



Synthesis, Cytotoxicity, and Molecular Docking Study of Complexes Containing Thiazole Moiety

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Abstract: The ligand 5-methyl-2-phenyl-4-[(E)-1,3-thiazol-2-yl-diazenyl]-2,4-dihydro-3H-pyrazol-3-one (Dy) has been synthesized by diazo coupling reactions of 5-methyl-2-phenyl-2,4-dihydro-3H-pyrazol-3-one with 2-aminothiazole and ferric hydrogen sulfate (FHS), as a catalyst, under solvent-free conditions. A series of complexes of the ligand with Co(II), Ni(II), Cu(II), and Zn(II) ions are synthesized and structurally characterized by ¹H NMR, FTIR, and UV-Visible spectral techniques. The cytotoxic activity of the complexes and the uncoordinated ligand against human breast cancer (MCF-7) and chronic myelogenous leukemia cell line (human erythroleukemia) (K-562) cell lines exhibits good viability in the range of 50.16–55.16% at a concentration of >100-110 µg/mL as compared to the inhibition in the untreated cells. Further, the metal complexes and ligand were screened against antibacterial strains of *S. typhi*, *S. aureus*, and *E. coli*. Both the cytotoxicity and antioxidant studies are correlated with computational docking analysis and powder XRD studies reveals that all complexes are in crystalline nature.

Keywords: Azo dye, solvent-free, Fe(HSO₄)₃, powder X-ray diffraction.

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INTRODUCTION

Diazo coupling reactions is one of the most important class of organic reactions which is currently performed with liquid acid catalysts, such as H₂SO₄, HF, HBr, HCl, CF₃COOH, and complexes of BF₃ are frequently used in organic synthesis (1-3). This traditional approaches are inherently associated with environmental issues like pollution, catalyst waste, and issue in separation and recuperation (4). To overcome these issues an economical catalyst is employed (5). Current days the azo dyes grab much attention from synthetic organic chemists, due to its numerous groups of dyes consumed today like FMCG products such are textile, paper, coloring agents for foods and cosmetics industries. Other applications include emerging technologies like liquid crystals, organic photoconductors and non-linear optics (6-8). Azo dyes have got much attentions from different branches of chemistry and they show more prominent applications like colorants in the textile and digital printing, photography industries (9-11) liquid crystals (12) chiral switches in photochemistry (13) and for molecular recognition (14). Azo dyes exhibit important pharmacological activities like anti-inflammatory (15), plant growth regulating properties (16). Antipyretic (17) and antitumor (18, 19) are also reported. The metal complexes derived from azo dyes have been investigated by many workers (20-23) and their complexes have been extensively used in many industries like dyes, synthetic leather and vinyl polymers. The applications of metal complexes of dyes not limited for industries but also in the field of pharmacology such as inhibition of DNA, RNA, and protein synthesis, nitrogen fixation, and carcinogenesis (24, 25). 5-Methyl-2-phenyl-2,4-dihydro-3H-pyrazol-3-one is the biological potent organic moiety (26, 27). Many researchers work on metal chelate of 5-methyl-2-phenyl-2,4-dihydro-3H-pyrazol-3-one (28-32) but the catalytic synthesis of dyes from this organic moiety is not reported.

The aim of the present study was to synthesize the 5-methyl-2-phenyl-4-[(E)-1,3-thiazol-2-yl-diazenyl]-2,4-dihydro-3H-pyrazol-3-one (Dy) via the diazotization reaction of 5-methyl-2-phenyl-2,4-dihydro-3H-pyrazol-3-one with 2-aminothiazole using the iron(III) hydrogen sulfate, Fe(HSO₄)₃, as an economical catalyst and their metal chelates, screening for *in-silico* molecular docking studies followed by antimicrobial and cytotoxic activities.

EXPERIMENTAL

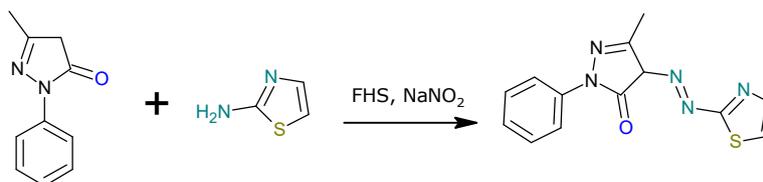
Materials and methods

All the chemicals used in this study were pure and AR grade. 2-Aminothiazole, ethanol, DMSO, were procured from Sigma–Aldrich and all the metal salts were obtained from Hi-Media and the solvents were used after purification by distillation (33). The human breast cancer (MCF-7) cell line was obtained from National center for cell science (NCCS), Pune, India. FTIR spectra were recorded as KBr pellets in the Bruker Alpha FT-IR Spectrometer, ¹H NMR were recorded on a Bruker DPX 400, δ values relative to the deuterated DMSO. Magnetic susceptibilities were

Shafeeulla R, Krishnamurthy, Bhojynaik, Manjuraj, JOTCSA. 2017; 4(3): 787-810. **RESEARCH ARTICLE** measured at room temperature by the Gouy method. The molar conductance measurements were measured in solution of the metal complexes in DMF (10^{-3} M) using Equip-tronics EQ-660A conductivity meter, Mass spectra are recorded in a Quattro LC, Micro Mass spectrometer. Elemental analysis is obtained from a Vario-Micro Qub elemental analyzer. Absorbance is measured using Systronics UV-VIS spectrophotometer-119, X-ray diffractometer (PHILIPS PW3710) using CuK α (1.5418 \AA) radiation operated at 45 kV and 25 mA is used in X-ray investigations.

Synthesis of 5-methyl-2-phenyl-4-[(E)-1,3-thiazol-2-yl-diazenyl]-2,4-dihydro-3H-pyrazol-3-one (Dy)

To a stirred mixture of NaNO₂ (2 mmol, 0.28 g) and silica gel (1 g), a suspension of an appropriate 2-aminothiazole (2 mmol) and ferric hydrogen sulfate (FHS) (10 mol%, 0.06 g) in water (12 mL) was added. After the mixture was stirred for about 20-30 min, the solution of 5-methyl-2-phenyl-2,4-dihydro-3H-pyrazol-3-one (2 mmol, 0.6 g) was added. The progress of the reaction was monitored by TLC using chloroform: methanol as eluent (9:1). After the completion of the reaction, the mixture was filtered and the residue was washed several times with hot water and product is recrystallized in ethanol. The synthetic route is represented in Scheme 1.

