

Synthesis, Characterization and Antibacterial Activity of Schiff Bases Derived from 4-Dimethylaminobenzaldehyde with Some Amino Acids and 4-Aminoantipyrine toward Cu (II), Ni (II), Co (II), Cd (II) and Mn (II) Ions

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Abstract: A new chain of Schiff base complexes made up of Cu (II), Co (II), Cd (II), Ni (II), and Mn (II) ions derivatives were obtained by the condensation of 4-dimethyl-aminobenzaldehyde, and 4-aminoantipyrin and some amino acids. The structures of all synthesized complexes were established on the basis of their elemental analysis (UV, FTIR, and ¹HNMR). The study of X-ray crystallographic shows that the compound (Z)-4-((4-(dimethylamino)benzylidene)amino)-1,5-dimethyl-2-phenyl-1,2-dihydro-3H-pyrazol-3-one (**L₅**) crystallized in monoclinic system with space group of C2/c and the unit cell dimensions are a= 17.7916, b= 6.8610, c= 29.7199, α =90.000 β =101.326 γ = 90.000., Z = 8 and V = 3557.20 Å³. The antibacterial activity of the complexes are tested against *Bacillus subtilis*, *Bacillus cereus* and *Pseudomonas aeruginosa*.

Key Words: Schiff base, 4-dimethyl-aminobenzaldehyde, 4-aminoantipyrin, Schiff base complexes, *Bacillus cereus*, *Pseudomonas aeruginosa* and *Bacillus subtilis*.

I. Introduction

Schiff bases are an important class of ligands in management science and has a wide range of application in many fields [1]. Schiff bases obtained from the salicylaldehydes are popular polydentate ligands coordinated in neutral forms. Recently, metal complexes of Schiff bases are gaining popularity because of their antibacterial, antifungal and antitumor activities [2-3]. El-ajaily et al. (4) studied the antibacterial activities of the Schiff base derived from the salicylaldehyde and cysteine and its Mn (II), Co(II), Ni (II), Cu (II) and Cd (II) complexes on some pathogenic bacteria. The study intends to create Schiff base complexes of Cu(II), Ni(II), Co(II), Cd(II) and Mn(II) ions which are obtained from the reaction of 4-dimethylaminobenzaldehyde, L-valine, L-tyrosine, L-glycine, L-asparagine or 4-aminoantipyrine). Their geometrical structures are verified using different methods. The antibacterial properties of the complexes have been tested against *Bacillus subtilis*, *Bacillus cereus* and *Pseudomonas aeruginosa*.

II. Experimental

2.1 Chemicals and reagents

All chemicals and reagents utilised in this study were laboratory pure purchased from Aldrich or BDH which include 4-dimethylaminobenzaldehyde, L-valine, L-tyrosine, L-glycine, L-asparagine, 4-aminoantipyrine, NaOH, C₂H₅OH, CH₃OH, [Cd(C₂H₃O₂)₂·2H₂O], [Co (C₂H₃O₂)₂·4H₂O], NiCl₂·6H₂O, Cu (CH₃COO)₂·H₂O, (MnCl₂·4H₂O), CHCl₃ and dimethyl sulfoxide (DMSO).

2.2 Synthesis of Schiff bases

The amino acid Schiff bases were prepared according to the following procedure:- NaOH (10 mmol, 0.4g) is dissolved in methanol (30 cm³) and the amino acids [L-valine, L-tyrosine, L-glycine, L-asparagine or 4-aminoantipyrine] (10 mmol) were added to it. The mixtures were stirred with a magnetic there at room temperature. When the mixtures turned homogeneous, a solution of 4-dimethylaminobenzaldehyde (10 mmol, 1.49 g) in ethanol (20 ml) was added. After 2 minutes the solutions was evaporated to 20% of its original volume and 1ml of acetic acid was added immediately. After 2 hours yellow crystals formed. The crystals were then filtered and washed with ethanol and recrystallized from hot methanol to give yellow crystals with the yield of ca. 75-85 % [4]. The proposed structures for the synthesized Schiff bases are shown in Scheme 1:

