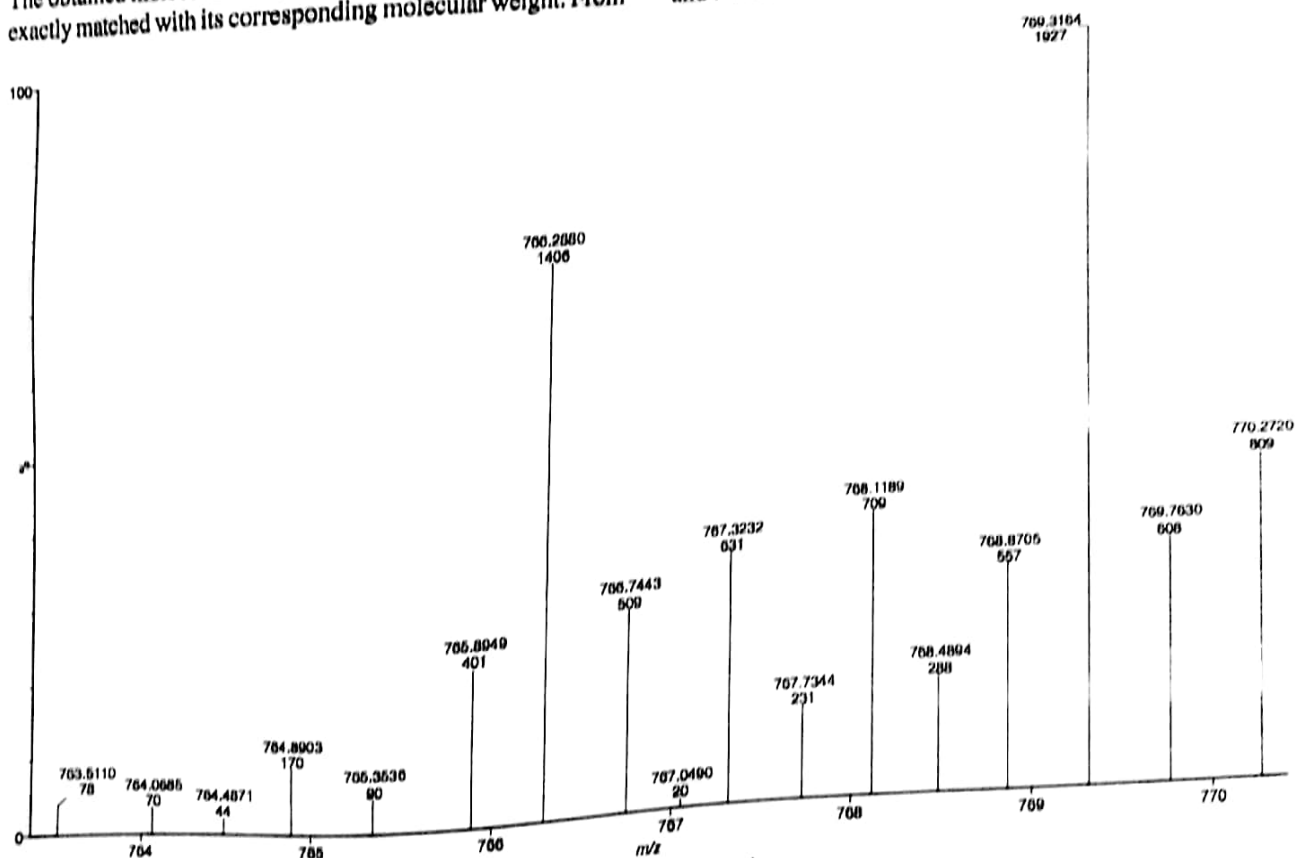


Fig. 3. ESI-MS of Ni complex

TABLE-4
STEPWISE THERMAL DECOMPOSITION STUDY OF QUINOLINE SCHIFF BASE METAL(II) COMPLEXES

Complex	Decomposition temp. (°C)	Weight loss (%)		Interference
		Observed	Calculated	
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Co]	200-240	12.92	13.22	C ₈ H ₈
	280-680	28.97	28.10	C ₁₁ H ₈ N ₂ O ₂
	720-980	23.88	23.80	C ₈ H ₈ N ₂ O ₂
	Residue	21.91	33.90	CoO + C ₇ H ₇ N ₂ O ₂ S
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Cu]	200-460	44.38	44.10	C ₁₀ H ₈ N ₂ O ₂ S
	550-980	16.56	16.80	C ₇ H ₇ N
	Residue	33.16	32.90	CuO + C ₇ H ₇ N ₂ O ₂ S
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Ni]	220-340	16.15	16.90	C ₇ H ₇ N
	350-640	21.65	22.15	C ₇ H ₇ NO ₂ S
	660-980	18.15	18.10	C ₇ H ₇ O ₂ S
	Residue	36.83	40.90	NiO + C ₁₀ H ₇ N ₂ O ₂