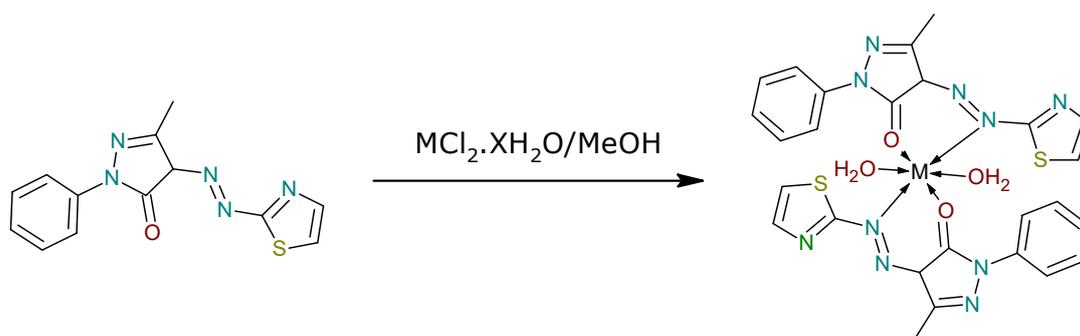


Scheme 1: Synthesis of 5-methyl-2-phenyl-4-[(E)-1,3-thiazol-2-yl-diazenyl]-2,4-dihydro-3H-pyrazol-3-one (Dy).

5-Methyl-2-phenyl-4-[(E)-1,3-thiazol-2-yl-diazenyl]-2,4-dihydro-3H-pyrazol-3-one (Dy): Brick red, Yield :89%, m.p. 148-150 °C. (Anal.: C, 54.7; H, 3.7; N, 24.5; O, 5.6; S, 11.2; C₁₃H₁₁N₅OS calc.: C, 55.6; H, 3.8; N, 23.4; %). FT-IR (ATR, cm⁻¹): $\nu(\text{C=O})$ 1668, $\nu(\text{C=N})$ 1594, $\nu(\text{N=N})$ 1505, $\nu(\text{C-S})$ 759. ¹H-NMR (400 MHz, CDCl₃, ppm): 7.91–7.20 (m, 7H, Ar-H.), 2.84 (s, 3H, CH₃).

Synthesis of metal complexes 1-4.

A methanolic solution of metal chlorides (0.461 g, 1.5 mmol) in 20 mL was added drop wise to a methanolic solution (20 mL) of the ligand (1.044 g, 3 mmol) with continuous stirring (Scheme 2). The resulting solution was refluxed for 4-6 h and the solution was reduced to half of its initial volume. It was then allowed to stand overnight in a refrigerator. A colored complex was precipitated, which was separated by filtration under vacuum. It was washed thoroughly with distilled water and cold methanol and dried *in vacuo* over fused CaCl₂ and was recrystallized from methanol (35, 36).



Scheme 2: Synthesis of metal complexes (M = Cu, Co, Ni, Zn).

Result for **1**: Light brown; yield 68%; mp, >300 °C; mw, 670.2 g/mol; found for C₂₆H₂₄CuN₁₀O₄S₂: C: 46.72; H: 3.62; N: 20.96, Calc. for C: 45.17; H: 3.28; N: 20.22. UV-Vis (DMF): λ_{max} (log ε) = 20,000. FT-IR (cm⁻¹):ν(C=O) 1603, ν(C=N) 1586, ν(N=N) 1581, ν(C-S) 755, ν(M-O) 509, ν(M-N) 477. ΛM = 130 Ω⁻¹.cm² mol⁻¹

Result for **2**: Black brown; yield 63%; mp, >300°C; mw, 665.6 g/mol; found for C₂₆H₂₄CoN₁₀O₄S₂: C: 47.16; H: 3.65; N: 21.10, Calc. for C: 47.30; H: 3.44; N: 20.96. UV-Vis (DMF): λ_{max} (log ε) = 14,705, 16,395. FT-IR (cm⁻¹):ν(C=O) 1630,ν(C=N) 1583, ν(N=N) 1492, ν(C-S) 762, ν(M-O) 511, ν(M-N) 473. ΛM = 128 Ω⁻¹.cm².mol⁻¹.

Result for **3**: Dark brown; yield 62%; mp, >300°C; mw, 665.3 g/mol; found for C₂₆H₂₄NiN₁₀O₄S₂: C: 47.09; H: 3.70; N: 21.72, Calc. for C: 47.08; H: 3.84; N: 21.36. UV-Vis (DMF): λ_{max} (log ε) = 15,000, 20,000, 25,00. FT-IR (cm⁻¹):ν(C=O) 1606, ν(C=N) 1586, ν(N=N) 1477, ν(C-S) 757, ν(M-O) 510, ν(M-N) 477. ΛM = 132 Ω⁻¹.cm².mol⁻¹

Result for **4**: Pale brown; yield 59%; mp, >300°C; mw, 672.0 g/mol; found for C₂₆H₂₄N₁₀O₄S₂Zn: C: 46.60; H: 3.65; N: 20.90, Calc. for C: 46.89; H: 3.46; N: 20.28. FT-IR (cm⁻¹):ν(C=O) 1607, ν(C=N) 1582, ν(N=N) 1485, ν(C-S) 757, ν(M-O) 509, ν(M-N) 474. ΛM = 125 Ω⁻¹.cm².mol⁻¹.

Molecular docking studies

Molecular docking for the synthesized compounds is done by as per our previous work (37, 38). For antioxidant and cytotoxic molecular docking study we use (PDB code: 3MNG) as antioxidant enzyme receptor and (PDB code: 2A91) as EGFR kinase domain was used throughout the work. Followed by butylated hydroxytoluene (BHT) and Actinoin as standards for docking studies. The docking results of both antioxidant and cytotoxic guided for wet analysis.

Antimicrobial activity

The antibacterial activity was determined using the agar disc diffusion method against four Gram-negative and Gram-positive bacterial strains. The nutrient agar (18 g/L) medium at 4 °C were maintained for microorganisms. The agar media was prepared by suspending 2 g nutrient

agar in 100 mL of distilled water. The pH maintained to neutral value and finally autoclaved, after some interval of time to bring down the media temperature to 45 °C. The test compounds were dissolved in DMF and 5 µL of each compound with final concentrations of 10.0 µg/disc, 25.0 µg/disc and 50 µg/disc was poured on 4-mm sterile paper discs and placed on nutrient agar plates. In each plate DMF served as negative control and the standard antibacterial drug cefexime (1 mg/mL) served as positive control.

