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Synthesis, Characterization, Single Crystal Structural Studies, Antibacterial Activity and DFT Investigations of 2-Chloro-5-Ethoxy-3,6-Bis(Methylamino)-1,4-Benzoquinone

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Abstract

A new compound 2-chloro-5-ethoxy-3,6-bis(methylamino)-1,4-benzoquinone (cemb) has been synthesized and characterized by various physicochemical techniques. The compound crystallizes in the triclinic system with the space group P-1 and contains extended hydrogen bonding interaction providing supramolecular architecture. The compound contains CH···π and oxygen···π interaction which stabilizes the crystal structure and is also responsible for the semiconducting behavior. The DFT calculations in the gaseous phase of cemb with B3LYP hybrid functional show good correlation with experimental results (single crystal XRD data) and are used to predict the molecular properties. Excluding the side ethyl group, the structure possesses C_s symmetry. The ring breathing mode of vibration is found to have frequency (551 cm⁻¹) which is much lower as compared to the corresponding mode of benzene (991 cm⁻¹). The bioefficacy of the compound has been examined against the growth of bacteria *in vitro* to evaluate antimicrobial potential. The photoluminescence properties indicate that the compound has characteristic fluorescence emissions. The HOMO and LUMO energy gap reveals that the energy gap reflects the chemical activity of the molecule.

Keywords: 1,4-Benzoquinone derivative; Crystal structure; Solid state electrical conductivity; Antibacterial activity; DFT calculations

Introduction

Naturally occurring compounds containing a quinone group are involved in various bioenergetic processes as electron-transport agents. Some quinones play important roles in the biochemistry of energy production and serve as vital links during electron transport. These compounds have attracted considerable attention because of their biological activity and chemotherapeutic values [1]. Various quinone containing oxygenated aromatic rings show biological activities [2]. Quinones are found in all respiring animal and plant cells where they function primarily as components of the electron transport chains involved in cellular respiration [3] and photosynthesis [4]. These compounds occur in other natural contexts and as pollutants [4-8]. Quinones form an important class of toxic metabolites and paradoxically can be either mutagenic and, therefore, are potentially carcinogenic, or effective anticancer agents [9]. Some quinones possibly play roles in the cellular defence since they effectively inhibit growth of bacteria, fungi and/or parasites [10-13].

Quinones are a large class of compounds endowed with rich and fascinating chemistry [14]. They are widely distributed in the natural world [15], being found in bacteria, plants and arthropods and hence, quinones are ubiquitous to living systems. Their role as electron transfer agents in primary metabolic processes like photosynthesis and respiration is vital to human life. A large number of chemical derivatives with 1,4-benzoquinone as the basic subunit exhibit prominent pharmacological applications such as antibiotic, [16,17] antitumor, [18-21] antimalarial, [22] antineoplastic, [23] anticoagulant [24] and herbicidal activities [25]. Coordination chemistry of quinones is also quite rich from the perspective of designing magnetic materials [26] and understanding photo physical properties [27]. The chemistry of quinones and hydroquinones is of interest because of their involvement in the electron and hydrogen transfer reactions in various processes including the biological energy conversion reactions such as respiration and photosynthesis [28-30]. For the antibacterial agents often used, quinone compounds are tetracyclines, new quinolones, anthracyclines, and mitomycins. More than 2000 naturally occurring quinones, likeanthraquinones, naphthoquinones, and benzoquinones are now known and are widely distributed in nature as pigments and intermediates in the cellular respiration and photosynthesis [31,32].

In view of the above, it is of great interest to investigate the derivatives of benzoquinone and to study the mode of bonding and structure. The present paper reports the synthesis, spectral, structural and DFT studies of 2-chloro-5-ethoxy-3,6-bis(methylamino)-1,4-benzoquinone (cemb).

Experimental Details

Chemicals and starting materials

Commercial reagents were used without further purification and all experiments were carried out in open atmosphere. The solvents were dried and distilled before use, following the standard procedures.

Physical measurements

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Carbon, hydrogen, nitrogen contents were estimated on a CHN Model CE-440 Analyser and on an Elementar Vario EL III Carlo Erba 1108. IR spectra were recorded in the 4000-400 cm⁻¹ region in KBr pellets on a Perkin Elmer/Varian Excalibur 3100 FT-IR spectrophotometer. ¹H and ¹³C NMR spectra were recorded in DMSO-d₆ on a JEOL AL300 FT-NMR spectrophotometer. The electrochemical experiments were performed with a CHI-660C (CH Instruments, USA), using threeelectrode system, platinum wire as counter electrode (area ~ 8.0 cm²) and Hg/HgO was used as the reference electrode. Electrical conductance of pressed pellets was measured by a conventional two-probe method in the 313-443 K range with contacts made on the pellet surfaces by means of silver paint. A Keithley 236 SMU was used to measure the sample resistance. The fluorescent data were collected at room temperature with a Varian CARYECLIPSE spectrophotometer. Six bacterial pathogens Salmonella typhi, Shigella flexneri, Staphylococcus aureus, Aeromonas hydrophila, E. coli and Enterococcus faecalis were used to test the antibacterial activity of the compound. Commercial antibacterial drugs, streptomycin sulphate and chloramphenicol (Himedia) in same concentration of 10-100 µg/disc are used to compare effectiveness of the test compounds.

Synthesis of cemb

For the synthesis of cemb, sodium acetate (1.148 g, 14 mmol) was added pinch by pinch within 20 min to a reaction mixture of chloranil (0.492 g, 2 mmol), which was added drop wise to methylamine hydrochloride (0.270 g, 4 mmol) in 15 ml of absolute ethanol with constant stirring for 1 h and refluxed 4h at 65°C. The solid separated was filtered off, washed out successively with ethanol-water mixture (50:50 v/v) and air dried. The precipitates were suspended in ethanol (20 mL) to which pyridine (1 mL) was added and refluxed for 1 h at 65°C. The resulting clear solution was filtered and kept for crystallization. Single crystals of cemb suitable for X-ray analysis were obtained by slow evaporation of the solutions over a period of 25 days (Scheme 1). Grey color crystals were obtained with: Yield: 72%; m.p>290°C; Anal. Found: C, 49.72; H, 5.46; N, 11.05%. Calc. for $C_{10}H_{13}ClN_2O_3(244.67)$: C, 49.09; H, 5.36; N, 11.45%. The crystal structure was confirmed by XRD.