[(C ₁₈ H ₁₇ N ₃ O ₃ S) ₂ Cd]	180-260	13.84	14.21	C ₇ H ₇ N
	280-540	25.03	25.30	C ₇ H ₇ N ₂ O ₂ S
	560-980	22.53	22.43	C ₇ H ₇ N ₂ O ₂ S
	Residue	31.19	37.90	CdO + C ₁₂ H ₇ NO

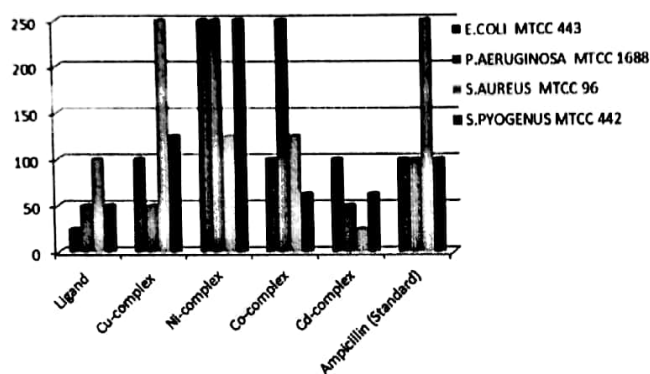


Fig. 4. Antibacterial activity of ligand and its metal complexes (MIC, mg mL⁻¹)

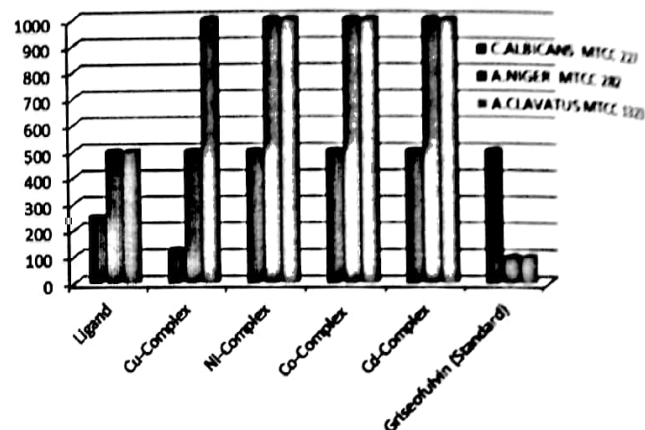


Fig. 5. Antifungal activity of ligand and its metal complexes (MIC, mg mL⁻¹)

308 100 and 1250 µg/mL using agar plate method. Among the
309 synthesized compounds ligand molecule has shown good
310 activity with all three fungal strains. In metal(II) complexes,
311 Cu(II) complex showed excellent activity against fungal strain
312 *C. albicans*. Co(II), Ni(II) and Cd(II) complexes exhibit good
313 activity against fungal strain *C. albicans* (Fig. 5).

