Solvent role in molecular structure level (TD-DFT), topology, and molecular docking studies on liquid 2', 4'-dicholoroacetophenone

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Abstract

The study investigated the structure, spectroscopy, and chemical reactivity of 2',4'-dichloroacetophenone using DFT and Hartree-Fock (HF) methods. The B3LYP and LC DFT (CAM-B3LYP) methods were applied to a 6-311++G (d, p) basis set combination. The bond angles and bond lengths are appraised in depth to understand the geometrical framework. The FT-IR and FT-Raman spectra of the respective compound were analyzed within the regions of 4000-400 cm⁻¹ respectively. The experimental spectra were compared and discussed in connection with the theoretical spectrum. The VEDA program was implemented to accomplish the vibrational assignments of all the fundamentals highlighted using the PED analysis. TD-DFT level was used to explore the essential electronic properties of LUMO and HOMO energies. To forecast reactive sites, the MEP surface was examined. NBO was computed according to the compound's reactivity and biological requirements. The Multi-Wave Function Analyzer 3.6 is designed to examine and compare topological inquiries, specifically focusing on the LOL and ELF. The detailed investigation specifies the drug likeness and molecular docking that assisted in the prediction of the physicochemical, pharmacokinetic, and biological availability of 2', 4'-dichloroacetophenone.

Keywords: Vibrational assignments; Drug-likeness; NCI; Molecular docking, 2', 4'-dichloroacetophenone.

1.0 Introduction

Acetophenones are the most prevalent type of aromatic carbonyl and exhibit intriguing physical, chemical, and biological properties (Balan et al., 2009). Natural sources of them include chicory, beef, bananas, apples, apricots, spearmint, watermelon, cauliflower, and cheese. They can also be acquired

using various synthesis techniques (Subramanian et al., 2008). Nevertheless, the acetophenone structure is used in the pharmaceutical and related industries. It's also worth noting that amino acetophenone derivatives have some level of local anesthetic activity (Huang et al., 2010). An orange-scented, colorless liquid known as acetophenone has a sweet and pungent flavor. Skin and eye irritation may result from brief, acute exposure to the acetophenone vapor. Twenty acetophenones were studied, and electronic and topological indices were used to correlate the compounds' antibacterial activity with their structural properties (GRIFFIN, 1968). Furthermore, diazonium salts with a dihydroxy acetophenone structure have anti-microbial activity (Sudarshan et al., 2024). Aside from that, substances between p-substituted acetophenone and benzoyl hydrazone have antifungal properties (Sivakumar et al., 2008). In contrast, substituted acetophenones are promoted as reagents for synthesis in certain organic processes. Acetophenone is one of the most basic aromatic ketones (Sudarshan et al., 2015). Numerous beverages and sweets with cherry flavors frequently contain it as a flavoring ingredient. Numerous plant species have been found to contain acetophenone, which is also a volatile component of honey. It has a role as a photosensitizing agent, an animal metabolite, and a xenobiotic. It is utilized in the field of medicine, and it was marketed as a hypnotic and anticonvulsant under the brand name Hypnone. The chemical formula [C8H6Cl2O] describes the structure of 2', 4'- dichloroacetophenone (24DCA) derived from benzene. It is used in the preparation of Schiff base derivatives. Chloroacetophenone is a chemical mace and is primarily used as a riot-control agent (tear gas) (Vinod P. Singh and Katiyar, 2009). It is also used in pharmaceuticals and was previously used as an alcohol denaturant. Based on a literature review; it has come to our attention that no researchers have tried any quantum chemical approaches for this molecule. A comprehensive structural and spectroscopic profile of the 24DCA molecule is the primary emphasis of the present exploration. Molecular geometrical parameters, spectroscopic techniques, reactivity descriptor investigations, and Mulliken, were performed by HF/B3LYP/CAMB3LYP techniques invoking 6-311++G (d, p) level. Computational studies were also assessed and the theoretical and experimental results were compared (Ramanan et al., 2016). Chemical stability, band gap energy, and biological presumption were studied for gas, ethanol, methanol, and dimethyl sulfoxide using HOMO-LUMO and UV-visible inquiry.