Three plates of each bacterial strain were prepared and incubated at 37 °C for 24 h. The antibacterial activity was determined by measuring the zones of inhibition (mm). The same procedure was followed for the antifungal assay against *A. niger*, *A. flavus*, *A. fumigates*, and potato dextrose agar (PDA) was the media (39). Fungal spores were spread on the surface of each agar plate instead of the bacterial culture and the plates were incubated at 28 °C for 24 h. In each plate DMF served as negative control and Fluconazole (1 mg/mL) was used as positive control. The experiments were carried out in triplicate. The lowest concentration at which inhibition was shown was considered as minimum inhibitory concentration (MIC) (40, 41).

Antioxidant activity

DPPH radical scavenging method

Free radical-scavenging capacities of the ligand (Dy) and their metal complexes were determined using the stable 2,2-diphenyl-1-picrylhydrazyl radical (DPPH). An aliquot of 25-75 mg concentrations of synthesized compounds in methanol was added to 3 mL of 0.004% w/v DPPH radical solution and each test tube was made up to final volume of 4 mL. butylated hydroxytoluene (BHT) was used as a standard and dissolved in methanol to get the same concentration as that of synthesized compounds. Each mixture was vortexed for a few seconds and allowed to stand in the dark for 10 minute at ambient temperature. The absorbance of each reaction mixture was measured at 517 nm against a blank.

Nitric oxide (NO) scavenging method

All the tested samples in various concentrations (25, 50 and 75 µg/mL) were prepared in DMF and made homogeneous. 0.75 mL of phosphate buffer saline (0.2 M, pH 7.4) and 0.5 mL of sodium nitroprusside (10 mM) were mixed together to generate nitric oxide radicals (NO) and this solution were added to the test compounds and incubated for 120 min at 25 °C. The reaction mixture of the above samples (1 mL each) was treated with 1 mL of Griess reagent (1% sulfanylamide, 2% H₃PO₄ and 0.1% naphthylene diamine dihydrochloride). Butylated hydroxytoluene (BHT) was used as a positive control. The absorbance was measured at 546 nm. The ability to scavenge from both radicals was calculated as follows:

$$\text{Scavenging ratio (\%)} = [(A_i - A_o) / (A_c - A_o)] \times 100\%$$

Where A_i is the absorbance in the presence of the test compound, A_o is absorbance of the

blank in the absence of the test compound, A_c is the absorbance in the absence of the test solution.

In-vitro anticancer activity - Cell culture

The human cancer cells (K-562 ATCC® CCL-243™) and (MCF7-ATCC® HTB 22™) were maintained in Modified Eagles Medium (MEM) supplemented with 10% FCS, 2% essential amino acids, 1% each of glutamine, non-essential amino acids, vitamins and 100 U/mL Penicillin-Streptomycin. Cells were sub-cultured at 80–90% confluence and incubated at 37 °C in a humidified incubator supplied with 5% CO₂. The stock cells were maintained in 75 cm² tissue culture flask.

Cell viability assay

The cytotoxicity effect of test samples was performed by 5-diphenyl-2H-tetrazolium bromide (MTT) assay. Briefly, cultured cells (1×10^6 cells/mL) were placed in 96 flat-bottom well plates, then the cells were exposed to different concentrations of prepared samples (1–100 µg/mL) and incubated at 37 °C for about 24 h in 5% CO₂ atmosphere. After 24 h incubation, MTT (10 µL) was added to the incubated cancer cells and further incubated at 37 °C for about 4 h in same environment. Thereafter, we dissolved 200 µL of formazan crystals in DMSO and monitored the absorbance at 578 nm with reference filter as 630 nm. The cytotoxicity effect was calculated as:

$$\text{Cytotoxicity}(\%) = 1 - \left(\frac{\text{mean absorbance of}}{\text{mean absorbance of - ve}} \right) \times 100$$

$$\text{Cell viability}(\%) = 100 - \text{Cytotoxicity}(\%)$$

RESULTS AND DISCUSSION

Mass spectrum

The mass spectrum of the ligand (Figure 1) obtained by using of Quattro LC, Micro mass spectrometer (ESI) showed the molecular ion peak at m/z 285.0 which coincide with the formula weight within the precision limit of ± 0.02 . Metastable ion(s) were not observed, thus representing the proposed fragmentation pattern of the ligand (Scheme 3).

¹H NMR spectrum

The ¹H NMR spectrum (Figure 2) of the ligand in DMSO-d show a doublet at 7.91 and 7.88 ppm shifts is for H1 and H2 protons respectively. The doublet at 7.59-7.58 ppm is assigned to the H6 proton. The other two protons H5 and H7 resonated together but was slightly distorted by coupling effect from H6 and H8 which appeared as a multiplet peak at 7.47 ppm and a doublet peak showed at 7.34 ppm assigned for H4 and H8 protons, at the extreme right of the spectrum a singlet signal appeared at 2.84 ppm is assigned for methyl group H3 proton. However, the chemical shifts and multiplicity patterns are in agreement with the structure of the compound (Figure 3).

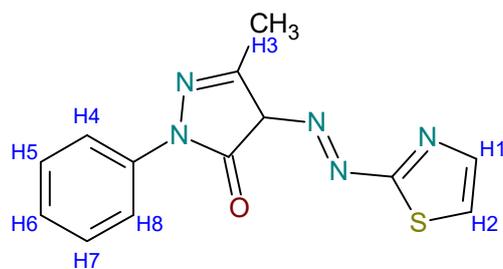


Figure 3: The structure of the ligand for H NMR.

FTIR spectroscopy

A comparative FTIR spectral study of the ligand and its metal complexes has been recorded as KBr pellets. Some significant IR bands for the uncoordinated ligand and their metal complexes have been listed in Table 1. The free ligand showed both a strong band and a weak band at 1668 cm^{-1} and 3090 cm^{-1} which were attributed to $\nu(\text{CO})$ and $\nu(\text{C-H})$ aromatic (42). The IR spectrum of the ligand displays bands at 1594 cm^{-1} which may be assigned to the $\nu(\text{C=N})$ thiazole ring and the strong band belongs to $\nu(\text{N=N})$ was found at 1501 cm^{-1} (43). The hypsochromic shift was observed for the $\nu(\text{CO})$ band at 1668 cm^{-1} assigned in the spectrum of the ligand. This carbonyl band showed reduction in intensities between 1668 and 1603 cm^{-1} region observed due to the coordination to central metal ions (44). However, the $\nu(\text{N=N})$ band at 1501 cm^{-1} in the free ligand appeared to have shifted to lower frequencies in the spectrum of the metal complexes due to donation of the lone pair of electron present on the nitrogen atom to the metal ions in the complex. After complexation, the peaks appeared almost in the same region between 1594 - 1586 cm^{-1} and 759 - 757 cm^{-1} shows that the electrons from the nitrogen and sulfur of the thiazole ring are not donated for bonding with metal ion (45, 46). All the complexes show broad bands in the region 3434 - 3439 cm^{-1} which may be assigned to $\nu(\text{O-H})$ of coordinated water (47). This suggests that the nitrogen and sulfur atom of the thiazole ring are not participated in the coordination. This was further supported from existence of new metal- ligand band in the complex spectrum ranging from 509 - 511 cm^{-1} for M-O band and 474 - 477 cm^{-1} for M-N band and the elemental analysis as shown in Table 2.

