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Design, DFT Calculations and Antimicrobial Activity of New Synthesized Piperazine Derivative

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Keywords	Abstract
Piperazine Derivatives DFT Calculation HOMO, LUMO Energies Antibacterial Activity MEP	The title compound (2,2'-(piperazine-1,4-diyl)bis(N'-((E)-5-chloro-2-hydroxybenzylidene)acetohydrazide) (5-CIPAH) was synthesized by reacting 1,4-Piperazinediacetic acid, 1,4-dihydrazide and 5-Chloro-2-hydroxybenzaldehyde. Mass spectrometry, ¹ H, ¹³ C-NMR, IR results of the synthesized compound were examined. Many information about physical and chemical properties of 5-CIPAH can be obtained by theoretical calculations. Density functional theory (DFT) is widely used theoretical method for predicting of chemical structures. The structure was optimized using DFT/6311G method with GAUSSIAN09. Frontier Molecular Orbitals (HOMO and LUMO) energies were calculated. Global reactivity descriptors and also electrophilic and nucleophilic regions were defined by molecular electrostatic potential surface. Antibacterial and fungal activity were evaluated.
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1. INTRODUCTION

The imine group (azomethine) of Schiff bases attracted much attention due to widely usage in biological studies and which was use in chemistry (Venkataramana et al., 2010; Yan et al., 2012; Gupta & Goklani, 2017; Ermiş & Durmuş, 2020). Piperazine ring is also an important class of N-heterocyclics having biological activities including antitumor (Xu et al., 2019), antiviral drugs (Hooshmand et al., 2021), antibacterial (Jalagari et al., 2019) and antifungal etc (Suryavanshi & Rathore, 2017) and pharmaceutical applications (Ullh et al., 2022). Piperazine derivatives have been widely used in the production of new drugs in recent years (Shaquiquzzaman et al., 2015).

In this paper, titled compound (**5-CIPAH**) was obtained and structural characterized by elemental analyses, IR, mass spectrometry, ¹H, ¹³C-NMR NMR methods. Antibacterial activity against six bacteria [*B.cereus* 709 ROMA, *S. aureus* ATCC 29213, *K. pneumonia* ATCC 13883, *P. aeruginosa* ATCC 27853, *S. aureus* ATCC 25923, *E. coli* ATCC 25922] was conducted. Microdilution (as MICs) and disc diffusion method were used to determine the antibacterial activity. Moreover, the **5-CIPAH** was evaluated for the activities against three fungal pathogens (*C. tropicalis* M002, *C. parapsilosis* M006 and *C. albicans* ATCC 80018) by agar well diffusion technique. The structure-activity relationship (SAR) of the **5-CIPAH** was also analyzed in the present work. Frontier molecular orbital energies (FMOs), Global reactivity descriptors and MEP were studied at B3LYP/6-311G level of theory.

2. MATERIAL AND METHOD

2.1. Physical Measurements

Used chemicals were purchased from Merck (with high purifity). Fourier Transform Infrared spectra of the was performed between 400 and 4000 cm^{-1} from KBr pellets on. NMR spectras were carried out by Bruker-Spectrospin Avance DPX-400 Ultra-Shield (using d_6 -DMSO solvent). Mass spectrometry was recorded on an Agilent 6470 QQQ LC-MS/MS @ 1290 INF HPLC. The m.p. values were determined using an Opti Melt apparatus. All reactions were watched using Merck silica gel (60 F 254) by thin-layer chromatography (TLC). The elemental analysis was implemented out on a LECO CHNS 9320 type analyzer.