To determine the reactive sites of the molecule, the molecular electrostatic potential (MEP) surface of the molecule was activated. The NBO analysis successfully demonstrated the hyperconjugation of different intra-molecular interactions in the molecule. To further enhance the topological examination of interatomic interactions within the ligand-enzyme pocket, non-covalent interactions (NCI) have been conducted. Furthermore, molecular docking and ADMET drug-likeness will aid in the justification of intramolecular drug-receptor interactions and provide a pathway for designing new molecules. The investigation also encompassed an examination of the process and biological activity of the 24DCA compound about its antifungal (IPCV), antioxidant (2V32), insulin inhibitor (IBCV), antiseptic (5ZEC), and antiviral (6UEI) attributes. A molecular docking study aimed at incorporating relevant proteins collected from the RCSB-protein data bank site.

2.0 Experimental details

A liquid sample of 2', 4'-dichloroacetophenone with a purity level of over 98-99% was purchased from Sigma-Aldrich Chemical Company. The sample was used to record FT-IR and FT-Raman spectra, without undergoing any further purification. An Alpha II Bruker FT-IR spectrometer was used to capture FT-IR spectra in the wave number range of 4000-400 cm⁻¹. The spectra were obtained utilizing a high-sensitivity, temperature-controlled DLaTGs detector, and the KBr pellet method. The resolution of the spectra was ± 2.0 cm⁻¹. The FT Raman spectrum was obtained using the BRUKER RFS 100/S apparatus, which utilizes a Nd: YAG laser source operating at a constant power of 150MW and an excitation wavelength of 1064 nm. The spectrum was recorded at room temperature

to measure Raman scattering. The spectrum measurements included a range of 4000-400 cm⁻¹, and the reported wave number is expected to have an accuracy within 1 cm⁻¹.

3.0 Details of computations

The Gaussian 09W software package was adopted to accomplish structural optimizations, energy, and vibrational frequency computations on the aforementioned compound leveraging the HF/B3LYP/CAM-B3LYP/6-311++G (d, p) basis set to characterize the molecular framework, electronic, and other properties (Frisch et al., 2009). Utilizing VEDA 4 software potential energy distribution calculations were used to interpret the theoretical vibrational wave number. Gauss View 5.0 was employed by scrutinizes like FMO (Frontier molecular orbital), NBO (Natural bond orbital), & MEP (Molecular electrostatic potential), and a variety of polar solvents, such as CH3OH (protic), ethanol, and DMSO (aprotic), deployed in UV–vis absorption measurements (N. Elangovan et al., 2024a). The topological traits of LOL (Localized orbital locator) and ELF (Electron localization function) were done with the aid of a Multiwfn software package and plotted by a visual molecular dynamics program (Lu and Chen, 2012). To ascertain the characteristics of drug-likeness, the Swiss ADME package was used. Furthermore, the Auto Dock 4.2 program was implemented to explore the biological roles of the molecule (Goodsell et al., 2021). The molecular graphics software PYMOL and Discovery Studio have been deployed to portray the protein-ligand combination.

4.0 Results and Discussion

4.1 Structural properties

The optimization of geometrical parameters are notably indispensable step in quantum chemical calculations. The structure after optimizing 2', 4'-dichloroacetophenone in addition to the atom numbering is portrayed in Fig. S1. The HF/B3LYP/CAM-B3LYP /6-311++G (d, p) methods were deployed to optimize the system. The optimized geometric parameters for the named chemical are shown