Antibacterial assay

The antibacterial assay was done according to the Bayer method [33]. The test compounds were dissolved in DMSO to a final concentration of 5 mg/mL. Sterilized Whatman no. 1 filter paper discs (5 mm) were impregnated with different volume (2, 4, 8, 12, 16 and 20 μ L) of compounds to get a final concentration of 10, 20, 40, 60, 80 and 100 μ g per disc. Sterilized paper disc loaded with the 10 μ L of DMSO was taken as a control. The bacterial test pathogens were spread on fresh Müller Hinton Agar (MHA) plates with the help of cotton swabs to form an even lawn of the test bacteria. The filter paper disc impregnated with the test compounds were placed on the surface of the MHA plates seeded with test bacteria and the plates were incubated in a BOD Incubator (Caltan-152, Narang Scientific Works, New Delhi, India) for 24 h at 37 ± 2°C. The inhibition zones around each disc were



Page 2 of 10

measured after 24 h of incubation.

Crystal structure determination

Crystals suitable for X-ray analysis of cemb were grown at the room temperature. The crystal structure data were collected on an Oxford Gemini diffractometer equipped with a CrysAlis CCD software using a graphite mono-chromated Mo K α (λ =0.71073 Å) radiation source at 293 K for cemb. Multiscan absorption correction was applied to the X-ray data collection and the structure was solved by direct methods (SHELXS-08) and refined against all data by full matrix leastsquare on F² using anisotropic displacement parameters for all nonhydrogen atoms. All hydrogen atoms were included in the refinement at geometrically ideal position and refined with a riding model [34]. The MERCURY package and ORTEP-3 for Windows program were used for generating structures [35,36]. The details of data collection, structure solution and refinement are listed in Table 1. Selected bond lengths and angles are given in Table 2. ORTEP diagram of cemb with atom numbering schemes is shown in Figure 1A. The single crystal X-ray diffraction studies indicate that the starting compound chloranil is converted to cemb during the course of reaction. The grown single crystals of cemb were analyzed by single crystal X-ray data. The bond lengths and angles in the ring are normal [37,38]. The structure is stabilized by intermolecular N-H-O and C-H-O hydrogen bonding occurring between hydrogen of substituted N-methyl amine group to oxygen of nearby molecule and $CH \cdot \pi$ interaction occurring between hydrogen of substituted ethoxy group to centroid of ring of a nearby molecule at a distance of 3.561 Å and Oxygen... π at a distance of 3.557 Å producing linear chain structure. The structure is also stabilized by

Empirical formula	C ₁₀ H ₁₃ CIN ₂ O ₃		
Formula weight	244.67		
Т (К)	293 (2)		
Λ (Μο Κα) (Å)	0.71073		
Crystal system	Triclinic		
Space group	P-1		
a (Å)	7.5866 (15)		
b (Å)	8.6898 (18)		
c (Å)	9.740 (2)		
α (°)	91.172 (18)		
β (°)	112.09 (2)		
γ (°)	111.71 (2)		
V (Å ³)	543.2 (2)		
Z	2		
D _{calc} (mg m ⁻³)	1.496		
μ (mm ⁻¹)	0.345		
F (000)	256		
Crystal size (mm ³)	0.24 x 0.21 x 0.18		
θ range (°)	3.11-29.00		
Index ranges	-10 ≤ h ≤ 9 -11 ≤ k ≤ 11		
Paflections collected	3570		
Independent reflections	1205		
Data/restraints/ parameters	2894/0/145		
$Goodness_of_{fit}$ on F^2	0.973		
Final R indices wR [I>2 σ (I)] (R)	0.0760.0.1516		
Final P indices $(all data)$	0.1561 0.1050		
Largest diff pook/bolo ($a^{\frac{1}{3}}$)	0.257 0.415		
	0.337, -0.415		

 ${}^{a}R_{1} = \Sigma ||F_{o}| - |Fc||\Sigma |F_{o}|, {}^{b}R_{2} = [\Sigma w (|F_{o}| - |F^{2}_{o}|)^{2}/\Sigma w |F^{2}_{o}|^{2}]^{1/2}.$ **Table 1:** Crystallographic data for cemb molecule.

Bond lengths (Å)			Bond angles (°)		
	Exp.	Cal.		Exp.	Cal.
CI-C11	1.737 (4)	1.74	C9-O3-C4	117.3 (4)	118.2
O3-C9	1.325 (6)	1.32	C3-N2-C8	129.3 (4)	129.5
O3-C4	1.414 (5)	1.42	N2-C3-C4	128.4 (4)	128.8
O1-C5	1.246 (4)	1.25	N2-C3-C2	112.9 (3)	112.3
N2-C3	1.326 (5)	1.34	O2-C2-C1	123.5 (4)	123.9
N2-C8	1.436 (5)	1.44	O2-C2-C3	116.5 (4)	116.8
O2-C2	1.249 (4)	1.26	N1-C6-C5	111.5 (3)	112.2
N1-C6	1.326 (5)	1.34	N1-C6-C1	129.2 (4)	129.4
N1-C7	1.445 (5)	1.45	C6-N1-C7	130.9 (4)	130.5
C5-C6	1.523 (6)	1.52	O1-C5-C4	123.2 (4)	123.6
C6-C1	1.376 (5)	1.36	O1-C5-C6	117.1 (3)	117.6

Table 2: Selected bond lengths (Å) and bond angles (°).



Figure 1: (A) ORTEP diagram of cemb molecule at 30% probability level. **(B)** NH···O, CH···O, CH···Cl, CH··· π and Oxygen··· π interactions leading to linear chain structure.

		1	1	
D-H A	d(D-H)	d(H A)	d(D···A)	<(DHA)
N2-H2 O1	0.859	2.155	2.917	147.59
N1-H1O2	0.860	2.162	2.907	144.69
C7-H01DO2	0.961	2.642	3.228	119.69
C9-H01HCI1	0.970	2.992	3.826	155.55

Table 3: Intermolecular hydrogen bond parameters [Å and °].



intramolecular hydrogen bonding, i.e., N-H…O, C-H…O and C-H…Cl (Figure 1b). The geometrical parameters relevant for these interactions are given in Table 3. The above observed intermolecular interactions have stabilized the molecular crystal packing which leads to the

formation of supramolecular architecture [39].

UV-Vis spectra

The experimental UV–vis spectrum of cemb in DMSO is shown in Figure 2. It shows a transition at 336.50 nm.

Photoluminescence emission spectra

The photoluminescence emission spectrum of cemb was recorded in DMSO at room temperature (Figure 3). On excitation with a 335 nm laser source at room temperature, a strong fluorescence emission peaked at 426 nm is observed. The photoluminescence property shows that the compound has characteristic fluorescence emissions. The main chromospheres in the compound are the carbonyl and halogen groups and its conjugation degree is further enhanced by the amine group. The solution state photoluminescence property of compound in DMSO indicates that increase in the dilution of solution resulted in a blue shift and decrease in intensity at various concentrations due to weakening of the CH··· π and CH···Cl interactions, but the luminescence property persists even in dilute solution due to existence of the intermolecular interactions [40].