2.2. Synthesis of 2, 2'-(piperazine-1,4-diyl)bis(N'-((E)-5-chloro-2-hydroxybenzylidene)aceto hydrazide (5-CIPAH)

The solution of 1,4-piperazinediacetic acid, 1,4-dihydrazide (0,9 g, 4.5 mmol) (Koparde et al., 2018) in 25 mL of water- ethanol (1-4 rate) was slowly added with hot solution (50 °C) of 5-Chloro-2-hydroxy-benzaldehyde (1.4g, 8.9 mmol) in 30 mL of ethanol and refluxed for 24 h. Reactions were watched by TLC. The precipitated product was cooled to room temperature an $\sim 23^\circ\text{C}$ to crystallized from the ethanol/water (4:1) mixture and dried in vacuo. The reaction equation was given Figure 1. Yield 87%. M.p. 198-199°C. LC-MS (100 eV, APCI): 507.8 ($M+H^+$, 100%), 507.2 ($M+$, 82%), IR (KBr) ν/cm^{-1} : 2931 cm^{-1} , C-H aliph; 3066, C-H arom.; 3187, N-H. Elemental analysis for $\text{C}_{22}\text{H}_{24}\text{Cl}_2\text{N}_6\text{O}_4$ (MW:507.37 g/mol) (Calc.%) C: 52.08; H: 4.77; N: 16.56; Cl: 13.97, O: 12.61. (Found %) C, 51.01; H, 5.11; N, 17.04; O, 12.41; Cl, 14.44.

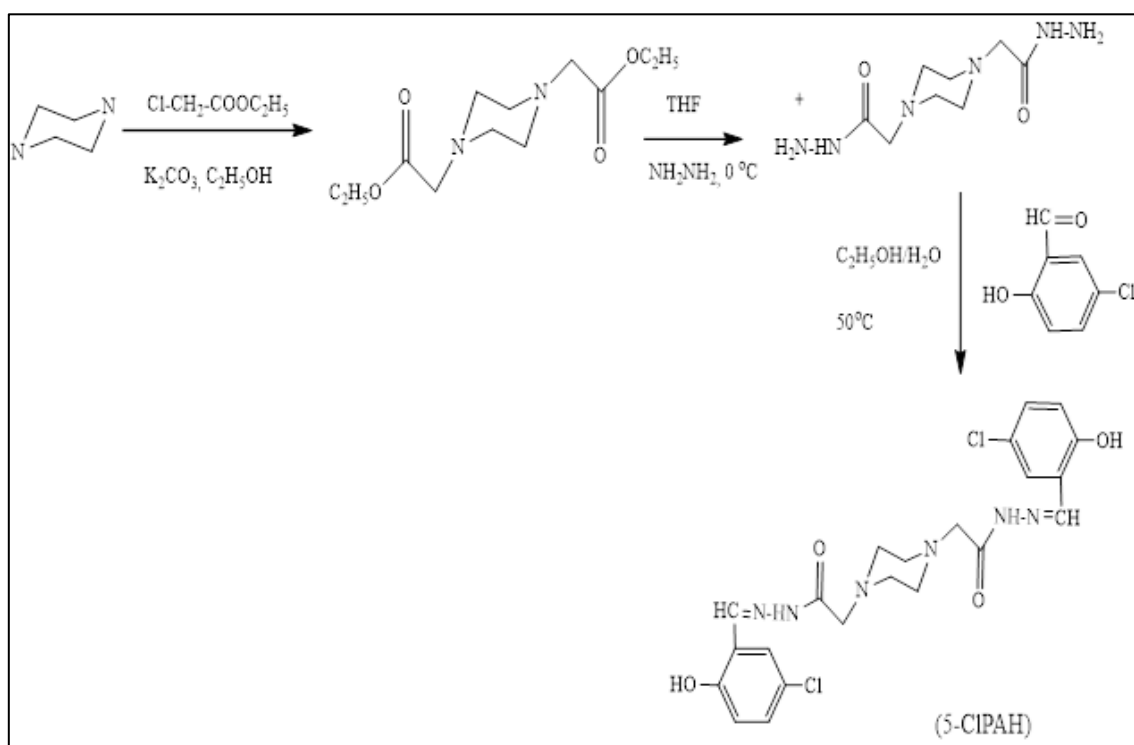


Figure 1. Preparation of 5-CIPAH

2.3. Procedure for Antibacterial Activity

Patogen bacterial cultures incubated for 24 hour at 37°C were acquired from Biology Department of Kırsehir Ahi Evran University.

The little discs 6 mm in diameter were treated with 50 μg Schiff bases and then were located on test plates. The paper discs containing 30 μg streptomycin and 50 μg Ampisilin were used as standard drugs. The **5-CIPAH** were tested against microorganisms three times and the activated zones were detected after a day (Luis Esaú et al., 2019).