Table 1: IR spectral data (cm⁻¹) of Ligand (L) and metal complexes.

Ligand/complex	$\nu(\text{C}=\text{O})$	$\nu(\text{C}=\text{N})$	$\nu(\text{N}=\text{N})$	$\nu(\text{C}-\text{S})$	$\nu(\text{M}-\text{O})$	$\nu(\text{M}-\text{N})$
Ligand (L)	1668	1594	1501	759	-	-
[CuL ₂ H ₂ O]Cl ₂ (1)	1603	1586	1481	757	509	477
[CoL ₂ H ₂ O]Cl ₂ (2)	1630	1583	1492	762	511	473
[NiL ₂ H ₂ O] Cl ₂ (3)	1606	1586	1477	757	510	477
[ZnL ₂ H ₂ O]Cl ₂ (4)	1607	1582	1485	757	509	474

Table 2: Elemental analyses, colors, melting points and yield of Ligand and metal complexes.

Ligand/Complex	Color	Melting point (°C)	Found (Calc.) %				Yield (%)	$\Delta M S \text{ cm}^2 \text{ mol}^{-1}$
			C	H	N	M		
Ligand (Dy)	Brick red	148-150	54.72 (55.65)	3.76 (3.86)	24.55 (23.46)	-	89	-
1	Light brown	>300	46.72 (45.17)	3.62 (3.28)	20.96 (20.22)	9.86 (9.62)	68	130
2	Black brown	>300	47.16 (47.30)	3.65 (3.44)	21.10 (20.96)	9.70 (9.89)	63	128
3	Dark brown	>300	47.09 (47.48)	3.70 (3.84)	21.72 (21.36)	9.67 (9.30)	62	132
4	Pale brown	>300	46.60 (46.89)	3.65 (3.46)	20.90 (20.28)	9.76 (9.48)	59	125

Electronic spectra and magnetic moment measurements

The UV-Vis spectrum of the ligand and their metal complexes are recorded in DMF and the values are reported in Table 3. The free ligand showed two recognizable absorption bands at 36,101 and 26,315 cm^{-1} would be due to the $\pi\text{-}\pi^*$ and $n\text{-}\pi^*$ transitions, respectively. The spectrum of the Cu(II) complex **1** showed a absorption band at 20,000 cm^{-1} which is attributed to the transition ${}^2E_g \rightarrow {}^2T_g$ due to distorted octahedral geometry (48, 49). Further confirmation was done by magnetic moment values 1.78 B.M. which is consistent with proposed distorted octahedral geometry for Cu(II) complex. The Co(II) complex **2** showed absorption band at 14,705 cm^{-1} and 16,393 cm^{-1} due to the transitions ${}^4T_{1g}(F) \rightarrow {}^4A_{2g}(F)$ and ${}^4T_{1g}(F) \rightarrow {}^4A_{2g}(P)(\nu_2)$, respectively, the Co(II) complex also showed magnetic moment at 4.56 B.M. These results suggested the presence of octahedral geometry for Co(II) complex. (50, 51). The Ni(II) complex **3** showed three bands at 25,000 cm^{-1} , 20,000 cm^{-1} and 15,384 cm^{-1} which are assignable to ${}^3A_{2g}(F) \rightarrow {}^3T_{2g}(F)(\nu_1)$, ${}^3A_{2g}(F) \rightarrow {}^3T_{1g}(F)(\nu_2)$ and ${}^3A_{2g}(F) \rightarrow {}^3T_{1g}(P)(\nu_3)$ transitions respectively. The spectral data indicate that the Ni(II) ion coordinated through octahedral geometry. The magnetic moment of Ni(II) complex is 2.74 B.M. These results suggested the presence of octahedral geometry for Ni(II) complex (52, 53).

Powder X-ray diffraction study of complexes

Powder X-ray diffraction graph of the complexes were recorded over the $2\theta = 0\text{-}80^\circ$ range. The major peaks of relative intensity greater than 10 % were indexed using a computer software. The diffraction data like angle (2θ), interplanar spacing (d) and relative intensity (%) have been summarized in (Table 4) from the data all **1-4** show sharp crystalline peaks indicating their crystalline nature. The XRD patterns of all complexes are very similar and suggest that the complexes have similar structure. The average crystallite sizes of the complexes d_{XRD} were calculated using the Sherrers formula [54]. **1**, **2**, **3** and **4** have a crystallite size of 28.25, 28.16, 15.04 and 42.06 nm, respectively, suggesting that the complexes are in a nano-crystalline phase and the X-ray diffraction data of **1** are shown in Table 5 and spectrum in Figure 4.

Table 3: Electronic spectral data (cm⁻¹) and magnetic studies for Ligand and its metal complexes.

Entry	Transitions (cm ⁻¹)	Transitions	M _{eff} (BM)
Ligand (Dy)	36,101 26,315	π-π* π-π*	-
1	20	² E _g → ² T _{2g}	1.78
2	16,395 14,705	⁴ T _{1g} (F) → ⁴ A _{2g} (P) (ν ₂) ⁴ T _{1g} (F) → ⁴ A _{2g} (F)	4.56
3	25,385 20,000 15,000	³ A _{2g} (F) → ³ T _{2g} (F) (ν ₁) ³ A _{2g} (F) → ³ T _{1g} (F) (ν ₂) ³ A _{2g} (F) → ³ T _{1g} (P) (ν ₃)	2.74
4	Diamagnetic		Diamagnetic

Table 4: PXRD spectral data of 1-4.

Complexes	2θ	d	Relative intensity	Full width at half maximum
1	26.42	3.37	100	0.32
2	21.32	4.16	100	0.20
3	19.74	4.49	100	0.56
4	18.61	4.76	100	0.20

Table 5: X-ray diffraction data of **1**.