Computational Details

All calculations were performed using the Gaussian 03 and Gauss View 4.1 software [41]. The structure optimization of cemb was done using DFT method with the functional B3LYP and basis set 6-311 ++ G(d,p) [42]. The input geometry of the molecule for the DFT calculations was generated from the single crystal X-ray investigation carried out in the present study. The HOMO-LUMO energies, MEP, UV-vis spectra, linear polarizability and first hyperpolarizability were also calculated by the DFT method [43,44]. To facilitate the vibrational analysis potential energy distributions (PEDs) were also computed using the GAR2PED software.

Results and Discussion

Chloranil react with methylamine hydrochloride in presence of sodium acetate and pyridine to yield compound cemb. There is substitution of two methylamine and one ethoxy group with replacement of three chlorine atoms through a condensation reaction, which is confirmed by X-ray diffraction studies. The Scheme 1 depicts the formation of cemb. The compound is air stable, non-hygroscopic crystalline solids, which is insoluble in common organic solvents but soluble in DMF, DMSO and can be kept in desiccators over a prolonged period without any sign of decomposition with melting point at >290°C.

Geometry optimization

The optimized geometry (optimized energy -2037.988 a.u) of cemb are shown in Figure 4(A). There is good agreement between the geometrical parameters obtained by X-ray crystallography to those generated by DFT method as shown in Table 2. The slight disagreement in the bond lengths and angles are due to the fact that the experimental results are for the solid phase and the theoretical calculations are for the gas phase [45]. In the solid state, the existence of a crystal field along with the intermolecular interactions connect the molecules together, which results in the differences in geometrical parameters between the calculated and experimental values.

UV-Vis spectra

The DFT calculations were performed for the gas phase and DMSO solvent. Calculations predict an intense electronic transition at 328.24 nm.

Page 3 of 10





Figure 4: (A) Optimized structure (B) MEP plot (C) ESP plot and (D) Contour map of cemb molecule.



Vibrational assignments

The cemb molecule is a 29 atomic molecule with 81 normal modes of vibration. The observed and calculated IR and Raman spectra of cemb are reproduced in Figures 5 and 6. The distribution of the normal modes could be divided under the following sections:- (i)



CH₃ group modes (27 modes) and CH₂ group modes (6 modes), (ii) C=O modes (6 modes), (iii) Ring modes (12 modes), (iv) C-NH(CH₃) modes (6 modes), (v) N-H modes (6 modes), (vi) N-CH₃ modes (6 modes), (vii) C-Cl modes (3 modes), (viii) C-O(CH₂)(CH₃) modes (3 modes), (ix) O-CH₂(CH₃) modes (3 modes) and (x) H₂C-CH₃ modes (3 modes). The calculated and observed vibrational frequencies with the corresponding PEDs and assignments for the same molecule are collected in Table 4. Assignments for several of the modes agree with the modes of 2,5-diamino-3,6-dichloro-1,4-benzoquinone [46]. In the following we discuss some additional modes briefly.

CH₃ group modes (27 modes) and CH₂ group modes (6 modes): The ring CH stretching vibrations fall in the region 3000-3100 cm⁻¹ [47-51]. The symmetric CH₃ stretching modes of the CH₃ groups, the CH₂ symmetric stretching group modes and the antisymmetric CH₂ stretching modes of the ¹⁸CH₃[#] and ¹⁴CH₃ groups are purely localized modes whereas the antisymmetric stretching modes of the ²⁶CH₃ and ²³CH₂ groups couple strongly. The assignments for these modes could be seen in Table 4. The antisymmetric deformation modes (") of the ²⁶CH₃, 18 CH₃ and 14 CH₃ groups are assigned at the frequencies 1442 (ν_{56}) and 1460 ($\nu_{_{59}}$), 1444 ($\nu_{_{57}}$) and 1466 ($\nu_{_{60}}$) and 1448 ($\nu_{_{58}}$) and 1467 ($\nu_{_{61}}$) cm⁻¹. The frequencies 1362 (ν_{52}), 1442 (ν_{54}) and 1427 (ν_{55}) cm⁻¹ correspond to the symmetric deformation modes () of the groups ${
m ^{26}CH}_{
m s}$, ${
m ^{14}CH}_{
m s}$ and ¹⁸CH₃ respectively. Out of the two rocking modes of the ¹⁴CH₃ group the perpendicular rocking is calculated to have magnitude 1107 (v_{42}) cm⁻¹ while the parallel rocking mode has magnitude 1154 (v_{45}) cm⁻¹. Similarly, the ${\rm ^{18}CH_3}$ and ${\rm ^{26}CH_3}$ groups also have two rocking modes at the frequencies 1173 $(\nu_{_{46}})$ and 1111 $(\nu_{_{43}})$ and 810 $(\nu_{_{33}})$ and 1103 $(\nu_{_{41}})$ cm⁻¹. These rocking modes are coupled with each other and many other modes. The torsional modes τ ($^{18}CH_3),$ τ ($^{14}CH_3)$ and τ ($^{26}CH_3)$ are assigned at 107 (ν_6) 171 (ν_{10}) and 232 (ν_{14}) respectively.

Out of the rest four modes of the CH₂ group the scissoring mode is calculated to have magnitude 1476 (ν_{62}) cm⁻¹, the wagging mode is found at 1384 ($_{53}$) cm⁻¹ while the twisting and the rocking modes have the magnitudes 1268 (ν_{49}) and 1141 (ν_{44}) cm⁻¹.

C=O modes (6 modes): The two carbonyl stretching modes couple strongly with the ring stretching modes to give rise to the computed frequencies 1661, 1639, 1595 and 1584 cm⁻¹; the former two frequencies being assigned to the "opc v (C=O) and "ipc v (C=O) modes whereas the latter two frequencies are assigned to ring stretching modes. The two γ (C=O) modes are assigned at 757 and 745 cm⁻¹ whereas the 2 β (C=O) modes are assigned as the frequencies 412 and 321 cm⁻¹ in