A serial concentration was prepared by thin out the stock solution with 20 %DMSO solvent: 1250, 625, 312.5, 156.25, 78.125, 39.06, 19.53, 9.76, 4.88 $\mu\text{g mL}^{-1}$. After incubation at 37°C for 24 h of the MIC of against microbial strain was defined by macroscopic observation (Koneman et al., 1997).

2.4. Antifungal activity (*in vitro*)

Candida albicans ATCC 80018, *Candida tropicalis* M002 and *Candida parapsilosis* M006 strains were obtained from Biology Department of Kırsehir Ahi Evran University. *In vitro*, antifungal activite was studied using agar well diffusion method (Suryavanshi & Rathore, 2017). The 50 μL of microbial suspension was disseminated over plate containing agar surface. The 5-CIPAH and standard drugs Ampisilin and Streptomycin were incubated at 303 K for 48 h.

2.5. Computation details

Molecular modeling of the compound was carried out by Gaussian 09 (Frisch et al., 2009) program using DFT/B3LYP (Prasad et al., 2022) density functional theory with 6-311G (Foresman & Frisch, 1996) base set and semi-empirical theory with AM1 (Austin Model 1) (Dewar et al., 1985) base set. HOMOs and LUMOs as frontier molecular orbitals computed by DFT method play a important role in the chemical reactivities, stabilities and electronic transition levels of molecules (Dennington et al., 2016).

3. RESULT AND DISCUSSION

3.1. Structure of the 5-CIPAH

IR spectrum of **5-CIPAH** is shown in Figure 2. As seen in Figure 2, a characteristic O-H stretching band is observed at 3344,40 cm^{-1} . Other stretching bands are observed at 3187 cm^{-1} (N-H), 3066 cm^{-1} (CH: aromatic), 1688 cm^{-1} (C=O; amide), 1512 cm^{-1} (C=C: aromatic), 1294 cm^{-1} (C-N).

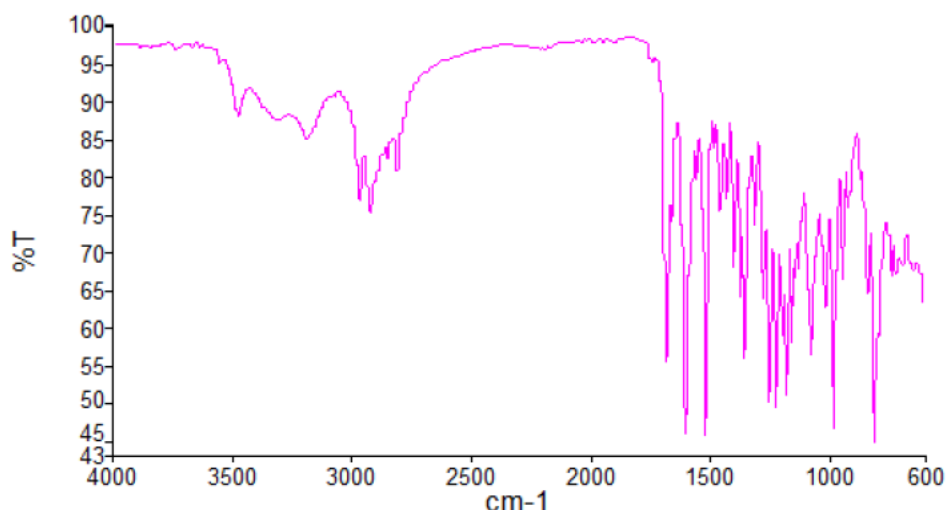


Figure 2. IR spectrum of **5-CIPAH**

The NMR spectrums (^1H , ^{13}C) of 5-CIPAH were obtained and interpreted in *d6* -DMSO. In ^1H -NMR, the protons of piperazine appeared at 2.38 (m, 4H) and 2.62 (m, 4H) ppm respectively. The protons belonging to N=CH (8.4 ppm), and C-NH (11.2–11.51 ppm) were detected a singlet. The aromatic-H peaks were observed between 6.94-7.60 ppm.