in Table S1, along with their correlation to the experimental XRD data (Demircioğlu et al., 2015). The studied compound includes two 'Cl' and one 'C=O' group with a benzene ring, along with the theoretical value we can figure out that the measured value of the bond length is marginally lower than the optimized value. The reason for the inclusion of isolated molecule theoretical computations in the gaseous phase is noteworthy. Simultaneously, the experimental outcome is a constituent of the solid-state molecule. The vibrational frequencies were measured primarily based on calculated geometrical parameters and it'll be a good approximation for the calculation of the above parameters. As an effect of the electronegative substituent on the benzene ring little distortion seems within the benzene ring and the angle is barely out of perfect hexagonal structures. Evaluating bond angles and bond length reckoned leveraging the B3LYP is better than HF and well correlates with the experimental value (Elkaeed et al., 2022). As a result of the hydrogen atoms being substituted with chlorine, oxygen, and CH3 groups, the phenyl rings are marginally deformed and deviate slightly from their hexagonal shape. For the benzene ring, the reckoned CC bond spacing ranges from 1.405Å to 1.379. The HF approach yielded the smallest CC bond length of 1.379 Å, whereas the prolonged CC bond length measured 1.514 Å by DFT. In the most extreme cases, they differ by 0.1Å within the same method of calculation. The C-Cl bond lengths vary between 1.760Å to 1.739Å, and when the observed values exhibit a strong concurrence with the values measured 1.736 Å & 1.732 Å (Abozeed et al., 2022). The estimation of this bond length is observed to be overestimated in the DFT approaches, while it is diminished in the HF method.

The C-H bond lengths of aromatic benzene rings and methyl groups were reckoned in the present work ranging from 1.072Å to 1.093 Å. However, the experimental values had been stated as 1.082, 0.930, 1.090Å, and 1.000, 0.860, and 0.910Å respectively. The C=O bonds have respective lengths of 1.189, 1.215, and 1.209 Å. The arrangement of bond angles makes it evident that the phenyl ring is asymmetrical. C2-C1-C12 < C1-C6-C5 < C1-C12-C14 < C3-C4-C5 < C1-C2-C3-C2-C3-C4. As a result of the substitution of Cl and CH₃ At 117.5,121.6,121.1° for HF 117.0,121.7, 121.1° for B3LYP, and

117.3,121.5,121.0° for CAM- B3LYP. The calculated bond angles that differ from experimental values are 116.7, 122.3, and 120.5°.

4.2 Vibrational spectral analysis

In organic chemistry, vibrational spectroscopy is frequently used to analyze to identify functional groups in organic compounds, as well as to investigate molecular confirmations, kinetics, and reaction mechanisms. The 2', 4'-dichloroacetophenone is composed of 17 atoms and has the molecular formula C8H6Cl2O. It has 45 normal vibrational modes, including stretching, bending, torsion, and a few mixed vibrations, and it is symmetrical in the C_s point group, classified into 29A' and 16A" species (Jeyavijayan, 2015a). The in-plane and out-of-plane vibrations are denoted by the A' and A" species, respectively. The reckoned frequencies, PED, and complete vibration assignments of 24DCA are documented in Table 2. The present research effort primarily aims to figure out the correct experimental wave number by comparing it to theoretically scaled wave numbers with PED adopting the aforementioned methodologies, along with the aid of normal mode studies (N. Elangovan et al., 2021b). According to the findings, the DFT values tend to exhibit less variation than HF values. Figs. 1 & 2 exhibit the correlation graph of the reported and predicted FT-IR and FT-Raman vibrational frequencies providing visual comparison.

4.2.1 Carbon - Hydrogen vibration

In heteroaromatic substances, C-H stretching is often detected in the $3100\text{-}3000\,\text{cm}^{-1}$ wavelength range. This region is renowned for its rapidity in detecting such vibrations. The nature of the substituent group interaction has an impact on the region that can be used to identify the C-H stretching vibration (Sundaraganesan et al., 2006). The Trifluoromethyl compound was studied by Balachandran et al., who observed the v(C-H) modes, in both spectra, with bands spanning $3075/3047~\&~3090/3020~\text{cm}^{-1}$. Within the research, the bands are seen in the recorded IR & Raman bands located 3087/~3004~&~3073/~3007