Peak no	2 θ	θ	Sin θ	h k l	d		Intensity	a in Å
					Cal	obs		
1	12.382	6.19125	0.1078	1 5 2	7.14243	7.1464	339.6	3.62
2	13.343	6.6719	0.1161	1 9 1	6.62998	6.6340	451.39	3.62
4	14.424	7.21225	0.1255	2 0 6	6.13561	6.1355	267.93	3.62
5	15.708	7.8543	0.1364	9 7 4	5.63681	5.6313	196.12	3.62
6	17.150	8.57515	0.1491	9 6 9	5.16609	5.1655	336.55	3.62
7	17.601	8.8005	0.1529	8 2 2	5.03479	5.0351	194.86	3.62
8	18.619	9.3097	0.1617	1 3 3	4.76165	4.7634	195.50	3.62
9	19.225	9.6128	0.1669	1 7 2	4.61285	4.6142	382.53	3.62
10	21.789	10.8947	0.1890	1 2 1	4.07554	4.0754	211.51	3.62
11	23.237	11.6189	0.2014	2 1 6	3.8247	3.8261	508.84	3.62
12	25.309	12.6547	0.2190	2 1 0	3.51614	3.5161	442.26	3.62
13	26.420	13.2102	0.2285	3 7 2	3.37075	3.3716	974.94	3.62
14	29.318	14.65905	0.2530	1 0 4	3.04385	3.0448	125.80	3.62
15	36.042	18.0214	0.3093	8 0 2	2.48987	2.4900	74.07	3.62
16	44.889	22.4446	0.3817	5 6 8	2.01759	2.0177	28.34	3.62

Antimicrobial activity

The ligand and its metal complexes were tested on their antimicrobial activity against bacterial (*Salmonella typhi*, *Staphylococcus aureus*, *Escherichia coli*) and fungal (*Aspergillus niger*, *Aspergillus flavus*, *Aspergillus fumigatus*) strains by the agar disc diffusion method and results are given in Table 6a and 6b. Cefixime and roxithromycin are standard drugs and were used as references. The highest inhibition activities were observed for **1**, **2** and **3**. While the ligand and **4** did not show significant activity comparable to that of the standard drugs Cefixime at concentrations of 50, 25 and 10 µg/disc which represent the sensitive nature of the compounds. However, these complexes also tested for the fungal strains, exhibited least activity compared to standard fluconazole except **2** and **3**.

Antioxidant studies: DPPH and Nitric oxide radical scavenging methods

DPPH radical scavenging results of the synthesized ligand and metal complexes showed a moderate activity. The metal complexes exhibited more radical scavenging inhibition effects than that of the ligand. The **1** exhibited potent antioxidant effect nearly close the standard BHT and have potential to be applied as free radical scavengers and s free radical scavengers and **2** and **4** complexes showed condensed antioxidant activity and **3** and the ligand showed moderate activity. We found that the inhibitory effect of the compounds tested on NO is concentration-dependent and suppression ratio increases with increasing sample concentrations. The **1** is the most effective among all the complexes. The **2**, **3**, and the ligand showed better activity, while the 4 does not show any nitric oxide activity. Both methods of antioxidant activities are shown in Figure 5.

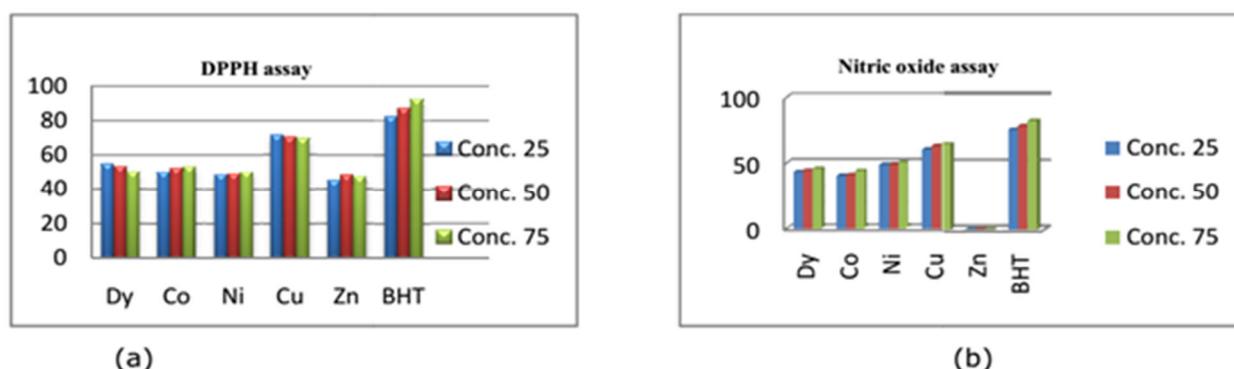


Figure 5: Antioxidant activity (a) DPPH method (b) Nitric oxide method.

Table 6a:- Antimicrobial activity of ligand and complexes.

Compound	Conc (µg/disc)	Zone of inhibition (mm)		
		<i>S. typhi</i>	<i>S. aureus</i>	<i>E. coli</i>
Ligand	25	12 ± 0.3	14 ± 0.8	11 ± 0.2
	10	10 ± 0.4	12 ± 0.3	09 ± 0.4
	50	22 ± 0.7	24 ± 0.8	23 ± 0.8
1	25	15 ± 0.5	20 ± 0.3	17 ± 0.6
	10	11 ± 0.4	17 ± 0.7	12 ± 0.3
	50	17 ± 0.2	19 ± 0.4	19 ± 0.9
2	25	16 ± 0.5	16 ± 0.3	15 ± 0.4
	10	10 ± 0.3	14 ± 0.3	11 ± 0.3
	50	22 ± 0.7	24 ± 0.8	22 ± 0.9
3	25	15 ± 0.8	18 ± 0.7	18 ± 0.5
	10	12 ± 0.5	15 ± 0.7	12 ± 0.7
	50	15 ± 0.7	15 ± 0.7	15 ± 0.3
4	25	12 ± 0.8	12 ± 0.7	13 ± 0.7
	10	09 ± 0.3	10 ± 0.7	11 ± 0.7
	50	25 ± 0.01	26 ± 0.1	24 ± 0.1
Cefixime	25	14 ± 0.1	23 ± 0.3	21 ± 0.3
	10	12 ± 0.1	19 ± 0.4	16 ± 0.1
	50	16 ± 0.5	20 ± 0.4	16 ± 0.8

Table 6b: Antifungal activity of ligand and complexes.