Scheme 1: Ligand structures

2.3 Synthesis of Schiff base complexes

The Schiff base complexes were produced by mixing 50 ml of methanol containing NaOH (20 mmol, 0.8 g) of the amino acids; [L-valine, L-tyrosine, L-glycine, L-asparagine or 4-aminoantipyrine] (20 mmol) with 50 ml of the methanol solvent in a flask and stirred at room temperature. A solution of 4-dimethylaminobenzaldehyde (20 mmol, 2.98 g) in (50 ml) of ethanol was then added into the solution. After 2 minutes, the metal ions (10 mmol) and the mixtures were stirred magnetically for 3 hours. The solution was evaporated to reduce 75% of the original volume and the residue was left to stand overnight. The products obtained were filtered and recrystallized from a methanol/ethanol (50%) mixture[4].

2.4 Determining antibacterial activities using the agar well diffusion method

Three pathogenic bacteria are exposed to Schiff bases complexes by utilising well diffusion technique [5]. Pathogenic bacteria namely *Bacillus subtilis*, *Bacillus cereus* and *Pseudomonas aeruginosa* are cultured on nutrient agar (NA) plates and incubated at 37 °C for 24 hours to test for purity colonies. Next, they were inoculated into nutrient broth (NB) for overnight culture. One sterile swab was dipped into the suspension of bacteria 104 and rolled separately on the surface of plates which contains nutrient agar (Oxoid). The plates were then dried in a laminar hood at room temperature for duration of 15 minutes. Then, 6 mm holes are made with a cork borer and 20 µL of nutrient agar were pipetted to cover the base of the hole to prevent leakage Schiff bases and Schiff bases complexes. 150 µL Schiff bases and Schiff bases complexes were added to each hole and the plates were incubated at 37 °C for 24 hours aerobically. Finally, after 24 hours the diameter of the growth inhibition area around each hole was measured using a ruler and recorded.

III. Results And Discussion

3.1 Microanalysis

Table 1 shows the elemental analysis data of Schiff based complexes with the formation of 1:1 [M:L] ratio. The findings are in agreement with the theoretical values. Thin-layer chromatography (TLC) method and elemental analyses. (C, H and N) have been utilised to analyse the purity of the Schiff based complexes.

Table (1): Elemental analysis data of Schiff bases and their complexes

Schiff bases complexes	C%	H%	N%	Mwt	Yield%
L1 (C ₁₄ H ₂₀ N ₂ O ₂)	68.01 68.00	7.69 7.65	11.33 11.31	248	75%
[CuL ₁ (Cl)(H ₂ O) ₃]3H ₂ O	29.60 28.98	6.83 5.83	7.97 7.51	454	65%
[CdL ₁ (OH)(H ₂ O) ₃]	39.07 39.00	6.05 5.98	6.51 6.48	430	68%
L2 (C ₁₈ H ₂₀ N ₂ O ₃)	69.27 69.26	6.41 6.40	8.97 8.91	312	78%
[MnL ₂ (OH)(H ₂ O) ₃]	47.26 47.12	5.69 5.70	10.50 10.47	457	56%
[NiL ₂ (OH)(H ₂ O) ₃]	46.87 46.80	5.64 5.65	10.42 10.39	460.86	70%
L3 (C ₁₂ H ₁₅ N ₃ O ₃)	57.83 57.75	6.02 5.98	16.86 16.45	249	85%
[MnL ₃ (OH)(H ₂ O) ₃]3H ₂ O	33.64 33.60	6.31 6.29	9.81 9.80	428	75%
[CdL ₃ (Cl)(H ₂ O) ₃]3H ₂ O	28.63 28.56	5.16 5.10	8.35 8.33	503	70%
L4 (C ₁₁ H ₁₄ N ₂ O ₂)	64.07 64.02	6.67 6.75	13.59 13.53	206	80%
[CuL ₄ (Cl)(H ₂ O) ₃]6H ₂ O	28.33 28.30	6.65 6.63	5.21 5.20	466	66%
[CdL ₄ (Cl)(H ₂ O) ₃]3H ₂ O	31.13 31.00	4.95 4.96	6.60 6.50	424	60%
L5 (C ₂₀ H ₂₂ N ₄ O)	71.83 71.83	6.63 6.63	16.75 16.75	334	79%
[CoL ₅ (Cl) ₂ (H ₂ O) ₂]	48.09 48.00	5.21 5.20	11.22 11.20	499	70%
[CuL ₅ (Cl) ₂] H ₂ O	49.33 50.13	4.93 4.12	11.51 11.63	486	65%