314 **Cytotoxicity:** The *in vitro* cytotoxicity of ligand and its
315 metal(II) complexes was investigated against A-549 (human
316 lung cancer) and MCF-7 (human breast cancer) cell lines and
317 results are tabulated in Table-5. Paclitaxel was used as the
318 standard drug during the activity. The ligand and its metal(II)

complexes showed inhibition of cell value IC₅₀ in the range 308
33.55-47.61 µM for A-549 and 30.95-46.81 µM for MCF-7 309
cell lines. The ligand and its metal complexes exhibited higher 310
activity against the A-549 cancer cell line and lowered in the 311
case of the MCF-7 cancer cell line compared to standard. The 312
obtained results showed that the most of the synthesized metal 313
complexes were found to be more active than their corresponding 314
ligand molecule (Fig. 6). The order of activity of all the 315
synthesized compounds against the A-549 cancer cell line is 316

TABLE-5
HYDROGEN BONDING INTERACTIONS OF β-TUBULIN WITH
VARIOUS QUINOLINE SCHIFF BASE LIGAND METAL(II) COMPLEXES

Complexes	Binding energy (kcal/mol)	Atoms involved in the bonding interactions	Distance atom pair	Angle	Figure
Cd-Metal	-11.32	Drg-1:H - O-THR276	1.76776	143.055	8A
		ARG278:N - HC-Drg	3.5812		
Co-Metal	-10.70	Drg-H - O-THR276	1.78816	149.203	8B
		ARG278-N - HC-UNL	3.58511		
Cu-Metal	-12.39	Drg:H - O-THR276	1.95704	123.983	8C
		Drg:CH - O-THR276	2.91936		
		Drg:CH - O-ALA233	3.63567		
		ARG278-N - HC-Drg	3.83767		
Ni-Metal	-11.39	Drg-CH - O-THR276	1.7098	157.537	8D
		ASP226:O - HC-Drg	3.15848		

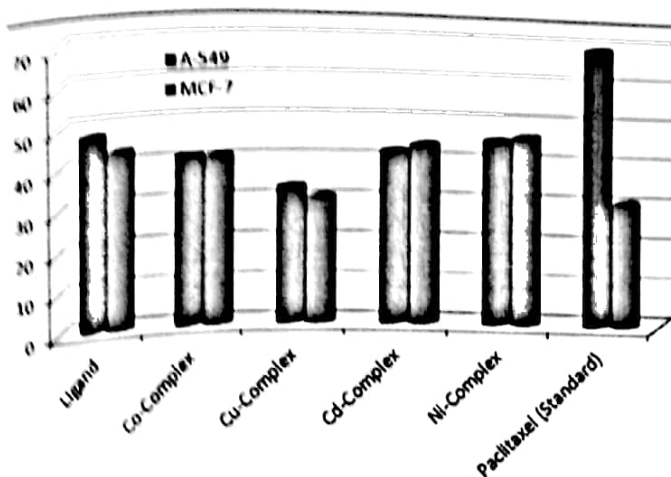


Fig. 6. In vitro cytotoxicity of ligand and metal complexes

328 Cu > Co > Cd > Ni-complex > ligand and for MCF-7 cancer
329 cell line Cu > Co > Cd > ligand > Ni-complex.

330 **Molecular docking of β -tubulin with drug-metal**
331 **complexes:** We employed molecular docking, to investigate
332 the interaction of metal complexes (Cd, Co, Cu and Ni) with
333 β -tubulin through AutoDock4.2 [43]. The lowest binding
334 energy conformation of metal complexes (Cd, Co, Cu and Ni)
335 with β -tubulin was found at -11.32, -10.70, -12.39 and -11.39
336 kcal/mol, respectively. All the metal(II) complexes show a
337 considerable binding energy and affinity with β -tubulin as
338 shown in Fig. 7. The analysis of β -tubulin-Cd metal complex
339 show the conventional hydrogen bonding interaction of residue
340 Thr276 (1.76 Å) and carbon hydrogen bonding interaction
341 with Arg278 (3.58 Å) (Table-5). The Asp226 show electrostatic
342 interactions, Thr220, Thr223, Phe272, Ser277, Gln282, Arg284
343 forms van der Waals interactions, Arg273 forms π -lone pair
344 and Asp226 makes π -anion type of interactions with Cd-metal
345 complex as shown in Fig. 8A. While, Leu217, Lys218, Leu219,
346 His229, Pro274, Leu371, Leu285, Pro360 forms hydrophobic