Compound	Conc (µg/disc)	Zone of inhibition (mm)		
		<i>A niger</i>	<i>A flavus</i>	<i>A fumigatus</i>
Ligand	50	14 ± 06	13 ± 0.8	11 ± 0.1
	25	12 ± 0.2	12 ± 0.5	10 ± 0.3
	10	10 ± 0.2	09 ± 0.4	09 ± 0.4
1	50	20 ± 0.6	18 ± 0.7	15 ± 0.8
	25	15 ± 0.2	14 ± 0.5	10 ± 0.1
	10	11 ± 0.4	10 ± 0.8	10 ± 0.5
2	50	17 ± 0.2	16 ± 0.7	15 ± 0.3
	25	16 ± 0.5	14 ± 0.5	13 ± 0.2
	10	10 ± 0.3	10 ± 0.2	11 ± 06
3	50	15 ± 0.8	14 ± 0.3	16 ± 0.1
	25	12 ± 0.5	11 ± 0.3	12 ± 0.6
	10	09 ± 0.3	08 ± 0.2	09 ± 0.6
4	50	16 ± 0.4	14 ± 0.7	12 ± 0.7
	25	13 ± 0.8	12 ± 0.5	11 ± 0.5
	10	10 ± 0.7	06 ± 0.8	07 ± 0.8
Fluconazole	50	24 ± 0.01	22 ± 0.1	23 ± 0.1
	25	16 ± 0.1	21 ± 0.3	22 ± 0.3
	10	10 ± 0.1	14 ± 0.4	10 ± 0.1

Molecular Docking study

The result of the docking studies have been carried between the human antioxidant enzyme receptor (PDB:3MNG) and the complexes are reported in Table 5a. The complexes exhibited comparable binding interaction energy (E-total) values in the in the range of -212.25 to - 327.01 kcal mol⁻¹. The complexes **1**, **2**, **3**, and **4** showed more negative binding values *i.e* 327.01, -280.78, -279.00 and -276.78 kcal mol⁻¹ respectively when compared with the standard butylated hydroxytoluene (E-total value -183.02 kcal mol⁻¹). The stability of the complexes **1** and **2** was evaluated by determining the hydrogen bonding interactions was evaluated by determining the hydrogen bonding interactions between amino acids of the protein molecules with the complex between amino acids of the protein molecules with the complexes. The result indicate that the complex **1** having the highest E-total values -337.07 kcal mol⁻¹ has greater binding affinity and better hydrogen bond interactions with the human antioxidant enzyme active site residues.

The molecular docking results showed in Figure 6 represent that the **1** has ability to form two hydrogen bonds with amino acid CYS72 with hydrogen bond distances 3.011 Å and 2.824 Å. Two more hydrogen bonds formed with amino acid PHE37 with bond distance of 2.751 Å and 2.520 Å. Single hydrogen bond can form between the **1** and the amino acids VAL39, LEU73 ALA71 and GLY38 of 2.945 Å, 2.944 Å, 2.660 Å, and 2.545 Å. The other interacting amino acids are CYS72, ALA71, GLY38, VAL70 and VAL35. The interaction of **2** represented in Figure 7 formed three hydrogen bonds with amino acids GLY38 with bond distances of 2.976 Å, 2.963 Å and 2.795 Å. Single hydrogen bond formed between the **2** and the amino acid ARG95, PHE37, SER74, LEU73, VAL70, CYS72, VAL35 with bond distance 3.064 Å, 3.057 Å, 3.200 Å, 2.967 Å, 2.733 Å, 2.638 Å and 2.654 Å The other interacting amino acids were CYS72, VAL35, HIS51, VAL35.

Table 5a- Antioxidant Docking scores.

Entry	Receptor PDB code	ΔG (kcal/mol) with MurB
Ligand	3MNG	-212.25
1	3MNG	-327.01
2	3MNG	-280.78
3	3MNG	-279.00
4	3MNG	-273.78
Butylated hydroxytoluene (STD)	3MNG	-183.02

Table 5b- Cytotoxic Docking Scores.

Entry	Receptor PDB code	ΔG (kcal/mol) with MurB
Ligand	2A91	-217.95
1	2A91	-323.66
2	2A91	-316.42
3	2A91	-313.96
4	2A91	-306.96
Actinoin (STD)	2A91	-235.99