Bold =Experimental values.

3.2 Infrared spectra

The infrared bands analysis of L-valine, L-tyrosine, L-asparagine and L-glycine Schiff bases show a band at 1506- 1630 cm⁻¹ rang assignable to νC=N of the azomethine[6]. The infrared spectral data of the same Schiff bases shows a band at 1576 to 1670 cm⁻¹ rang which proves the existence of CO groups of the L-glycine, L-valine, L-tyrosine, and L-asparagine[7]. The Schiff bases display a broad band at 3144-3397 cm⁻¹ rang which is due to the presence of ν(OH)[8]. The infrared spectra of the present complexes are compared with the free Schiff bases to establish the changes that have taken place during the complexation. Infrared spectral values of complexes portrays bands in the range of 1515 to 1602 cm⁻¹ assignable to ν(C=N) of the azomethine[9]. A slight change is noticed at ν(C=N) band after complexation. The infrared spectral data of all metal complexes display a band at 1609 to 1631 cm⁻¹, which indicate the existence of C=O group of the carboxyl group. This band is present in the higher and lower frequencies compared to the original position in free ligand that is 1576 to 1670 cm⁻¹. The complexation process uses the oxygen of carboxylic groups of the, L-valine, L-tyrosine, L-asparagine and L-glycine and nitrogen atoms of azomethine groups, the ligands are bidentate. A broad band which ranging from 3091 to 3466 cm⁻¹ verifies the existence of water molecules in the separated complexes. δOH disappears and new bands will appear in the spectra of complexes molecules around the range of 469 to 688 cm⁻¹ and 528 to 660 cm⁻¹ are assign to ν(M-O) and ν(M-N) bonding [10]. For the L₅ the infrared spectra of these complexes display a broadband at 3292-3466 cm⁻¹ because of the presence of water molecules [11]. The absorption bands at 1610, 1529 cm⁻¹ are dedicate to the existence of ν(C=N) group of azomethine. The change to a lower frequency show how this group is effected by complexation [12]. Another, complexation site that takes part in coordination is the (C=O) group. The proved can be observed at 1613 cm⁻¹ in the spectrum of free ligand which may be attributed to (C=O) of a carbonyl group. The shifting from a higher frequency and *vice versa* in the spectra of the complexes suggests the participation of (C=O) group in coordination with the metal ion through the oxygen atom of a carbonyl group of 4-aminoantipyrine [13]. New bands which are not observed in the free Schiff bases are observed at 464-632 and 628-522 cm⁻¹ due to ν(M-O) and ν(M-N) vibrations. This provides a good proof for the sharing of the oxygen and nitrogen atoms during complexation [14].