In ^{13}C NMR of Schiff base, piperazine ring peak and azomethine CH=N carbon peak were observed at 53.60 ,163.5 ppm, respectively. The aromatic-C peaks were also observed between 107.7-156.5 ppm.

The mass spectrum of 5-CIPAH is presented in Figure 3. Molecular ion $[\text{M}+\text{H}]^+$ peak is observed as base peak at 507.8 (m/z). The main peaks (82%), $[\text{M}^+]$ at (m/z) 507.2 and (75%), $[\text{M}+3\text{H}]^+$.at (m/z) 509.8 are observed.

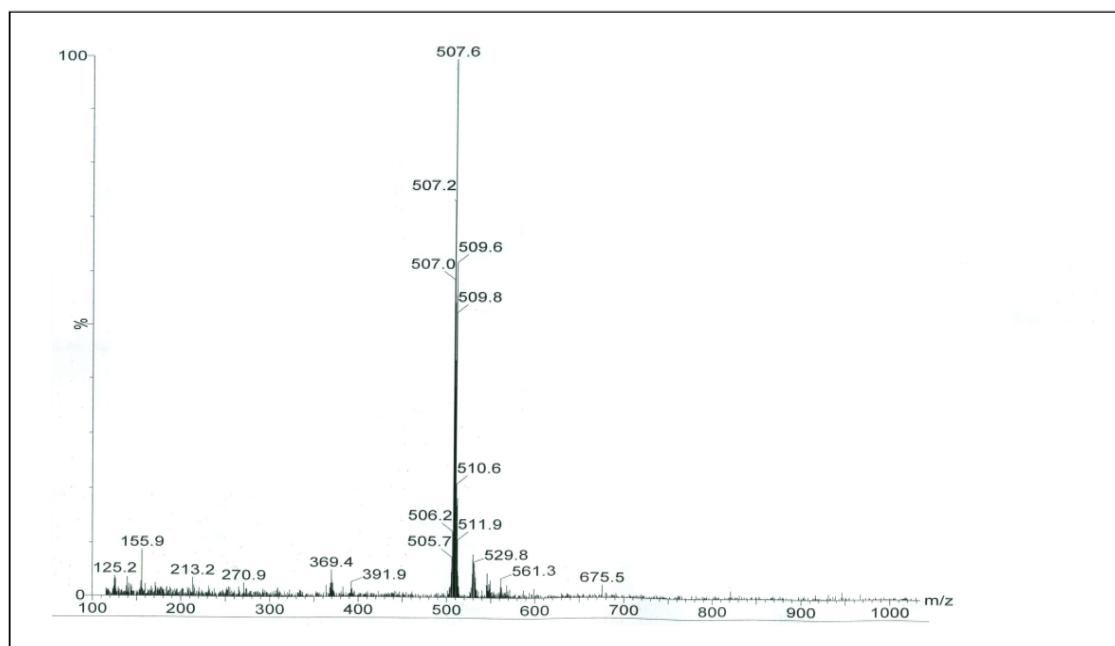


Figure 3. Mass spectrum of 5-CIPAH

3.2. Antimicrobial activity results

The tested strains selected for this study were categorized as three Gram negative bacteria (*E. coli*, *K. pneumoniae*, *P. aeruginosa*), three Gram positive bacteria (*S. aureus**, *B. cereus*, *S. aureus*) and three fungi (*C. tropicalis*, *C. albicans* and *C. parapsilosis*). The 5-CIPAH was dissolved in DMSO (20%) at proper concentration. The antibacterial results were given in Table 1, Table 2 by MICs and disc diffusion, respectively. The positive control used in this study as antibacterial and antifungal agents is Ampicillin, and Streptomycin and were compared (Figure 4).

As the disc diffusion technique results show that the compound exhibits a moderate inhibition effect against tested all bacteria. The compound was significant activity against *P. aeruginosa* ATCC whereas Ampicillin and Streptomycin, the drugs used as standart, have been determined a bit active (10-12 mm) against the six bacteria.