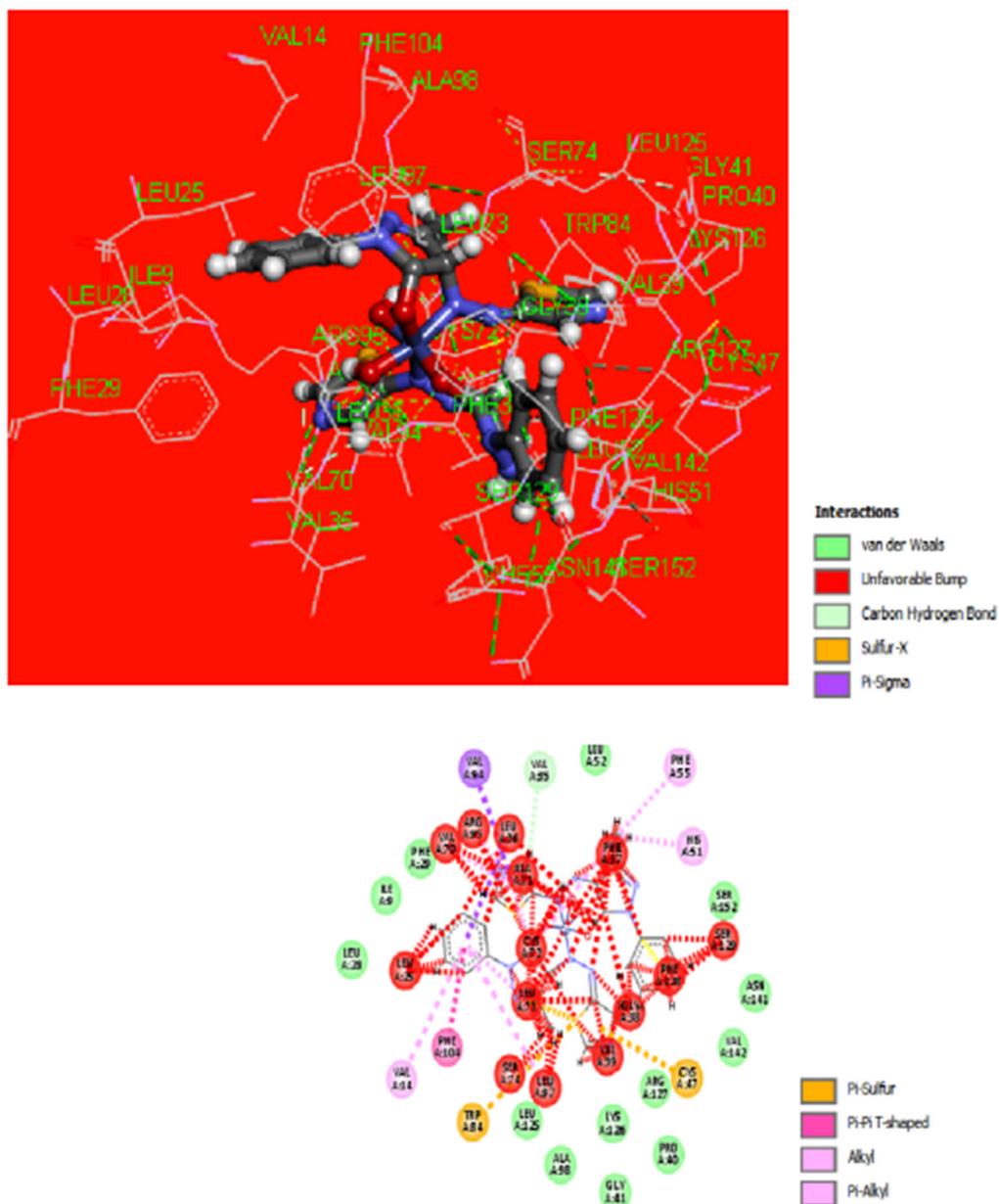


Figure 6: Interaction of 1 with amino acids of 3MNG (top) 3D-structure of complex (ball and stick model Oxygen-red, Nitrogen-blue) protein receptor (stick model) (bottom) 2D-structure of complex.

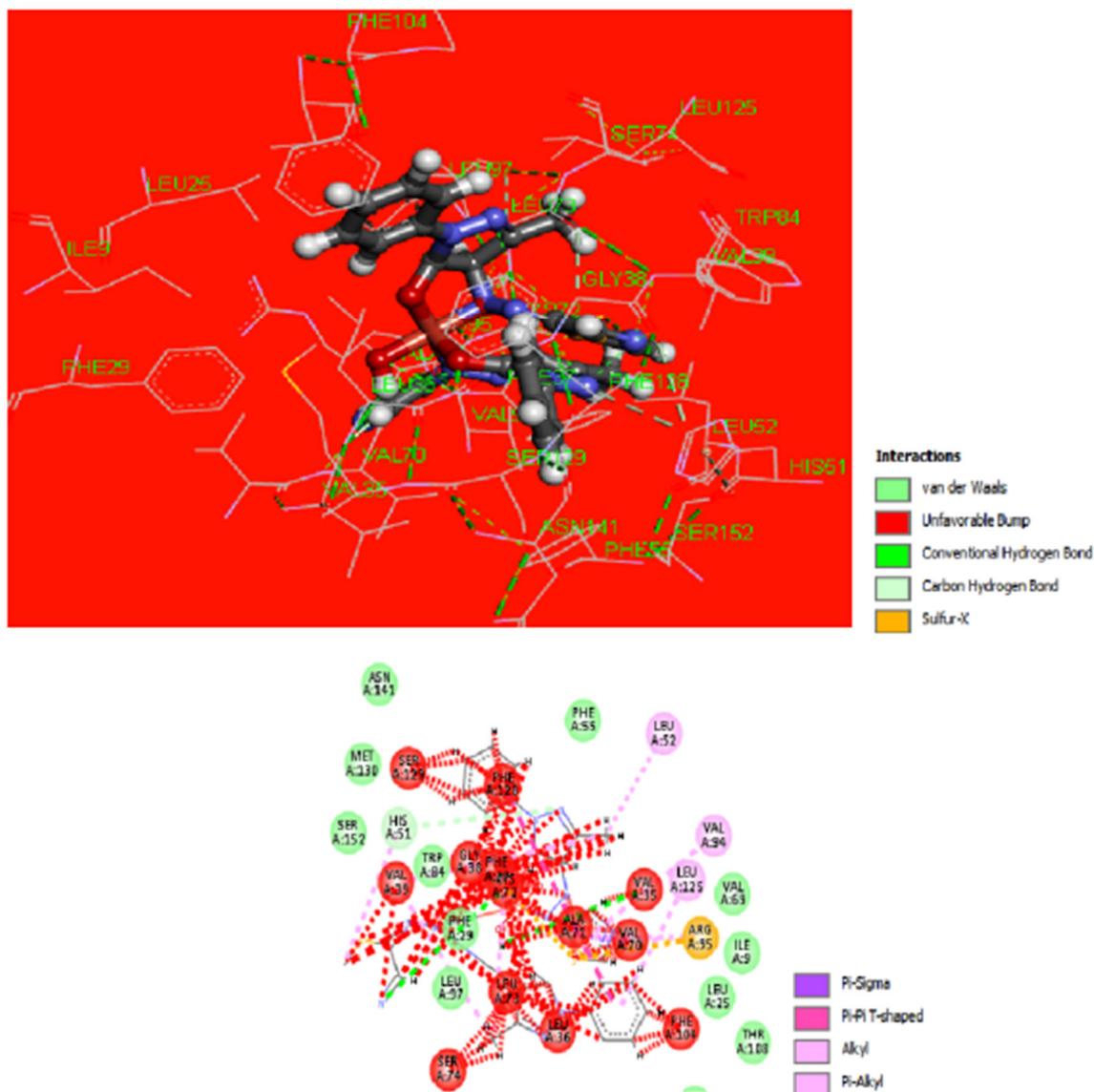


Figure 7: Interaction of 2 with amino acids of 3MNG (top) 3D-structure of complex (ball and stick model Oxygen-red, Nitrogen-blue) protein receptor (stick model) (bottom) 2D-structure of complex.

However, the 3 ($-279 \text{ kcal mol}^{-1}$) and the 4 ($-276.78 \text{ kcal mol}^{-1}$) have good docking energy minimum values than the standard but were not able form good hydrogen bonding interactions with human antioxidant enzyme receptor than **1** and **2**.