Page 5 of 10

	IR					
C No Theoretical wave		l wave	Experimental wave	Vibrational assignment (%/RED)		
5. NO.	number	1	number	Vibrational assignment (%PED)		
	Unscaled	Scaled*				
1	31	31	-	$\delta(O_{22}-C_{23})[59] + \Box(R)[14] + \gamma(C_1-N_8)[9] + \gamma(C_4-N_7)[9]$		
2	37	36	-	$T(O_{22}-C_{23})[31] - \delta(C_{23}-C_{26})[19] + T(C_1-N_8)[8] + \beta(C_3-O_{22})[8] + \Box(R)[8]$		
3	56	55	-	$δ(O_{22}-C_{23})[70] + τ(C_1-N_8)[9] + □(R)[7]$		
4	63	62	-	□(R)[35]+τ(C ₄ -N ₇)[15]+δ(O ₂₂ -C ₂₃)[17]		
5	82	81	-	$T(C_{2}-O_{2})[25] + \delta(C_{2}-C_{2})[22] + v(C_{2}-O_{2})[21] + \delta(O_{2}-C_{2})[10] + \Box(R)[9]$		
6	108	107	_	$\frac{1}{3} = \frac{1}{22} + \frac{1}{23} = \frac{1}{26} + \frac{1}{23} = \frac{1}{26} + \frac{1}{22} = \frac{1}{22} + \frac{1}{23} = \frac{1}{22} + \frac{1}{23} +$		
-	100	107	-	$\frac{\gamma(n_7 + n_2)}{(1 + 1)^2} = \frac{\gamma(n_7 - n_3)}{(1 + 1)^2} = \gamma(n_7 - n_3)$		
1	129	128	-	$\frac{\gamma(N_8 - H_{13})[32] + \Box(R)[27] + \tau(C_1 - N_8)[13] + \tau(N_8 - C_{14})[10] + \Box(R)[9]}{(3 - 1)^{10}}$		
8	147	146	-	$\gamma(C_4 - N_7)[38] + \Box(R)[22] + \tau(C_4 - N_7)[9] + \tau(N_7 - C_{18})[8]$		
9	168	167	-	$\gamma(C_{3}-O_{22})[38] + \Box(R)[17] + \gamma(C_{4}-N_{7})[17] + \delta(O_{22}-C_{23})[8]$		
10	173	171	-	$T(N_8 - C_{14})[60] + \Box(R)[10] + T(C_1 - N_8)[9] + \Box(R)[9]$		
11	190	189	-	$\beta(C_4 - N_7)[27] + \beta(C_6 - Cl_9)[20] + \delta(N_8 - H_{13})[10] + \beta(C_1 - N_8)[10] + \delta(N_7 - H_{12})[9]$		
12	220	218	-	$\gamma(C_{e}\text{-}Cl_{o})[44] + \Box(R)[12] + \Box(R)[11]$		
13	224	222	-	$\beta(C_{c}-Cl_{c})[29] + \gamma(C_{c}-Cl_{c})[21] + \beta(C_{c}-N_{c})[20]$		
14	235	233	-	$T(C_{1}, C_{2}) = \frac{1}{2} \left[\frac{1}{2} + \frac{1}{$		
15	250	257		$\frac{1}{23} = \frac{2}{26} \frac{1}{163} + \frac{1}{2} \frac{1}{26} = \frac{1}{2} \frac{1}{163} + \frac{1}{2} \frac{1}{$		
10	209	207	-	$\frac{1}{(O_{23}-O_{26})[10] + p(O_{6}-O_{19})[13] + 0(10_{8}-O_{14})[10] + v(O_{4}-O_{5})[1]}{(O_{10}-O_{10})[10] + p(O_{6}-O_{10})[10] + p(O_{6}-O_{10})[1$		
16	271	269	-	$\alpha(R)[22] - O(N_7 - C_{18})[11] - O(N_8 - H_{13})[11] - \beta(C_6 - CI_9)[10]$		
17	324	321	-	$\gamma(C_3 - O_{22})[28] - \beta(C_5 - O_{10})[15] - \beta(C_2 - O_{11})[15] + \alpha(R)[14] - \delta(N_8 - H_{13})[10]$		
18	335	332	-	$\beta(C_3 - O_{22})[45] - \gamma(C_3 - O_{22})[13]$		
19	356	353	-	$\delta(C_{23}-C_{26})[26] + \gamma(C_3-O_{22})[17] + \delta(O_{22}-C_{23})[17] - \nu(C_6-Cl_9)[10] - \alpha(R)[7]$		
20	379	376	-	$\gamma(C_3 - O_{22})[34] + \delta(O_{22} - C_{23})[24] - \gamma(C_6 - Cl_9)[7]$		
21	416	412	-	$\beta(C_r-O_{r,r})[21] - \beta(C_s-O_{r,r})[20] + \delta(N_s-H_{r,r})[9]$		
22	461	457	-	$\alpha(R)[35] = \delta(N - H)[10] = \delta(N - H)[9] = \alpha(R)[8] = \beta(C - O)[7]$		
23	512	507		$\frac{V(1)[00]}{V(1-7)} = \frac{V(1-7)}{V(1-7)} = $		
23	512	507	-	$\frac{\gamma(C_4 - N_7)[13] - 0(C_{22} - C_{23})[12] + 0(N_8 - \Pi_{13})[10] + 0(C_3 - 0_{22})[1]}{\gamma(C_3 - 0_{22})[1]}$		
24	538	533	-	$\gamma(C_4 - N_7)(20) + \gamma(C_1 - N_8)(20) + O(O_{22} - C_{23})(15) + \gamma(C_2 - O_{11})(10) + \Box(R)(10)$		
25	557	551	569	$\delta(N_7 - H_{12})[15] + \beta(C_2 - O_{11})[11] - \nu(C_2 - C_3)[11] - \nu(C_3 - C_4)[8] - \nu(C_6 - Cl_9)[8]$		
26	613	608	-	$\gamma(C_{3}-O_{22})[29] - \Box(R)[20] - \delta(O_{22}-C_{23})[10] + \delta(C_{23}-C_{26})[26] + \gamma(C_{4}-N_{7})[7] - \gamma(C_{1}-N_{8})[7]$		