cm⁻¹, While stretching modes were envisioned at 3090, 3077, 3010, 3088, 3075, 3005, 3086, 3074, and 3006 cm⁻¹, respectively. The bands appeared with medium and weak intensities except for one band in FT-Raman and the expected region and contributing 98 % of PED. The vibration mode numbers of 1, 2, and 3 is also supported by experimental data and aforementioned works of literature. The bands arising from δ(C-H) mode linked to υ(C-C) mode are seen as an array of spectra in the range 1465-1281 cm⁻¹. In this compound, the above vibrations are seen in the region at 1465, 1372, 1235 cm-1 (IR), and 1375, 1237 cm⁻¹(Raman), respectively. The corresponding computed values, which show good correlation with experimental data, reckoned theoretically at 1468, 1378, 1240 1465, 1374, 1237, 1466, 1375, 1242 cm⁻¹ (Mode nos: 10, 13, and 16). The aromatic C-H out-of-plane bending mode appears in the range of 900-667 cm⁻¹, and it is calculated theoretically, can be seen in the following configuration: 750, 635, 515 751, 630, 510,752, 638, 512cm⁻¹. Similarly, the experimental peaks at 511 and 632 cm⁻¹ are correspondingly identified. All calculated wave numbers are well consistent with those obtained through experimentation and are figured in Table 1.

4.2.2 Methyl group vibrations

The vibrations of methyl groups are often associated with substituents that contribute electrons in aromatic chains. The inclusion of a CH₃ group in a compound entails the generation of one symmetric stretching vibration and two asymmetric stretching vibrations, and usually, asymmetric stretching vibrations are recorded at higher frequencies than those of the symmetric mode. In general, CH₃ exhibits asymmetric and symmetric stretching vibrations with wavenumbers of 2962 and 2872 cm⁻¹, respectively. The robust band at 2862 cm⁻¹ was attributed to the CH3 symmetric stretching vibration of 5-chloro-2-hydroxyacetophenone. Referring to the above workers in the present case weak to medium bands observed at 2850/2924cm⁻¹ (IR) and a very sturdy peak found at (Raman) at 2925cm⁻¹ shows asymmetric stretching vibrations of CH₃ whereas a faint band observed at 2833cm⁻¹ (Raman) assignable

to symmetric stretching mode. These assignments found support from the calculation followed by Gaussian showed the peaks at $v_{ass}CH_3 = 2929$, 2854, 2925, 2851, 2925 &2850 cm⁻¹ respectively, with mode nos; 4 and 5 having considerable PED values of 97 %. The υ_{ss}CH₃ is reckoned at 2839, 2834, and 2833 cm⁻¹(mode no; 6), nearly 98 % of PED impacts are confirmed, and its corresponding counterpart couldn't be detected in the FT-IR spectrum. The CH₃ deformation is expected in the region 1485–1400 cm⁻¹. The molecule in question, the medium and weak in-plane bending methyl deformation modes CH₃δ_{ipb} observed at 1422/1430cm⁻¹ in both spectra line up with the reckoned values at 1430,1425, and 1423 (mode no; 12) with PED 72 %. Thus, the band observed (calculated) at 1446 (1454, 1450, 1445 cm⁻¹) ascribed to out-of-plane bending vibration. These two bending are quite similar in the assigned values during the experiment and computations. The computational data for symmetrical methyl deformation modes CH₃sb was obtained at 1360, 1356, and 1354 cm⁻¹ with PED 72 % (mode no; 14). Theoretical calculations using the B3LYP method at 1356 cm⁻¹ are perfectly in line with measured values of 1356 cm⁻¹ in the FT-IR spectrum. The methyl rocking vibration bands are typically seen in the vicinity of 1045 and 970 cm⁻¹, respectively, in aromatic compounds. The methyl group rocking vibrations of the compound under consideration are spotted at 1040 & 1079 experimentally, agree with the theoretical computations at $\delta_{ipr} = 1047, 1042, 1045 \text{cm}^{-1} \text{(mode: no21)}$ and $\delta_{opr} = 1083, 1080, 1082 \text{ cm}^{-1} \text{(mode no: 20)}$ with PED 68 & 63% respectively.