347 interactions with drug-Cd complex as shown in Fig. 8A. Next,
348 analysis of β -tubulin-Cd metal complex is stabilized by
349 hydrogen bonding interaction of Thr276 (1.78 Å) and carbon
350 hydrogen bonding interaction with Arg278 (3.58 Å) shown in
351 (Fig. 8B), similar to β -tubulin-Cd metal complex (Fig. 8A).
352 Also, β -tubulin-Co metal complex shows non bonded inter-
353 action such as Thr220, Thr223, Ser277, Gln282 forms van der
354 Waals, Asp226 forms π -anion, His-229 makes π - π T-shaped
355 interaction and Leu217, Lys218, Leu219, Pro274, Leu286,
356 Leu371 and Pro360 forms hydrophobic alkyl and π -alkyl types
357 of interactions with drug-Cd as shown in Fig. 8B.

358 The analysis of β -tubulin-Cu metal complex shows the
359 hydrogen bonding interactions of Thr276 (1.95 and 2.91 Å)
360 (Table-5) and carbon bonding interactions Ala233 (3.63 Å)
361 and Arg278 (3.83 Å) shown in (Fig. 8C). In addition, Arg278
362 makes π -donon bonding interaction. Leu371 makes π -sigma
363 bonding, Asp226 forms π -anion, His229 forms π - π T-shaped
364 interactions with drug-Cu metal complex and Leu217, Leu219
365 and Pro274 forms hydrophobic alkyl and π -alkyl type of
366 interactions as shown in Fig. 8C. Next, analysis of β -tubulin-
367 Ni metal complex show bonding interactions with Thr276 (1.70
368 Å) and carbon bonding interactions Asp226 (3.15 Å) as shown
369 in Fig. 8D. Leu217, Thr220, Thr223, His229, Arg284, Gln282,
370 Gly370 and Leu371 forms van der Waals interactions. Asp226
371 forms π -anion, Lys281 forms amide- π stacked as shown in
372 Fig. 8D. While, Leu219, Arg278, Pro360 forms hydrophobic
373 type of alkyl and π -alkyl type of interactions as shown Fig. 8D.

374 Conclusion

375 A novel quinoline Schiff base ligand and its metal(II)
376 complexes were successfully prepared and characterized. The
377 synthesized Schiff base and its metal(II) complexes were
378 screened for antibacterial, antifungal and cytotoxicity activities.
379 Among the prepared compounds, the Schiff base ligand and
380 its Cd(II) complex showed an excellent activity against all four