27	650	644	641	$\gamma(C_3 - O_{22})[21] - \alpha(R)[12] - \gamma(N_7 - H_{12})[10] - \gamma(C_1 - N_8)[8]$		
28	677	670	-	$\gamma(C_3 - O_{22})[36] + \tau(C_4 - N_7)[20] + \gamma(N_7 - H_{12})[13] - \Box(R)[8]$		
29	707	700	-	$V(N_{o}-H_{1,0})[38] + T(C_{o}-N_{o})[36] - V(C_{o}-C_{c})[8]$		
30	753	745	-	$\Box(R)[32] + v(C - O)[30] - v(C - O)[14] + v(C - O)[7]$		
31	764	757	752	$v(C_{-}O_{-})[27] + \Box(R)[21] + v(C_{-}O_{-})[17] - v(C_{-}N_{-})[14] - \Box(R)[11]$		
32	704	784	102	$\frac{P(C_5 - C_{10})[21] + P(C_2 - C_{11})[11]}{P(C_4 - C_2)[12]} + \frac{P(C_4 - C_2)[12]}{P(C_4 - C_2)[12]} + \frac{P(C_4 - C_2)[21]}{P(C_4 - C_2)[12]} + \frac{P(C_4 - C_2)[22]}{P(C_4 - C_2)[12]} + \frac{P(C_4 - C_2)[22]}$		
32	040	044	-	$p(O_5^{-}O_{10}/L^{24}) + v(O_6^{-}O_{19}/L^{20}) + p(O_4^{-}v_7/L^{0}) + p(O_3^{-}O_{22}/L^{7}) = v(v_7^{-}O_{18}/L^{7})$		
33	010	011	-	$0(\bigcup_{26} - \prod_{27})[37] = 0(\bigcup_{23} - \prod_{25})[37] + 0(\bigcup_{23} - \prod_{25})[11]$		
34	834	826	-	$\beta(C_1 - N_8)[18] + \beta(C_2 - O_{11})[13] + \beta(C_3 - O_{22})[9] - \nu(C_6 - CI_9)[9]$		
35	871	863	890	$v(O_{22}-C_{23})[34] + \delta(C_{26}-H_{27})[22] + \gamma(C_3-O_{22})[11]$		
36	983	974	-	$v(N_8-C_{14})[14] + v(C_1-C_2)[14] + \beta(C_1-N_8)[8] - \beta(C_3-O_{22})[7]$		
37	1012	1002	-	$v(C_{23}-C_{26})[17] + \beta(C_4-N_7)[9] + v(N_8-C_{14})[9] + v(C_4-C_5)[8] - \delta(C_{26}-H_{27})[8] + \beta(C_3-O_{22})[7] + v(N_7-C_{18})[7]$		
38	1032	1022	-	$v(C_{23}-C_{26})[22] - v(N_{7}-C_{46})[21] - v(O_{22}-C_{23})[12] - v(N_{8}-C_{46})[11]$		
39	1075	1065	965	$v(N_{r}-C_{r_{0}})[18] - v(O_{r_{0}}-C_{r_{0}})[15] - v(C_{r_{0}}-C_{r_{0}})[12] + v(C_{r_{0}}-C_{r_{0}})[9] - \alpha(R)[7]$		
40	1119	1088	1128	$v(C_{-}O_{-})[19] - v(O_{-}-C_{-})[17] - v(C_{-}C_{-})[14] + v(N_{-}C_{-})[13] + \delta(C_{-}H_{-})[7]$		
41	1134	1103	-	$\delta(C, -H)$ [41] = v(C, -C,)[18]		
12	1130	1109		$\frac{\delta_{26} + \delta_{27} + \delta_{17} + \delta_{23} + \delta_{26} + \delta_{23} + \delta_{26} + \delta_{27} + \delta_{26} + \delta_{27} +$		
40	11.00	1100	-	$\frac{1}{2} \frac{1}{15} \frac{1}{10} $		
43	1142	1111	-	0(U ₁₈ -H ₁₉)[//] = 0(U ₁₈ -H ₂₁)[8]		
44	1173	1141	-	$\delta(C_{23}-H_{25})[50] + \delta(C_{26}-H_{27})[29] + \delta(C_{23}-H_{25})[8]$		
45	1187	1154	-	$\delta(C_{14}-H_{16})[33] + \delta(C_{14}-H_{15})[11] + \delta(C_{14}-H_{16})[7]$		
46	1206	1173	-	$\delta(C_{18}-H_{20})[35] - \delta(N_7-H_{12})[9] + \delta(C_{18}-H_{20})[7] - v(C_3-O_{22})[7]$		
47	1268	1234	-	$v(C_{1}-N_{8})[13] + \beta(N_{8}-H_{13})[10] - \beta(C_{1}-N_{8})[9] + v(C_{5}-C_{6})[8] - v(N_{8}-C_{14})[7] + \beta(C_{5}-O_{10})[7]$		
48	1294	1259	-	$\delta(C_{23}-H_{23})[33] - v(C_4-N_7)[14] - \beta(N_7-H_{12})[9] + v(C_3-O_{23})[8]$		
49	1304	1269	-	$\delta(C_{n-}H_{n-})[39] + v(C_{r-}N_{n-})[13] + \beta(N_{n-}H_{n-})[9] - v(C_{n-}O_{n-})[9]$		
50	1337	1301	1332	v(C - C)[39] = v(C - C)[13] = v(C - N)[11] = v(C - C)[8] + R(C - O)[7]		
51	1350	1322		$\frac{1}{2} = \frac{1}{2} $		
51	1309	1022	-	$\frac{v(c_2 - c_3)(c_3) - v(c_3 - c_2)(c_3) - v(c_4 - c_2)(c_3) + p(c_2 - c_4)(c_3)}{v(c_4 - c_2)(c_3 - c_2)(c_3$		
52	1401	1362	1405	$0(U_{26}-H_{27})[00] - 0(U_{23}-H_{24})[31]$		
53	1423	1384	-	$0(C_{23}-H_{24})[45] + 0(C_{26}-H_{27})[31] + v(C_{23}-C_{26})[11]$		
54	1463	1423	-	$\delta(C_{14}-H_{15})[72] - \delta(C_{18}-H_{19})[14]$		
55	1468	1428	-	$\delta(C_{18}-H_{19})[70] + \delta(C_{14}-H_{15})[16]$		