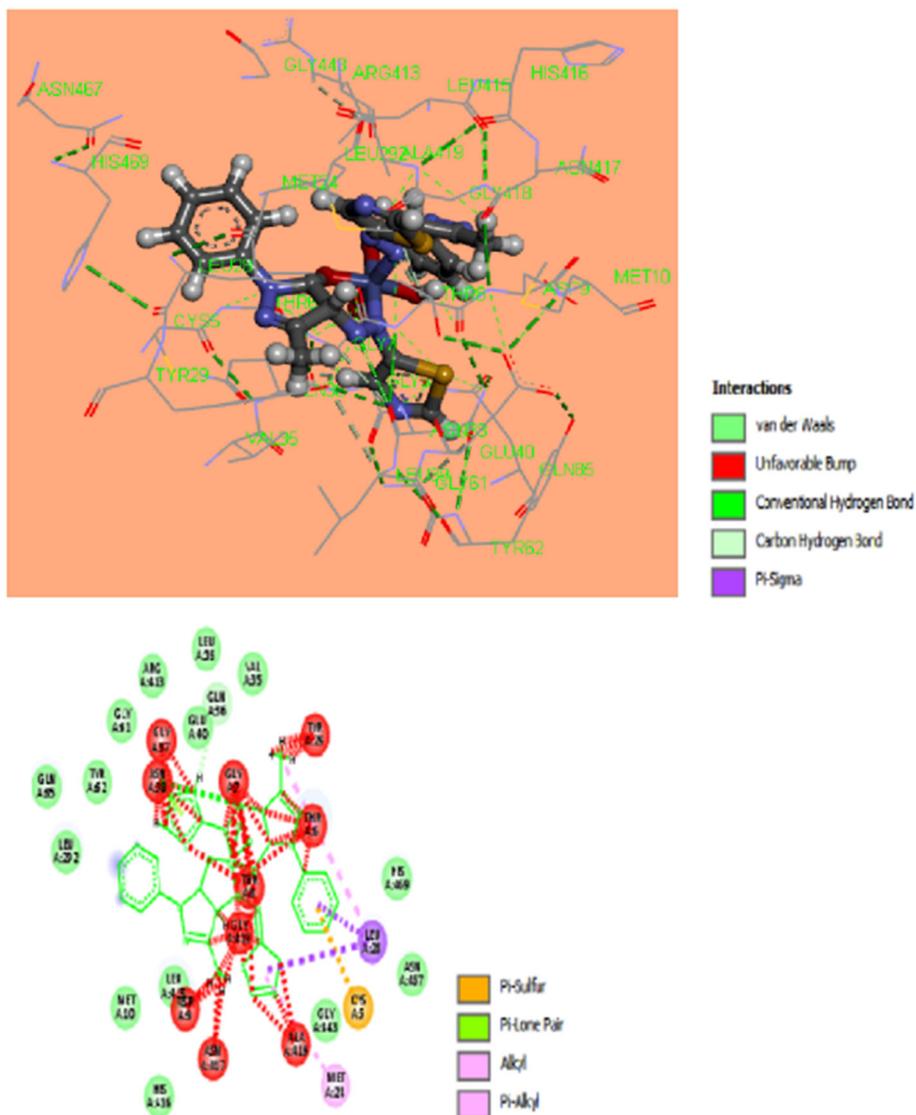


Figure 8: Interaction of 1 with amino acids of 2A91 (top) 3D-structure of complex (ball and stick model Oxygen-red, Nitrogen-blue) protein receptor (stick model) (bottom) 2D-structure of complex.

2.330 Å. Single hydrogen bonds were formed between the complex 1 and the amino acids THR6, GLY418 and ALA419 of 2.762 Å, 3.198 Å and 3.061 Å, respectively. The other interacting amino acids were ASN38, THR8 and GLN36 as shown in (Figure 8). The 2 formed two hydrogen bonds with amino acids GLY37 with bond distance 2.955 Å and 2.782 Å. A single hydrogen bonds were formed between the 2 and the amino acids ASN38, THR6, LEU39, GLY61, GLY418, ALA419, GLY7 and TYR420 with bond distance 3.193 Å, 3.020 Å, 2.995 Å, 2.904 Å, 2.577 Å, 2.764 Å and 2.538 Å. The other interacting amino acids were GLN36, ASN38 and TYR62 as shown in Figure 9. Even though the 3 (-313.42 kcal mol⁻¹) and 4 (-306.96 kcal mol⁻¹) are having good docking energy values than the standard actinoin (-260.51 kcal mol⁻¹) but these complexes are not forming good hydrogen bonding interactions with EGFR tyrosine kinase receptor than 1 and 2. The 3 and 4 interaction with active sites of EGFR tyrosine kinase with amino acid residues are LEU415, CYS32, TYR29, CYS5, and LEU28, VAL35, GLU40, ALA419, THR8. The binding value of 1 and 2 showed better cytotoxicity property than the remaining of complexes. The obtained results provide a sufficient explanation and good compromise between docking scores and *in-vitro* results of antibacterial and cytotoxicity activity.

Cytotoxicity

The results of cytotoxic activity is tabulated in Table 7 the reliable criteria for judging the efficacy of any anticancer drug are prolongation of life span, improving the clinical, haematological, biochemical profile, and reduction in viable tumor cell count in the host (41). In order to evaluate the biological effects of the ligand, and its 1, 2, 3 and 4 on cancer cells to treat human breast cancer (MCF7) and K-562 cell lines at the concentrations of 80, 90, 100, 110, 120, >130 and >200 µg/ mL for 48 h. The untreated cells were used as control. Cell growth inhibition was analyzed by the MTT assay and the results showed that the complexes and the ligand exhibited an inhibitory effect on the proliferation of MCF7 and K-562 cell lines in a dose-dependent manner. Among them, **1** and **2** showed the most potent inhibitory effect on the growth of both the MCF-7 and K-562 cells lines compared to the uncoordinated ligand.

Table 7: Cytotoxicity activity of ligand and complexes.

Tested samples	Inhibitory effect on cell lines (IC ₅₀)	
	MCF-7	K-562
Ligand	>130 µg/mL (50.38%)	>130 µg/mL (51.02%)
1	110 µg/mL (55.16%)	100 µg/mL (53.42%)
2	120 µg/mL (54.79%)	110 µg/mL (53.23%)
3	110 µg/mL (51.19%)	100 µg/mL (53.36%)
4	80 µg/mL (50.16%)	90 µg/mL (51.12%)

Note. Each value represents mean ±SD. The data were analyzed by Student's t test.

CONCLUSION

New biologically active Co(II), Ni(II), Cu(II) and Zn(II) ion complexes with 5-methyl-2-phenyl-4-[(E)-1,3-thiazol-2-yl-diazenyl]-2,4-dihydro-3H-pyrazol-3-one (Dy) are synthesized and structurally characterized. Except the Zn(II) complex, tentatively octahedral geometry is proposed for the other metal complexes. The newly synthesized Co(II) and Cu(II) metal complexes are potent antibacterial and antifungal agents against three bacteria (*Salmonella typhi*, *Staphylococcus aureus*, and *Escherichia coli*) and three fungi (*Aspergillus niger*, *Aspergillus flavus*, and *Aspergillus fumigates*). The synthesized uncoordinated ligand and metal complexes are screened for antioxidant activity, among them, Cu(II) Co(II) and Ni(II) complexes show fairly good activity in DPPH and Nitric Oxide scavenging. The results of computational docking studies promote to evaluate *in vitro* cytotoxic activity, The Cu(II) Co(II) and Ni(II), complex exhibited more *in vitro* cytotoxic activity than the unbound ligand and all complexes are of crystalline nature.

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