Table 1. Measured diameter (mm) of the compounds

Compounds	Gram-negative			Gram-positive		
	<i>K.pneumoniae</i> ATCC 13883	<i>E.coli</i> ATCC 25922	<i>P.aeruginosa</i> ATCC 27853	<i>B. cereus</i> Roma 709	<i>S. aureus</i> ATCC 29213	<i>S.aureus</i> ATCC 25923
5-CIPAH	15	15	16	10	10	8
Streptomycin	12	20	10	15	18	19
Ampicillin	12	16	12	15	18	18

<10: weak; >10: moderate; >16: significant Ampicillin (50 µg/disk) Streptomycin (30 µg/disk)

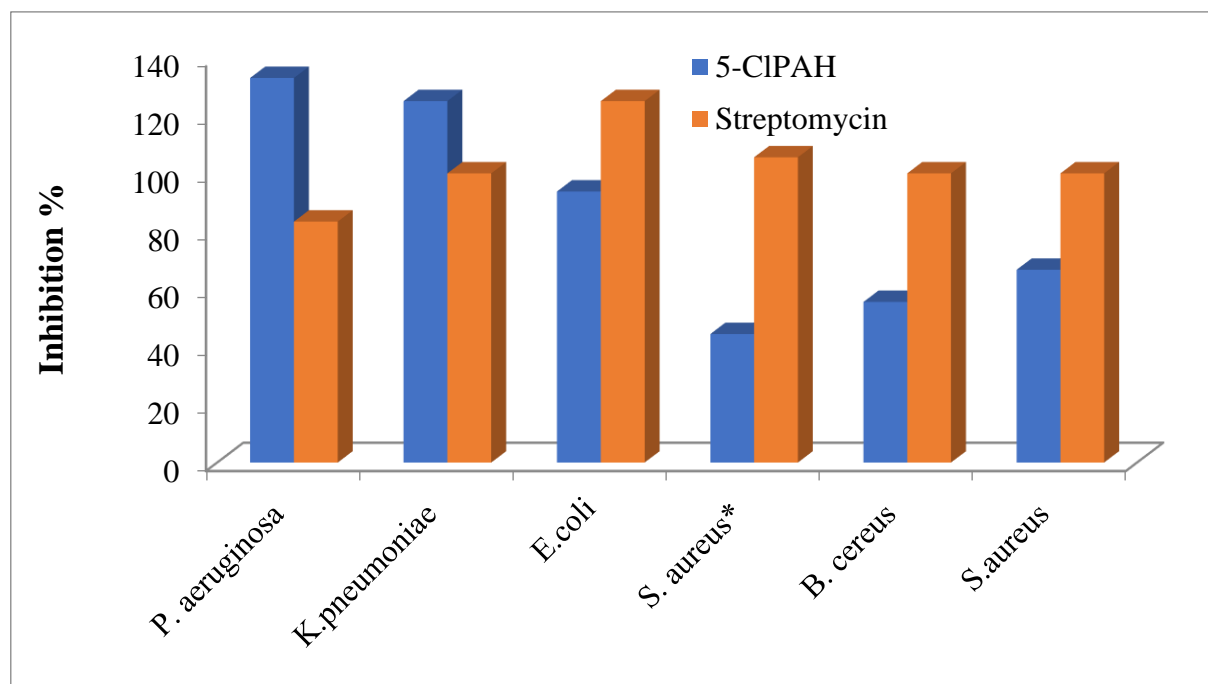


Figure 4. Percentage of inhibition of 5-CIPAH and Streptomycin against Ampisilin

The percentage of inhibition for 5-CIPAH molecule and standard drug Streptomycin showed in Figure 4. The 5-CIPAH molecule and Streptomycin show good activity against Gram-negative *K. pneumoniae* (125% 100%), respectively (Ampisilin was accepted 100% inhibition).

The 5-CIPAH molecule showed excellent inhibiting activity against bacterial strains with MIC value less than that of standard drugs Ampisilin and Streptomycin. The 5-CIPAH molecule was excellent inhibition (MIC=78.12 $\mu\text{g/mL}$) against *S. aureus* ATCC 25923 and *S. aureus* ATCC 29213.