Page 6 of 10

56	1482	1442	-	$\delta(C_{26}-H_{28})[61] - \delta(C_{26}-H_{29})[28] - \delta(C_{26}-H_{27})[8]$		
57	1485	1444	-	δ(C ₁₈ -H ₂₁)[87] - δ(C ₁₈ -H ₁₉)[8]		
58	1489	1449	1470	δ(C ₁₄ -H ₁₇)[91] - δ(C ₁₄ -H ₁₅)[7]		
59	1501	1460	-	$\delta(C_{26}-H_{29})[52] + \delta(C_{28}-H_{28})[23] - \delta(C_{23}-H_{24})[10] - \delta(C_{26}-H_{27})[7]$		
60	1507	1466	-	$\delta(C_{18}-H_{20})[66] - \delta(C_{14}-H_{16})[11] - \delta(C_{18}-H_{20})[7]$		
61	1508	1467	-	$\delta(C_{14}-H_{16})[72] + \delta(C_{18}-H_{20})[11] - \delta(C_{14}-H_{16})[7]$		
62	1517	1476	-	$\delta(C_{23}-H_{24})[86] + \delta(C_{26}-H_{29})[6]$		
63	1557	1514	1554	β(N ₈ -H ₁₃)[39] - β(N ₇ -H ₁₂)[13]- ν(C ₂ -O ₁₁)[9]		
64	1564	1522	-	$\beta(N_7 - H_{12})[35] + \beta(N_8 - H_{13})[13] + v(C_4 - N_7)[8]$		
65	1628	1584	-	$v(C_1-C_6)[23] - v(C_3-C_4)[20] - v(C_2-O_{11})[11] - v(C_1-N_8)[10]$		
66	1639	1595	1602	$v(C_1-N_8)[15] - v(C_3-C_4)[14] - v(C_1-C_6)[14] - v(C_5-O_{10})[8] - v(C_2-O_{11})[8] + v(C_4-N_7)[7]$		
67	1685	1639	-	$v(C_2 - O_{11})[32] + v(C_5 - O_{10})[29] - v(C_3 - C_4)[13] + v(C_4 - N_7)[7]$		
68	1707	1661	1692	$v(C_5-O_{10})[36] - v(C_2-O_{11})[18] + \alpha(R)[9] + v(C_1-N_8)[8] - v(C_4-N_7)[7]$		
69	2992	2910	2842	v(C ₂₃ -H ₂₄)[94]		
70	3033	2950	2989	$v(C_{18}-H_{19})[69] + v(C_{18}-H_{21})[14] + v(C_{18}-H_{20})[11]$		
71	3034	2951	3015	$v(C_{26}-H_{29})[35] + v(C_{26}-H_{27})[34] + v(C_{26}-H_{28})[26]$		
72	3049	2966	3042	$v(C_{14}-H_{16})[40] + v(C_{14}-H_{15})[34] + v(C_{14}-H_{17})[26]$		
73	3081	2997	-	$v(C_{23}-H_{25})[71] + v(C_{26}-H_{29})[15]$		
74	3101	3017	-	$v(C_{26}-H_{27})[53] - v(C_{26}-H_{29})[23] + v(C_{23}-H_{25})[12] - v(C_{26}-H_{28})[9]$		
75	3102	3017	-	$v(C_{18}-H_{21})[39] + v(C_{18}-H_{20})[31] - v(C_{18}-H_{19})[27]$		
76	3110	3026	-	$v(C_{26}-H_{28})[59] - v(C_{26}-H_{29})[25] + v(C_{23}-H_{25})[11]$		
77	3114	3029	-	v(C ₁₄ -H ₁₆)[52] - v(C ₁₄ -H ₁₅)[47]		
78	3127	3042	-	v(C ₁₄ -H ₁₇)[72] - v(C ₁₄ -H ₁₅)[19] - v(C ₁₄ -H ₁₆)[8]		
79	3135	3049	-	v(C ₁₈ -H ₂₀)[55] - v(C ₁₈ -H ₂₁)[44]		
80	3479	3385	3356	v(N ₈ -H ₁₃)[99]		
81	3515	3419	-	v(N ₇ -H ₁₂)[99]		

The numbers (in bracket) after the modes are the % potential energy calculated using normal coordinate analysis. The abbreviations are: v=stretching, ω =wagging, r=twisting, ρ =rocking, σ =scissoring, γ =out-of-plane deformation, β =in-plane deformation, α =planar ring deformation, ϕ =non-planar ring deformation, δ =deformation. The modes in bold letters in the last column are the assigned modes.

*With the scale factor of 0.9906 for calculated wave numbers lower than 1100 cm⁻¹ and the scale factor 0.9728 for larger wave numbers.

Table 4: Vibrational assignments of cemb.

accordance with the assignment reported in literature [52].

Ring modes (12 modes): In addition to the two ring stretching modes mentioned above (section 5.1.2) the frequencies 1321 (ν_{51}), 1300 ($_{50}$), 974 (ν_{36}) and 551 (ν_{25}) cm⁻¹ have been identified as the ring stretching modes. The lowest frequency mode 551 (ν_{22}) cm⁻¹ corresponds to the ring breathing mode. Out of the three planar ring deformation frequencies 641 (ν_{27}), 507 ($_{23}$) and 457 (ν_{22}) cm⁻¹ the first frequency appears to be due to the trigonal phenyl ring bending mode whereas the latter two correspond to the modes 6 (a,b) of the benzene molecule. The three normal modes of vibration corresponding to the non-planar ring deformations are calculated to have magnitudes 608 (ν_{26}), 146 ($_{8}$) and 128 (ν_{7}) cm⁻¹. All the 12 normal modes of the ring are strongly coupled with the other modes.

(C-NHCH₃) modes (6 modes): The frequencies 1233 ($_{47}$) and 1258 (ν_{48}) cm⁻¹ corresponds to ν (C₁-NHCH₃) and ν (C₄-NHCH₃), respectively. The two planar C-NHCH₃ bending mode are found to have widely differing magnitudes 826 ($_{34}$) cm⁻¹ and 188 (ν_{11}) cm⁻¹. The non-planar bending modes are calculated to have frequencies 167 ($_{9}$) cm⁻¹ and 3 1(ν_{1}) cm⁻¹.

N-H modes (6 modes): The wave numbers 3419 (ν_{81}) cm⁻¹ and 3384 ($_{80}$) cm⁻¹ correspond N₇-H and N₈-H stretching vibrations. The N-H stretching vibrations are pure and highly localized modes. The β (N₇-H) and β (N₈-H) modes are calculated to have magnitudes 1521 (ν_{64}) cm⁻¹ and 1514 ($_{63}$) cm⁻¹, respectively. The out-of-plane deformations corresponding to the modes γ (N₇-H) and γ (N₈-H) are correlated to the calculated frequencies 670 ($_{78}$) cm⁻¹ and 700 ($_{79}$) cm⁻¹ respectively.

 ${}^{\tiny @}\,^{n}C$ means C atom with number n; ${}^{\tiny @}opc{=}out$ of phase coupled and ipc=in phase coupled

N-CH₃ modes (6 modes): The stretching modes corresponding to ν (N₇-CH₃) and ν (N₈-CH₃) are found at the frequencies 1064 (₃₉) cm⁻¹ and 1002 (ν_{37}) cm⁻¹ respectively. The twisting modes are calculated to have magnitudes 62 (ν_1) and 55 (ν_3) corresponding to the modes τ (N₇-CH₃) and (N₈-CH₃), respectively while the angle bending modes are assigned at 268 (ν_{16}) and 257 (ν_{15}) wave numbers corresponding to the modes α (C₄-N₇-CH₃) and α (C₇-N₈-CH₃).

C-Cl modes (3 modes): In benzene derivatives C-Cl stretching is observed in wide spectral range 700-1050 cm⁻¹ [53]. The present calculations place the C-Cl stretching mode at frequency 783 (ν_{32}) cm⁻¹. The modes corresponding to the planar deformation mode β (C-Cl) and non-planar deformation mode γ (C-Cl) are at frequencies 222 (ν_{13}) and 218 (ν_{12}) cm⁻¹ respectively.