4.2.3 Carbon-Chlorine vibrations

The C-Cl stretching mode furnishes typically sturdy bands at 765-505 cm⁻¹. Compounds comprising multiple chlorine atoms exhibit exceptionally vivid bands as a result of the asymmetric and symmetric stretching modes. Additionally, there's a possibility that vibrational interaction with other groups might intensify an absorption shift to 840 cm⁻¹. In this study, mode no:(25,26) is ascribed to v=(
C-Cl) stretching vibrations in a strong band at 817 cm⁻¹ (IR) band, and these outcomes obtained match

precisely with the experimental data of 812, 800 & 800 (Nickless et al., 2016). Experimentally, the CCl stretching mode did not exhibit a Raman spectrum, and its matched harmonic wave numbers at v(C-Cl) of 845,843,843,825, 820, 822 cm⁻¹ respectively, it has notable PED values such as 66%, 68%, respectively. In the current exploration, the weak bands reported at $\delta(C\text{-Cl}) = 533,433\text{cm}^{-1}(Raman)$, and the reckoned values lies in the range $\delta(C\text{-Cl})$ of 540-436 cm⁻¹, (mode nos: 32 & 34) respectively. With adequate PED and substantiated by experimental data of 429 cm⁻¹(IR) and their related harmonic frequencies are; 434-400 cm⁻¹ with a PED contribution of 58% showed the $\gamma(C\text{-Cl})$ vibrations.

4.2.4 Carbon=Oxygen vibrations

The frequencies of carbonyl stretching exhibit a strong sensitivity to stimuli that influence the characteristics of the carbonyl group. Their specific frequency is distinctive to the particular type of carbonyl molecule that is under scrutiny. The frequency of carbonyl bond stretching has been the subject of particularly through correlations. Conjugation may enhance the strength of these bands, offering assistance in both the detection of Raman lines and the rising infrared band intensities. Since the aforementioned group is extremely polar (C+=O-), It typically generates a prominent infrared absorption spectrum. Typically, the carbonyl C=O absorption contributes significantly to the spectra which are most prominent between 1800 and 1690 cm⁻¹. A very prominent infrared band linked to the υ(C=O) vibration of 2', 4' dibromo acetophenone, was observed at 1692cm⁻¹ in IR and the calculated value is 1693cm⁻¹ an intense peak observed at υ(C=O) at 1695 cm⁻¹(IR) and 1693 cm⁻¹ (Raman) a theoretically predicted frequency at 1694cm⁻¹ is assigned to this mode, respectively (Jeyavijayan, 2015b). In the current study, it has been demonstrated that the (C=O) stretching frequencies display notable bands in 24DCA. These bands appear to have a significant impact on the projected frequencies of mode number 7, specifically at 1696, 1695, and 1698 cm⁻¹, respectively. The IR spectra of the compounds exhibited an intense absorption peak measured at 1697&1693cm⁻¹ (Raman) as a medium band, being over 72% of the total,

as can be seen in the PED column. The observed picture in this $\upsilon(C=O)$ spectral range is quite similar to those predicted in the cited literature. The FT-Raman spectrum of a molecule showed the C=O group's in-plane and out-of-plane bending vibrational modes at 1106 cm-1 and 900 cm-1 respectively. A tight agreement is seen between the reckoned value and the measured frequencies. This C=O molecule was unaffected by other vibrational modes, as the PED contribution of the vibration is 66 and 68 correspondingly.