Table 2. The MICs ($\mu\text{g/mL}$), (mM) of antibacterial activity of 5-CIPAH and standard drugs

Bacteria strain	MICs $\mu\text{g/mL}$ (mM)		
	5-CIPAH	Ampisilin	Streptomycin
Gram- negative			
<i>P. aeruginosa</i> ATCC 27853	156.25 (0.308)	93.75 (0.268)	93.75 (0.161)
<i>E. coli</i> ATCC 25922	156.25 (0.308)	93.75 (0.268)	46.87 (0.0806)
<i>K.pneumoniae</i> ATCC 13883	156.25 (0.308)	93.75 (0.268)	93.75 (0.161)
Gram-positive			
<i>S. aureus</i> ATCC 29213	78.125 (0.154)	46.87 (0.134)	46.87 (0.0806)
<i>B. cereus</i> Roma 709	156.25 (0.308)	93,75 (0.268)	93.75 (0.161).
<i>S. aureus</i> ATCC 25923	78.125 (0.154)	46,87 (0.134)	46.87 (0.0806)

The antifungal stud of the 5-CIPAH molecule was carried out *Candida tropicalis* M002, *Candida parapsilosis* M006 and *Candida albicans* ATCC 80018 fungal strains (Table 3) (Rahman et al., 2001). The antifungal activity results of Schiff base and standard drugs; Ampisilin and Streptomycin were exhibited in Table 3 (Figure 5). From the antifungal activity results, 5-CIPAH exhibited excellent inhibition against *Candida albicans*, *Candida parapsilosis* and *Candida tropicalis* with disck 10-22 mm than the standard drugs Ampisilin (7-8 mm), Streptomycin (10-12mm). As seen in Figure 5, The 5-CIPAH showed excellent activity against *C.albicans* (214.28 %) *C. parapsilosis* (314.28%) and *C. glabrata* (125%).

Table 3. Inhibition zone diameter (mm) of 5-CIPAH and standard drugs

Compounds	Zone of Inhibition (mm)		
	<i>Candida tropicalis</i> M002	<i>Candida parapsilosis</i> M006	<i>Candida albicans</i> ATCC 80018
5-CIPAH	10	22	15
Streptomycin	12	11	10
Ampisilin	8	7	7

The HOMOs and LUMOs which are responsible for chemical reactivities are known as the FMO. HOMO is the highest energy filled molecular orbital and LUMO is the lowest energy vacant molecular orbital. HOMO represents the desire to donate an electron while LUMO represents the desire to gain an electron (Fleming, 2010; Benabid et al., 2020). HOMO and LUMO have been carefully evaluated to understand the ligand reaction and to specify the reactive sites. These orbital energy levels are used to determine intermolecular charge transfers, and to calculate Global reactivity descriptors such as electron affinity, electronegativity, electrophilicity index, potential chemical reactivity and ionization (Ayers & Parr, 2008; Sultan et al., 2016; Oueslati et al., 2019). The energies of HOMO and LUMO orbitals of **5-CIPAH** molecule are -8,01eV and -5.52 eV, respectively. The HOMO and LUMO orbital energy gap ($\Delta E: E_{\text{HOMO}}-E_{\text{LUMO}}$) calculated using B3LYP/6311G method is 2,49 eV (Table 4, Figure 6). The values of global softness (S), ionisation potential (IP=-HOMO), electronegativity (χ), electron affinity (EA=-LUMO), chemical hardness (η), and electrophilicity index (ω) for **5-CIPAH** molecule at B3LYP/6311 basis are given in Table 4.