3.3 Electronic spectra

The electronic spectrum of Cu (II)-L¹ complex show bands in the wavelength range of 265-340 nm (8833-11333 cm⁻¹) because of ²E_g → ²T_{2g} transition and an octahedral geometry is suggested. At the same time, the spectral data of Cu (II)-L⁴ Schiff based complex show two bands at 295 and 650 nm (9833 and 21667cm⁻¹) which are accredited to charge transfer transitions. The intensity of these two bands supports the presence of an octahedral structure. For Cu (II)-L⁵ complex, the spectrum displays bands in the range of 300 to 680 nm (10000-22667 cm⁻¹) because of charge transfer and ²T_{2g} → ²E_g transitions which propose a square planar geometry[15]. The electronic spectrum of Co (II)-L⁵ Schiff based complex of [Co L₅ (H₂O)₂(Cl)₂] illustrate two bands at 320 and 515nm (10667 and 17167cm⁻¹). The first band happens because of ⁴T_{1g} (F) → ⁴T_{2g} (F) transition and the second one occurs due to ⁴T_{1g} (F) → ⁴T_{1g} (P) transition. These data illustrate an octahedral geometry. The cadmium complexes display only the charge transfer transitions that can be associated with the charge transfer from ligand to metal and *vice versa*, no d-d transition are expected for d10 Cd(II) complexes [16]. In the electronic spectra of both complexes which are recorded in (DMSO) solution, an absorption band in the range of 270 to 420 nm is noticed and this may be linked with a π → π* transition originating mostly in the azomethine chromophore (imine π → π* transition). Cd (II) complexes are colourless and diamagnetic as expected for d10 system. For Ni (II)-L² Schiff based complex, the spectrum displays bands in the range of 299 to 560 nm (9967-18667 cm⁻¹) which are linked to ³A_{2g} (F) → ³T_{2g} (F) and ³A_{2g} (F) → ³T_{1g} (P) transitions. These data prove the presence of an octahedral geometry. The electronic spectra of the Mn(II)-L₂ complex illustrates absorption bands at 298nm (17819cm⁻¹), 575nm (19167cm⁻¹) which is a feature of octahedral geometry that correspond to ⁶A_{1g} → ⁴T_{2g} (G), ⁶A_{1g} → ⁴E_{1g}, ⁴A_{1g} (G) transitions. The electronic spectra of the Mn(II)-L₃ complex shows absorption bands at 295nm, 380nm (9833-12667 cm⁻¹) which is a feature of octahedral geometry that correspond to ⁶A_{1g} → ⁴T_{2g} (G), ⁶A_{1g} → ⁴E_{1g}, ⁴A_{1g} (G) transitions[17].

Table 2: Infrared assignments (cm⁻¹) and electronic absorption (nm, cm⁻¹) of Schiff bases complexes.

Ligand/Complexes	v(OH) (H ₂ O)	v(COO ⁻)	v(C=N)	v(M-O)	v(M-N)	v(NH ₂)	λ _{max} nm	cm ⁻¹	Expected geometry
L1	3395	1670	1583	-	-	-	253	8433	-
[CuL ₁ (Cl)(H ₂ O) ₃]3H ₂ O	3387	1631	1553	653	550	-	300	10000	octahedral
							340	11333	
[CdL ₁ (OH)(H ₂ O) ₃]	3289	1620	1594	469	629	-	280	9333	octahedral
							305	10167	
L2	3144	1605	1582	-	-	-	249	8300	-
[MnL ₂ (OH)(H ₂ O) ₃]	3397	1609	1513	648	528	-	325	10833	octahedral
							575	19167	
[NiL ₂ (OH)(H ₂ O) ₃]	3394	1603	1515	645	521	-	320	10667	octahedral
							560	18667	
L3	3197	1670	1630	-	-	3097	285	9500	-
[CdL ₃ (Cl)(H ₂ O) ₃]3H ₂ O	3391	1658	1595	608	553	-	299	9967	octahedral
						2914	385	12833	
[MnL ₃ (OH)(H ₂ O) ₃]3H ₂ O	3363	1641	1602	631	596	2914, 2806	299	9967	octahedral
							380	12667	
L4	3397	1576	1506	-	-	-	343	11433	-
[CuL ₄ (Cl)(H ₂ O) ₃]6H ₂ O	3369	1610	1515	688	545	-	300	10000	octahedral
							650	21667	
[CdL ₄ (Cl)(H ₂ O) ₃]3H ₂ O	3581	1627	1550	681	660	-	305	10166	octahedral
							370	12333	
L5	3172	1613	1586	-	-	-	343	29155	-
[Cu L ₅ (Cl) ₂] H ₂ O	3466	1615	1610	464	628	-	362	27624	
							280	33670	square planar
							517	19342	
[Co L ₅ (H ₂ O) ₂ (Cl) ₂]	3292	1647	1529	632	522	-	320	10667	octahedral
							515	17167	