C-O(CH₂)(CH₃) modes (3 modes): The C₃-O(CH₂)(CH₃) bond gives rise to 3 normal modes of vibration as: a ν [C₃-O(CH₂)(CH₃)], a β [C₃-O(CH₂)(CH₃)] and a γ [C₃-O(CH₂)(CH₃)] modes. The wavenumbers 1088 (ν_{40}), 332 (ν_{18}) and 376 (ν_{20}), correspond to the above three modes respectively.

 $\text{O-CH}_2(\text{CH}_3)$ modes (3 modes): Three normal modes of vibrations ν (O₂₂-CH₂(CH₃)), τ (O₂₂-CH₂(CH₃)) about C₃-O₂₂ bond and α (C₃-O₂₂-CH₂(CH₃)) arise due to O₂₂-CH₂(CH₃) bond. The stretching mode is found at the frequency 862 (₃₅) cm⁻¹. The twisting mode is calculated to have magnitude 81 (₅) cm⁻¹ while the angle bending mode is assigned at

533 (v_{24}) wave number.

5.3.10 C(H₂)-C(H₃) modes (3 modes): Three normal modes of vibrations ν (C-C), τ (C-C) and α (O-C-C) arise due to H₂C-CH₃ modes. The stretching mode corresponding to ν (C₂₃-C₂₆) is found at the frequency 1022 cm⁻¹ (₃₈). The twisting mode is calculated to have magnitude 36 (ν_2) cm⁻¹ while the angle bending mode is assigned at 352 (ν_{19}) wave number.

Molecular electrostatic potential and contour maps

The molecular electrostatic potential (MEP) mapped surfaces illustrate the charge distributions of molecules three dimensionally which give clear signature of the interactions of the molecules. The electrostatic potential is used to find the reactive site of a molecule. The MEP mapped surface of the molecules are calculated by DFT/B3LYP/6-311G (d,p) method at the 0.02 isovalues and 0.004 density values. In color scheme of MEP plot, the red region show atoms with lone pairs or negative electrostatic potentials; the intensity of the color is proportional to the absolute value of the potential energy. The positive electrostatic potentials are shown in blue/yellow color region and characterize the polar hydrogen in O-H bonds. The Green areas cover parts of the molecule where electrostatic potentials are close to zero (C–C, C–N and C–S bonds). The MEP mapped surface of the cemb is shown in Figure 4(B).

Molecular electrostatic potential and electrostatic potential are the useful quantities to illustrate the charge distributions of molecules are used to visualize variably charged regions of a molecule. Therefore, the charge distributions can give the information about how the molecules interact with another molecule. At any given point r (x, y, z) in the vicinity of a molecule, the molecular electrostatic potential (MEP), V(r) is defined in terms of the interaction energy between the electrical charge generated from the molecule electrons and nuclei and a positive test charge (a proton) located at r. The molecular electrostatic potential is related to the electronic density and a very useful descriptor for determining sites for electrophilic attack and nucleophilic reactions as well as hydrogen-bonding interactions [54]. In Figure 4b and 4c, whereas electrophilic reactivity is presented by the negative (red) regions, nucleophilic reactivity is shown by the positive (blue) regions of MEP. The red region is localized on the vicinity of oxygen and reflects the most electronegative region (excess negative charge). On the other hand, the nucleophilic reactivity of the molecules is localized on the hydrogen atoms. In this respect, the compound (cemb) is found to be useful to both bonds metallically and interact intermolecularly. Based on these results, it can be ascertained that how the change of the chlorine group affects the molecular properties of the compound. One can see from the Figure 4b that the blue color on methyl amine is found to be the predominant because of the positive charge distribution on the molecule. Moreover, according to the ESP Figure 4c, the most negative ESP is spread over the oxygen and chlorine atoms, indicating the delocalization of electrons over the oxygen and chlorine atoms of the molecule which also reveal extended conjugation of the benzene ring with these groups [55]. Furthermore, the total charge density contour map is given in the Figure 4d. A contour plot is a two-dimensional XY plot of a three-dimensional XYZ surface showing lines where the surface intersects planes of constant elevation (Z). The contour maps are used to show lines of constant density or brightness, such as electrostatic potentials and are drawn in the molecular plane. The electron rich red lines are around oxygen and nitrogen whereas electron deficient region are shown by greenish-yellow lines. The contour maps are also Page 7 of 10

calculated by same basis sets at 0.004 density values at same level of calculations of the MEP mapped surface of the molecules [56].

Frontier molecular orbital analysis

The analysis of the wave function indicates that the electron absorption corresponds to the transition from the ground to the first excited state and is mainly described by one-electron excitation from the HOMO to LUMO. Both the HOMO and LUMO are the main orbitals taking part in the chemical reaction. The HOMO energy characterizes the capability of electron donating; The LUMO characterizes the capability of electron accepting [57]. The HOMO and LUMO, calculated at the B3LYP/6-311G (d, p) level can be seen in Figure 7 for the gas phase. The positive phases are marked as red and the negative ones as green. Figure 7 shows distribution and energy levels of the HOMO and LUMO orbitals. For the HOMO of cemb, electron density are localized on all atom except on methyl of ethoxy group with E_{HOMO} =-0.21294 a.u. For the LUMO, electron density are localized on the entire atom except on ethoxy group with E_{UMO} =-0.10412. The value of the energy separation between the HOMO and LUMO is 0.10882 a.u (2.961068 eV). In the contour map, the lines with red colors are regarded as the negative phase and the lines with green colors are the positive phase. The contour HOMO-LUMO diagram is in accordance with the frontier orbital. The frontier orbital gap helps to characterize the chemical reactivity, optical polarizability and chemical hardnesssoftness of a molecule [58]. The small HOMO-LUMO energy gap means low excitation energy for the compound, a good stability and a low chemical hardness. The molecules having a large energy gap are known as hard, and molecules having a small energy gap are known as soft molecules. The soft molecules are more polarizable than the hard ones, because they need small energy for excitation. The hardness value of a molecule can be determined by the formula [59]

$$\eta = \{-E_{HOMO} + E_{LUMO}\}/2$$

where E_{HOMO} and E_{LUMO} are the energies of the HOMO and LUMO molecular orbitals. The value of η of cemb, 0.05441 a.u (1.480534 eV), indicates for the compound should be a soft material.

Global reactivity descriptors

On the basis of Koopman's theorem, global reactivity descriptors [60] are calculated using the energies of Frontier molecular orbitals E_{HOMO} , E_{LIMO} and given by Equations (a)–(e).

$$\chi = -(E_{\text{HOMO}} + E_{\text{LUMO}})/2 = (I + A)/2 \text{ (a)}$$
$$\mu = -\chi = (E_{\text{HOMO}} + E_{\text{LUMO}})/2 = -(I + A)/2 \text{ (b)}$$



Pharm Anal Acta ISSN: 2153-2435 PAA, an open access journal

Page 8 of 10

$$\eta = (-E_{\text{HOMO}} + E_{\text{LUMO}})/2 = (I - A)/2 \text{ (c)}$$

where *A* and *I* are the ionization potential and electron affinity of the compounds respectively. Electron affinity refers to the capability of a ligand to accept precisely one electron from a donor. However, in many kinds of bonding *viz.*, covalent hydrogen bonding, and partial charge transfer occurs. Softness (*S*) is the reciprocal of hardness and is a property of compound that measures the extent of chemical reactivity.