4.2.5 C-C vibrations

The aromatic ring's ring stretching vibrations are quite distinctive, so they play an important part in the benzene and derivative spectrum. The substitution of aromatic rings in benzene derivatives has a consequence on the many ring vibrations. The six carbon atoms in each ring experience coupled vibrations known as skeletal vibrations. Ring stretching vibrations, notably C-C and C=C vibrations, which are alternatively referred to as semi-circle stretching vibrations, generate distinct bands in the infrared and Raman spectra, covering the spectrum of 1430-1650cm⁻¹. Substitution of the aromatic ring changes most of the ring modes. The specific location of these modes is more influenced by the substitution pattern around the ring system rather than the nature of the substituent. The frequency is undoubtedly diminished by the heavy halogens. (Bhat and Ahmad, 2016). The band with different intensities explored here was found to match with 1584, 1552, 1268,1139, 1094, 962, 869, and 679cm ¹/FT-IR, and the same type of bands were noted at 1585, 1270, 1144, 963 and 682cm⁻¹/FT-Raman spectrum and they were appointed to CC stretching vibrations. Theoretical calculations indicate that the C-C stretching vibrations account for 75%, 75%, 70% 65% 71% 70%, and 68% 59%, respectively. The spectral bands at 1590, 1588, 1584, 1555, 1552, 1550, 1275, 1270, 1269, 1146, 1145, 1142, 1099, 1098, 1095,965, 963, 962, 874, 872, 686, 685, 682 cm⁻¹ are reported. The observed data and all calculated frequencies line up perfectly. The benzene ring's out-of-plane and in-plane bending vibrations are noticed below 1002 cm⁻¹. In this instant study, the ring out-of-plane and in-plane and bending modes are affected to a great extent by the substituents and produce medium to weak bands at 596/575cm⁻¹ (IR) and weak bands at 600/ 567 cm⁻¹ (Raman) ascribe to C-C in-plane bending and corresponding theoretically fitted band at 604-574cm⁻¹ for 24DCA. The scaled theoretical wave number corresponding to in-plane bending vibrations goes well with their available Experimental observations. Additionally, 15 C-C out-of-plane bending modes computed at 327, 322, 320, 250, 245, and 241 cm⁻¹ with PED contribution accounting for 53% & 55%. The (δ_{ring}) and (β_{ring}) bending modes in the titled molecule are greatly affected by the substituent and theoretically produce bands at 298-45 cm⁻¹, not been found experimentally.