3.4 ¹HNMR spectra

The ¹HNMR spectra data of L₁ which is recorded in DMSO-d⁶ shows signals at 0.833 , 3.044 and 2.083 ppm that is corresponding to (CH₃)₂, (CH₃)₂N and CH groups. The multiplet signals at 6.782 to 7.691 ppm are caused by aromatic protons. The -OH of carboxyl group emerged at 10.00 ppm. The data obtained proof the connection between these two groups. Nevertheless, the --NH₂ group vanishes when L₁ is formed, signifying the presence of azomethine group at 8.52 ppm. The ¹HNMR spectra data of L₂ which is recorded in DMSO-d⁶ shows signals at 2.83 , 2.96 and 4.63 ppm that corresponds with -CH₂, (CH₃)₂N and -CH groups. The signal at 5.91 ppm corresponds to OH Aromatic. The multiplet signals at 6.77 -7.68 ppm are caused by aromatic protons. The signal at 9.90 ppm corresponds with azomethine proton. ¹HNMR spectra data of the L₃ which is recorded in DMSO-d⁶ shows signals at 7.18, 3.075 and 3.36 ppm corresponding to NH₂, (CH₃)₂N and CH groups. The multiplet signal at 6.69 to 7.72 ppm are due to aromatic protons. The -OH of carboxyl group appears at 9.664 ppm. These data support the involvement of these two group. Nevertheless, the --NH₂ group will vanish when

L₃ is formed, indicating the presence of azomethine group at 8.03 ppm. The ¹H-NMR spectrum of L₄ is recorded in DMSO-d₆ at room temperature and it shows C₆H₅ multiplet at 6.75 to 7.68 ppm, -CH₂ at 3.89 ppm, N-(CH₃)₂ at 3.009 ppm and -OH proton appears at 9.996 ppm[18]. Similarly, the --NH₂ group vanishes with the formation of L₄ which indicates the presence of azomethine group at 9.996 ppm. ¹HNMR spectra data of L₅ which is recorded in DMSO shows signals at 3.101, 2.97 and 4.49 ppm corresponding to N-CH₃, (CH₃)₂N and CH₃ groups[19]. At the same time, the --NH₂ group vanishes with the formation of L₅ which indicates the presence of azomethine group at 9.416 ppm. There is an absence of -OH proton due to complexation. A considerable change is observed in all other signals in this complexes[20].

3.5 X-ray crystallographic study of the compound L⁵

The X-ray investigation of the compound showed that the compound L₅ crystallized in monoclinic system with space group of C2/c and the unit cell dimensions are a= 17.7916, b= 6.8610, c= 29.7199, α=90.000 β= 101.326 γ= 90.000., Z = 8 and V = 3557.20 Å³. Figure 1 shows the molecular structure with the numbering scheme of the compound, the molecule is discrete and the asymmetric unit consist two molecules. The phenyl rings in each molecule (C1—C6) and (C21—C26) make a dihedral angle of 56.4(7) and 56.4(8)° with the pyrazol heterocyclic ring, respectively. The bond lengths and angles are in normal ranges and comparable to those in N,N'-bis(2,3,4-trimethoxybenzaldimine)-1,4-diaminocyclohexane[21] (Table 3) .

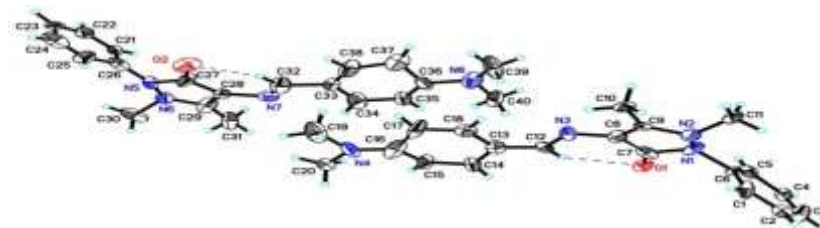


Fig.1. ORTEP diagram of the L₅ drawn at 50% probability displacement ellipsoids. The dashed line indicates the intramolecular hydrogen bond.