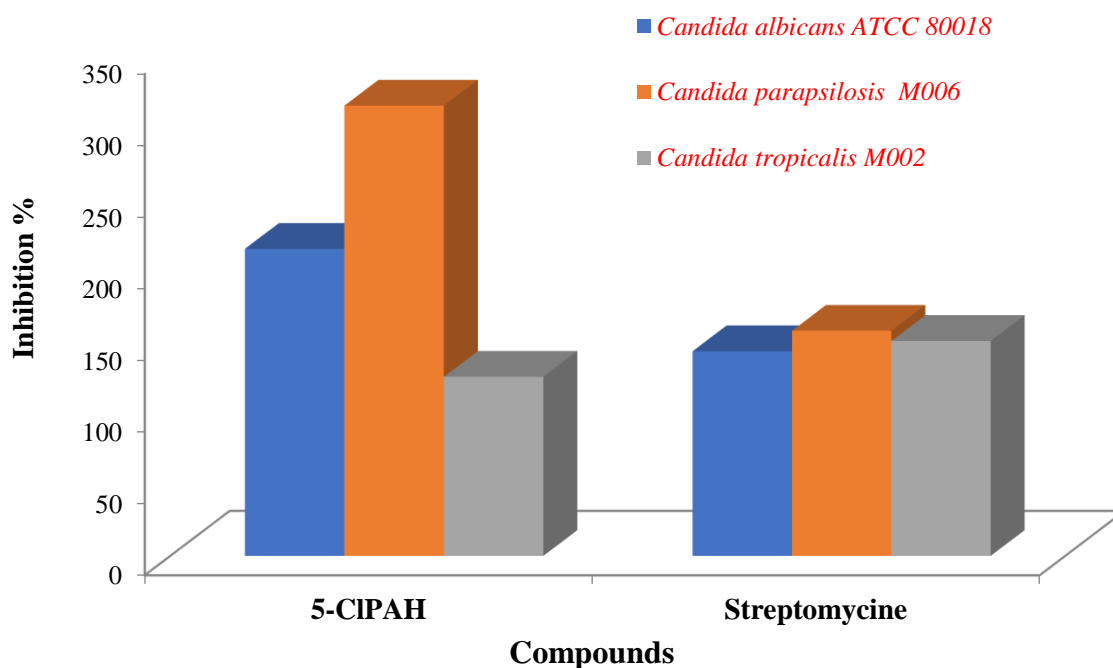


Figure 5. Percentage of inhibition of 5-CIPAH and Streptomycin against Ampisilin

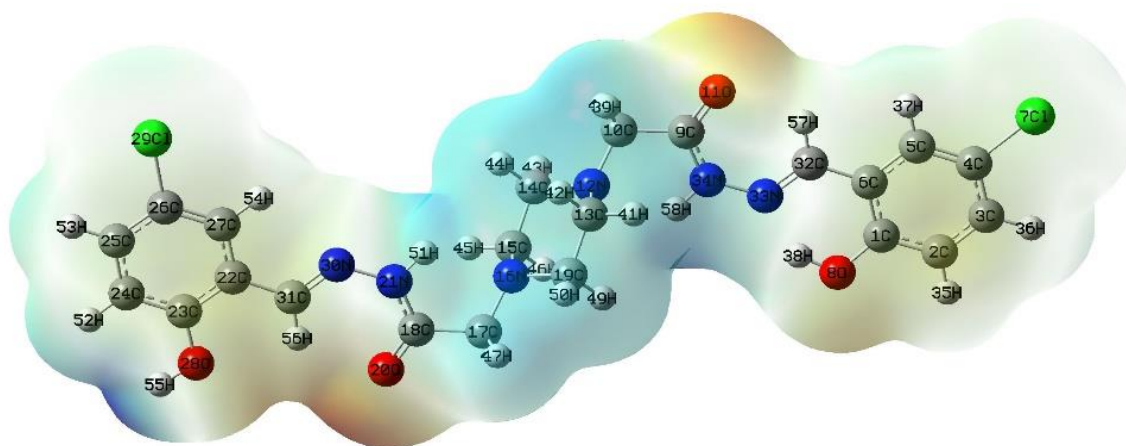


Figure 7. Derived MEP surface analysis representation of 5-CIPAH molecule

4. CONCLUSION

In this study, 5-CIPAH molecule with the chemical formula $C_{22}H_{24}Cl_2N_6O_4$ was synthesized and characterized. DFT based calculations were used to theoretically analyze the 3D optimized structure of 5-CIPAH molecule. The HOMO and LUMO energies were used to predict quantum chemical properties like hardness, electrophilicity and softness. Stable structure of 5-CIPAH molecule was obtained by PES analysis. The HOMO energy gap value for 5-CIPAH molecule was found to be -8.01 eV at B3LYP/6-311 G level of theory. In vitro biological tests show that 5-CIPAH molecule have moderate efficacy as antibacterial and antifungal reagents.

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CONFLICT OF INTEREST

The author declares no conflict of interest.

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