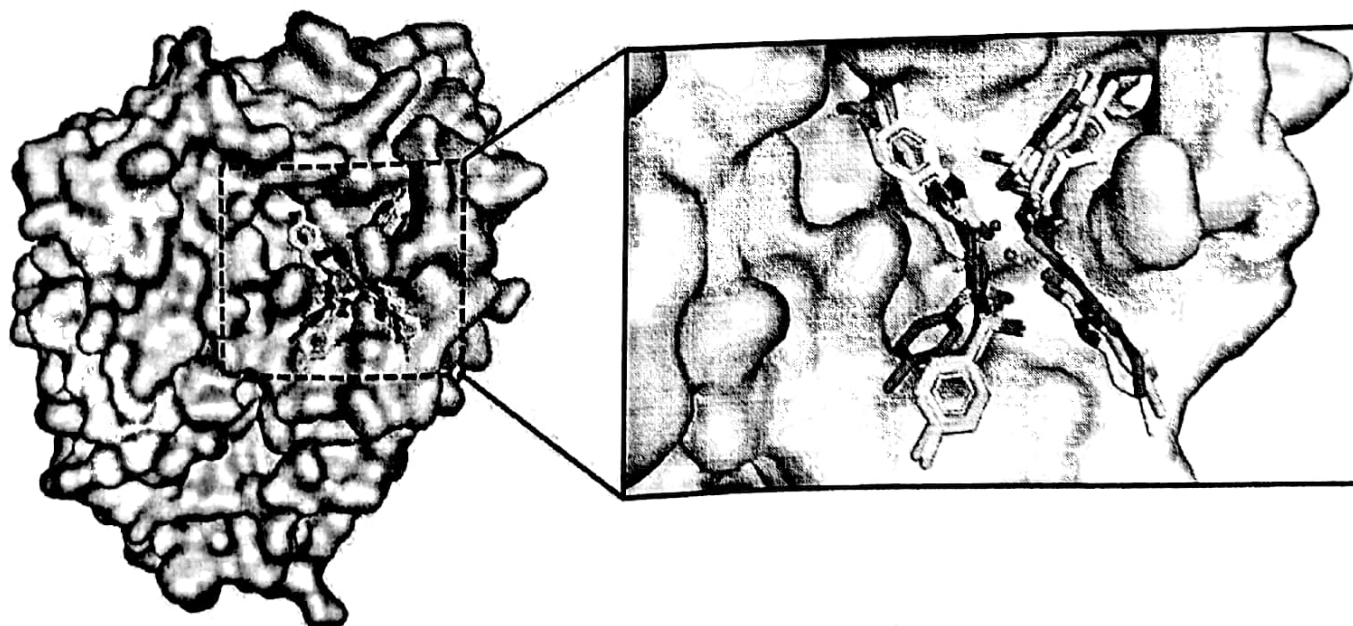


Fig. 7. Molecular docking of β -tubulin with drug-metal complexes. Here, (A) shows the overlapped docked conformation of Cd (green), Co (cyan), Cu (magenta) and Ni (yellow) metal complexes, the atoms such as N, O and H are shown in blue, red and white colour, respectively. (B) Zoomed view of β -tubulin binding pocket with overlapped conformation of metal complexes