Table 3. Selected Bond Lengths (Å) and Bond Angles (°) for Compound L⁵

Compound L ⁵			
Bond	Dist.	Angle	(°)
O1-C7	1.237(17)	N2-N1-C7	112.1(9)
O2-C27	1.256(18)	N2-N1-C6	117.2(10)
N1-N2	1.415(17)	C7-N1-C6	121.9(12)
N1-C7	1.438(16)	C9-N2-N1	101.5(10)
N1-C6	1.488(11)	C9-N2-C11	127.5(10)
N2-C9	1.349(14)	N1-N2-C11	114.6(9)
N2-C11	1.431(18)	C12-N3-C8	123.4(12)
N3-C12	1.279(17)	C16-N4-C20	124.4(17)
N3-C8	1.389(13)	C16-N4-C19	121.4(17)
N4-C16	1.317(19)	C20-N4-C19	113.8(12)
N7-C32	1.19(2)	C27-N5-N6	107.3(10)

Both molecules showed the presence of C12—H12...O1 and C32—H32...O2 intramolecular hydrogen bond, as a result a pseudo-six membered O1...H12/C12/N3/C8/C7 and O2...H32 /C32/N7/C28/C27 rings are formed (Table 4). In the crystal structure, the molecules are linked by C10—H10A...O1 and C32—H32...O2 intermolecular hydrogen bonds to form dimers. (Fig.2).

Table 4. Hydrogen Bond Lengths (Å) and Bond Angles (°) for Compound L⁵

Compound	D—H...A	D—H	H...A	D...A	D—H...A	Symmetry code
L ₅	C12—H12...O1	0.93	2.38	3.059(15)	130	
	C32—H32...O2	0.93	2.25	2.97(2)	133	
	C10—H10A...O1	0.96	2.48	3.360(15)	153	x,-1+y,z
	C20—H20C...O1	0.96	2.56	3.491(19)	164	-1/2+x,1/2+y,z

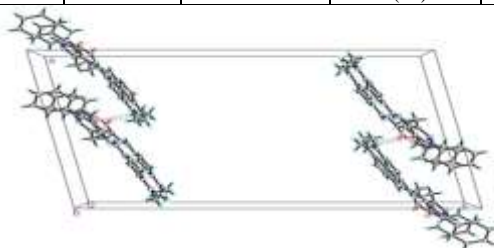


Fig. 2. Molecular packing of L₅ viewed down the b axis. Dashed lines denote C—H...O intermolecular hydrogen bonds.

3.6 Antibacterial Activity

The findings of the antibacterial activities are displayed in Tables 5-9. Schiff based complexes displayed good activities against three kinds of bacteria namely *Bacillus subtilis*, *Bacillus cereus* and *Pseudomonas aeruginosa*.

3.6.1 Antibacterial Activity of L^1 , Cd (II) and Cu (II)- L^1 complexes

The findings of antibacterial activities are displayed in Table 5. Schiff based complexes display good activity against three kinds of bacteria namely *Bacillus subtilis*, *Bacillus cereus* and *Pseudomonas aeruginosa*. Complexes which are obtained from L^1 show the highest inhibitory influence against the bacteria being tested. Basically, the inhibition area of the complexes increases along with their concentrations. The highest inhibition area is obtained by L^1 .Cu against *Bacillus subtilis* whereas the minimum inhibition are is obtained by L^1 against *Bacillus cereus* and *Pseudomonas aeruginosa*. Hence complexation increases the antimicrobial activity. Such increased activity of the metal complexes can also be explained on the basis of chelation theory [4]. According to this, the chelation reduces the polarity of the metal atom mainly because of the partial sharing of its positive charge with donor group and possible π -electron delocalisation over the whole ring. This increases the lipophilic character of the metal chelate system which favours its permeation through lipid layer of the cell membranes.