 $S = 1/\eta$ (d)

Recently Parr et al. [61] have defined a new descriptor to quantity the global electrophilic power of the compound as electrophilicity index (ω), which defines a quantitative classification of the global electrophilic nature of a compound. Parr et al. have proposed electrophilicity index (ω x) as a measure of energy lowering due to maximal electron flow between donor and acceptor. They defined electrophilicity index (ω) as follows

 $\omega = \mu^2 \,\mathrm{S} = \mu^2 / \eta \;(\mathrm{e})$

The energies of frontier molecular orbitals (E_{HOMO} , E_{LUMO}), energy band gap (E_{HOMO} , E_{LUMO}), electro negativity (χ), ionization potential (A), electron affinity (I), chemical potential (μ), global hardness (η), global softness (S) and global electrophilicity index (ω) for cemb are listed in Table 5.

In-vitro antibacterial activity

Six bacterial pathogens Salmonella typhi, Escherichia coli, Shigella flexneri, Staphylococcus aureus, Aeromonas hydrophila and Enterococcus faecalis were used to test the antibacterial activity of the cemb. The activity of cemb and commercial antibacterial drugs streptomycin sulphate and chloramphenicol were tested against the bacterial pathogens and the inhibition zones (cm) of the compounds were presented in the Table 6. The test results of the compound have also been compared with those of the standard drugs and results show significant activity of the compound against S. aureus than all other test pathogens. The cemb displays higher zone of inhibition (3.5 cm) against S. aureus at 100 µg/disc and does not display any zone of inhibition against Aeromonas hydrophila and E.coli at all concentration. A clearing zone around the disc indicates the inhibitory activity of the compound on the organism (Figure 8a-8d). The results are given in Table 6 which clearly indicate that the inhibitions by the compound is lower as compared to the standard drugs and activity increases with increasing concentration of the compound (since concentration plays a vital role in increasing the degree of inhibition) which is almost comparable to the size of the reference antibiotics. 1,4-Benzoquinones is a crossconjugated enediones, have ability to accept one electron initially, forming the semiquinone anion radical, followed by a further electron to give the dianion, is the dominant feature of quinone chemistry. It is this reversible reduction process that accounts for the biological activity [62]. Apart from this, other factors such as solubility, conductivity and intermolecular interaction may also be the possible reasons for activity. Furthermore, free HNCCO moiety increases the delocalization of π -electrons over the molecule of cemb and enhances the lipophilicity

Molecular properties	Values in eV		
Electronegativity (χ)	0.317		
Chemical potential (µ)	-0.317		
Global hardness (η)	0.545		
Global softness (s)	1.835		
Electrophilicity Index (ω)	-0.184		

Table 5: Reactivity parameters of cemb.



Figure 8: Antibacterial activity of cemb (A) Inhibition zone (IZ) of cemb against *S. aureus* (Center disc=DMSO, -ve control) (B) IZ against *S. flexneri* (C) IZ against of cemb against *S. typhi* (D) No inhibition of cemb against *E. coli*

	Concentration	Compound	Standard drug	
Test pathogen	of compound (µg/disc)	(cemb)	Streptomycin sulphate	Chloramphenicol
	10	-	2.0	2.4
	20	-	2.2	2.5
Salmonella	40	-	2.4	2.6
typni	60	0.9	2.8	2.9
	80	1.2	3.1	3.2
	100	1.3	3.5	3.7
	10	0.6	2.0	2.6
	20	0.8	2.4	2.9
Shigella flexneri	40	0.9	2.6	3.0
_	60	1.1	3.0	3.3
	80	1.3	3.3	3.5
	100	1.5	3.6	3.7
	10	1.8	2.0	2.2
	20	2.0	2.1	2.4
Staphylococcus	40	2.4	2.3	2.5
aureus	60	2.7	2.6	2.8
	80	2.8	2.9	3.2
	100	3.5	3.6	3.8
	10	-	2.2	1.6
	20	-	2.3	1.9
Enterococcus	40	-	2.5	2.0
faecalis	60	0.7	2.6	2.1
	80	0.9	3.0	2.3
	100	1.2	3.2	2.5

No zone was observed *inhibition zone is the average of the diameter of zone from two sides in cm. (*A. Hydrophila, E. coli* does not show activity).

 $\ensuremath{\text{Table 6:}}$ Antibacterial assay of the cemb and standard drugs against human bacterial pathogens.

Page 9 of 10

which leads to the breakdown of the permeability barrier of the cell and thus retards the normal cell processes. The mode of action may involve the formation of a hydrogen bond through the nitrogen /oxygen atom in the compound with the active centers of the cell constituents, resulting in interference with the normal cell process, i.e., prolonging the lag phase of the growth cycle and increasing the generation time of the organisms so that it takes each organism more time to complete cell division [63].

Conclusions

This paper reports the synthesis, spectral and crystal structure investigations of a new compound: 1,4-benzoquinone derivative, 2-chloro-5-ethoxy-3,6-bis (methylamino)-1,4-benzoquinone (cemb), formed via condensation reaction. It was found that during the reaction, chlorine group of the compound is replaced with ethoxy group generating a new molecule which is stabilized via weak intermolecular interactions. The results of antibacterial activity shows that the new compound cemb inhibit growth of Gram-positive bacteria. Keeping in view the single molecule approximation, the X-ray crystallographic results are well reproduced by DFT method. From the vibrational analysis it turns out that the ring breathing mode (551 cm⁻¹) and the trigonal ring bending mode (641 cm⁻¹) are considerably reduced as compared to the benzene modes. In the present case one γ (C=O) is found in the range 700-800 cm⁻¹ while the other γ (C=O) and two β (C=O) are found to be in the range 300-400 cm⁻¹ while for 2,5-diamino-3,6-dichloro-1,4-benzoquinone [46] the two y (C=O) and one β (C=O) were found in the range 700-800 cm⁻¹ and the other β (C=O) in the range 300-400 cm⁻¹. Inspite of the low symmetry (C₁) of the molecule the modes of the identical groups couple strongly. According to the MEP, the most negative ESP is spread over the oxygen and chlorine atoms, indicating the delocalization of electrons over the oxygen and chlorine atoms of the molecule which also reveal extended conjugation of the benzene ring.

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Page 10 of 10

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