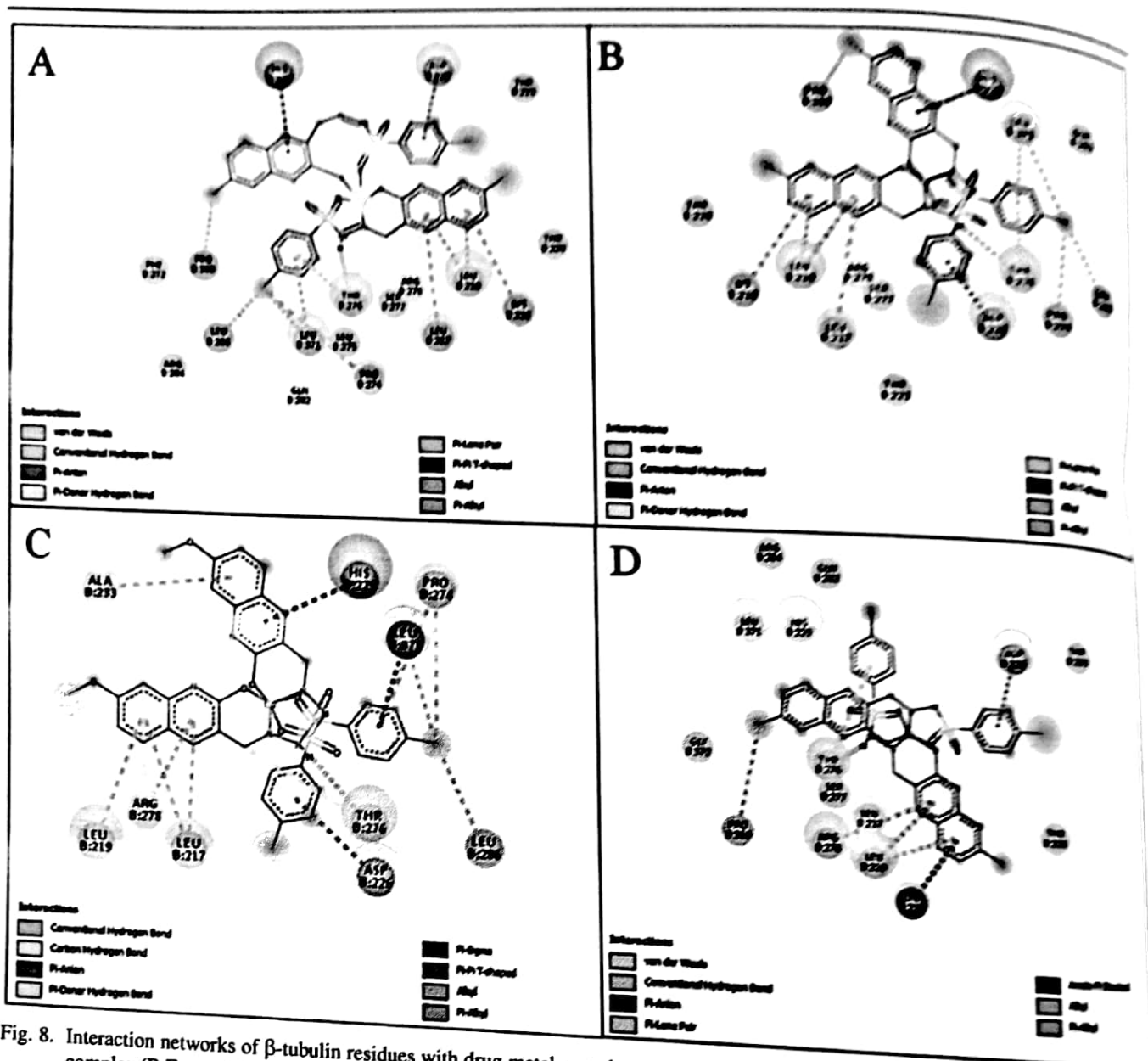


Fig. 8. Interaction networks of β -tubulin residues with drug metal complexes. (A) Interaction network of β -tubulin residues with Cd-metal complex (B.E. = -11.32 kcal/mol), (B) Interaction network of β -tubulin residues with Co-metal complex (B.E. = -10.70 kcal/mol), (C) shows interaction of β -tubulin with Cu-metal complex (B.E. = -12.39 kcal/mol) and (D) Interaction network of β -tubulin with Ni-metal complex (B.E. = -11.39 kcal/mol). The β -tubulin shows higher binding affinity for Cu-metal complex compare to other tubulin and metal complexes, this result is in agreement with the experimental observation. The 2D interaction images were generated using the Discovery studio Visualizer [46]

381 bacterial strains. In case of antifungal activity ligand and Cu(II)
 382 complex exhibited good activity against *A. clavatus* fungal
 383 strain. Other compounds were found to be less active against
 384 both bacterial and fungal strains. Furthermore, all the comp-
 385 ounds were screened for *in vitro* cytotoxicity against two human
 386 cancer cell lines and results showed that the Cu(II) complex
 387 was found to be more active among the prepared compounds.
 388 The Schiff base ligand, Co(II), Ni(II) and Cd(II) compounds
 389 showed excellent activity against the human lung cancer cell
 390 line (A-549). From the overall study, it was concluded that all
 391 compounds have excellent cytotoxicity properties compared
 392 to the standard drug paclitaxel. Ligand and Cd(II) complex
 393 has excellent antibacterial activity compared to the standard

drug Ampicillin. Furthermore, the binding modes and interactions of metal complexes with β -tubulin receptor protein are confirmed by molecular docking study. The docking study revealed that all the metal(II) complexes show excellent binding affinity at the paclitaxel site of the β -tubulin. Hence, it is concluded that prepared compounds possessed excellent cytotoxicity properties and could be used as potential lead for cancer treatment.

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CONFLICT OF INTEREST

The authors declare that there is no conflict of interests regarding the publication of this article.

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