Table 5. Diameter of inhibition zone for antibacterial screening of L^1 , Cd(II) and Cu(II)- L^1 complexes. Inhibition zone diameter in mm

Ligand/Complexes	Concentration	<i>Paeruginosa</i>	<i>Bacillus subtilis</i>	<i>Bacillus cereus</i>
L^1	5mg/ml	0	4	0
	10mg/ml	0	6	0
	20mg/ml	0	8	0
[CdL ₁ (OH)(H ₂ O) ₃]	5mg/ml	4	16	4
	10mg/ml	10	19	10
	20mg/ml	19	24	18
[CuL ₁ (Cl)(H ₂ O) ₃]3H ₂ O	5mg/ml	8	16	6
	10mg/ml	14	19	9
	20mg/ml	22	25	16

Notes: SD inhibition zone = \pm 1 mm (biological replicates, 3).

3.6.2 Antibacterial of L^2 , Ni(II) and Mn(II)- L^2 complexes

In order to test the antibacterial activity of L^2 , Ni (II) and Mn (II)- L^2 complexes, more than one test organism is utilised to increase the probability of detecting the antibiotic potential of the tested materials. The sensitivity of a microorganism to antibiotics and other antimicrobial agents are determined by the assay plates which are incubated at 37 °C for a day (for bacteria). The compounds which are tested display biological activities against different kinds of Gram-positive and Gram-negative bacteria. The findings are displayed in Table 6. When the biological activity of the Schiff bases and their metal complexes are compared the findings show that the biological activity of Ni (II) and Mn (II) complexes is higher than Ligand. The biological activity of the complexes are Ni (II) > Mn (II).

Table 6. Diameter of inhibition zone for antibacterial screening of L^2 , Ni(II) and Mn(II)- L^2 complexes. Inhibition zone diameter in mm

Ligand/Complexes	Concentration	<i>Paeruginosa</i>	<i>Bacillus subtilis</i>	<i>Bacillus cereus</i>
L^2	5mg/ml	10	11	9
	10mg/ml	13	15	12
	20mg/ml	15	16	14
[NiL ₂ (OH)(H ₂ O) ₃]	5mg/ml	10	13	12
	10mg/ml	14	16	15
	20mg/ml	17	25	18
[MnL ₂ (OH)(H ₂ O) ₃]	5mg/ml	20	10	9
	10mg/ml	20	12	11
	20mg/ml	22	13	12

Notes: SD inhibition zone = \pm 1 mm (biological replicates, 3).

3.6.3 Antibacterial Activity of L^3 , Cd (II) and Mn (II)- L^3 complexes

The antibacterial features of L^3 , Cd (II) and Mn (II)- L^3 on test bacteria (two Gram-positive and one Gram-negative) is displayed in Table 7. The findings show that the compounds display antibacterial features against two Gram-positive and one Gram-negative namely *Bacillus cereus*, *Bacillus subtilis* and *Paeruginosa*. Nevertheless, the antibacterial features of Cd(II) and Mn(II) complexes is higher than Ligand. Hence complexation increases the antimicrobial activity. Such increased activity of the metal complexes can also be explained on the basis of chelation theory [10]. According to this, the chelation reduces the polarity of the metal atom mainly because of the partial sharing of its positive charge with donor group and possible π -electron delocalisation over the whole ring. This increases the lipophilic character of the metal chelate system which favours its permeation through lipid layer of the cell membranes.

Table 7. Diameter of inhibition zone for antibacterial screening of L^3 , Cd(II) and Mn(II)- L^3 complexes. Inhibition zone diameter in mm

Ligand/Complexes	Concentration	<i>Paeruginosa</i>	<i>Bacillus subtilis</i>	<i>Bacillus cereus</i>
L^3	5mg/ml	4	4	4
	10mg/ml	7	8	6
	20mg/ml	9	18	13
[CdL ₃ (Cl)(H ₂ O) ₃]3H ₂ O	5mg/ml	4	17	14
	10mg/ml	8	19	17
	20mg/ml	19	29	24
[MnL ₃ (OH)(H ₂ O) ₃]3H ₂ O	5mg/ml	4	4	4
	10mg/ml	6	5	7
	20mg/ml	10	8	13

Notes: SD inhibition zone = ± 1 mm (biological replicates, 3).

3.6.4 Antibacterial Activity of Cd (II), L^4 and Cu (II)- L^4 complexes

The antibacterial features of Cd (II), L^4 and Cu (II) - L^4 on three bacteria (two Gram-positive and one Gram-negative) is displayed in Table 8. The findings show that compounds portray antibacterial features against two Gram-positive and one Gram-negative which are notably *Bacillus cereus*, *Bacillus subtilis*, and *Paeruginosa*. Nevertheless, the antibacterial features of Cd (II) and Cu (II) complexes are higher than ligand.

Table 8. Diameter of inhibition zone for antibacterial screening of L^4 , Cu (II) and Cd (II)- L^4 complexes. Inhibition zone diameter in mm

Ligand/Complexes	Concentration	<i>Paeruginosa</i>	<i>Bacillus subtilis</i>	<i>Bacillus cereus</i>
L^4	5mg/ml	4	4	4
	10mg/ml	5	8	12
	20mg/ml	9	12	14
[CuL ₄ (Cl)(H ₂ O) ₃]6H ₂ O	5mg/ml	13	19	6
	10mg/ml	18	24	12
	20mg/ml	24	29	14
[CdL ₄ (Cl)(H ₂ O) ₃]3H ₂ O	5mg/ml	14	16	6
	10mg/ml	19	28	10
	20mg/ml	29	31	20

Notes: SD inhibition zone = ± 1 mm (biological replicates, 3).

3.6.5 Antibacterial Activity of L^5 , Co (II) and Cu (II)- L^5 complexes

Table 9 displays the findings obtained for antibacterial activities. The Schiff base complexes which are being examined show good response towards three kinds of bacteria namely *Bacillus cereus*, *Pseudomonas aeruginosa* and *Bacillus subtilis*. Complexes which are obtained from L^5 show the highest inhibitory influence towards the bacteria which are being investigated. Basically, the inhibition area of the complexes increases as their concentration increase. The highest inhibited area is obtained by L^5 .Co against *Bacillus subtilis* whereas the least inhibited area is obtained by L^5 against *Bacillus cereus*. Therefore, complexation enhances the anti-microbial activity. This increased activity of metal complexes may also be explained on the basis of the chelation theory [22]. The study chelation reduces the polarity of the metal atom mainly due to the sharing of its partial positive charge with the group of donors and can electronic delocalization throughout the cycle. This increases the lipophilicity of the metal chelate system that facilitates its permeation through the lipid bilayer of cell membranes.

Table 9. Diameter of inhibition zone for antibacterial screening of L^5 , Cu(II) and Co(II)- L^5 complexes. Inhibition zone diameter in mm

Ligand/Complexes	Concentration	<i>Paeruginosa</i>	<i>Bacillus subtilis</i>	<i>Bacillus cereus</i>
L^5	5mg/ml	10	11	8
	10mg/ml	13	13	10
	20mg/ml	15	15	12
[Cu L ₅ (Cl) ₂] H ₂ O	5mg/ml	10	8	8
	10mg/ml	24	23	23
	20mg/ml	29	27	26
[Co L ₅ (H ₂ O) ₂ (Cl) ₂]	5mg/ml	20	16	13
	10mg/ml	24	20	18
	20mg/ml	25	22	20

Notes: SD inhibition zone = ± 1 mm (biological replicates, 3).

IV. Conclusion

Schiff base amino acid and 4-Aminoantipyrine complexes were prepared and characterized by based on the elementary analysis. X-ray, infrared, electronic absorption and ¹H NMR. An in vitro investigation carried out on the synthesized complexes showed that all the complexes displayed very good results against antibacterial activity of the selected microorganisms which are being examined.

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