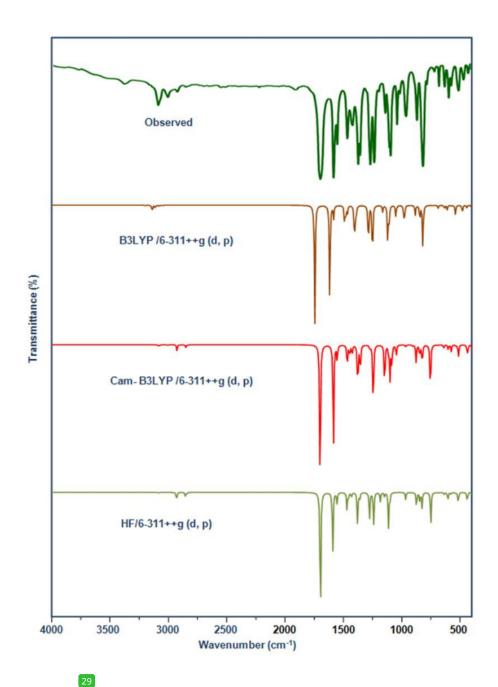


Fig. 1 Observed and Calculated FTIR spectrum of 24DCA

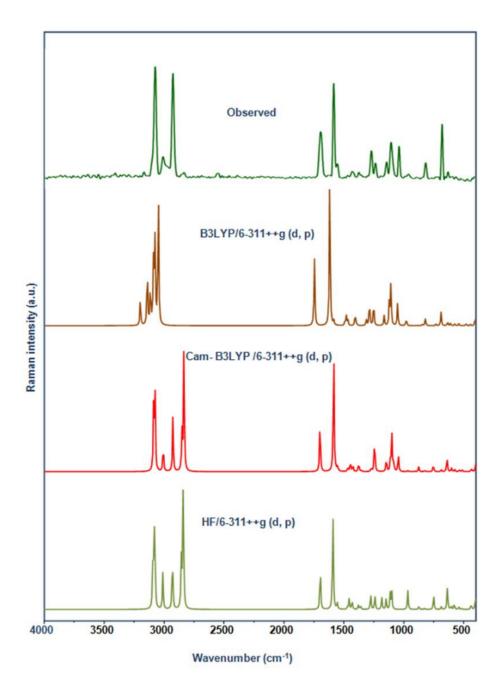


Fig. 2 Observed and Calculated FT-Raman spectrum of 24DCA

4.3 Frontier molecular orbital analysis

The energy of the atomic orbital and atomic contribution coefficients constitute significant in articulating the reactivity of molecular properties. The orbitals have an enormous effect on the overall molecular behavior. Molecular orbital offers essential perspectives into the bonding and other chemical properties of molecules. HOMO serves as the orbital responsible for supplying electrons, whereas LUMO, which occupies a lower energy level, operates as the orbital responsible for receiving electrons termed as Frontier molecular orbital. It has the potential to function as a as a straightforward indicator of kinetic stability. Both possess the ability to absorb and release electrons. These specific orbitals allow the molecule to have its strongest interactions with itself (Diya et al., 2023). The ligand s' optimized structure was employed to figure out the electron cloud shapes and electronic transition energies of the substance, the graphical representation of the FMOs depicts the outcomes of the calculation done in this work on gas, Dimethyl sulfoxide, ethanol, and methanol solvents respectively, also depicted in Fig. 3. The green and red regions respectively represent negative and positive charges. The primary focus of the current investigation on the 24DCA compound is on the HOMO charge distribution, with particular emphasis on the chlorine & hydrogen atoms throughout the whole ring system. By contrast the dislocation of the LUMO- π nature, specifically the benzene ring, across all C-C bonds, is symbolized by the HOMO

LUMO transition. This transition proposes an electron distribution movement from the chlorine atom to the C-C bond of the benzene ring and the CO-CH₃ group. The determination of the compound's transition energy from HOMO→LUMO with the shortest to the highest energy gap: 4.458 eV (ethanol), 4.461 (methanol), 4.464 (DMSO), 4.467 (water), 4.546 (Gas). As the header compound's gas phase had a wide energy gap, we may infer that the molecule was more stable than compared to other solvents with chemical harness 2.273. Water was discovered to have the greatest chemical hardness (2.233) among all solvents. The header molecule has a softness index of 0.439 (Gas), 0.447 (methanol), 0.430 (DMSO), 0.447 (water), and 0.449 (ethanol) which means that the molecule under exploration possessed non-toxic traits. The 24DCA complex exhibits higher Electronegativity (χ) values across all phases 5.212 for gas, 5.182 for water, 5.179 for DMSO, 5.174 for ethanol, and 5.177 for methanol, indicating that the screened molecules function as Lewis's acids (Elangovan and Sowrirajan, 2021). Having higher electron-accepting indices (ω +) in the water phase seems to be linked to a better ability to handle electron density. Moreover, the ability to donate electrons (ω -) in the water phase was higher (8.882) compared to other solvent phases. The reckoned global reactivity parameters are documented in Table S2.

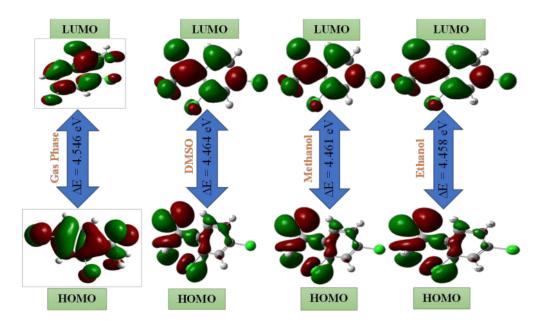


Fig. 3 FMO surface map of 24DCA

4.4 UV spectral analysis

U-V visible spectra were examined in different solvents to study solvent effects on electronic transitions. The most precise theoretical approach to evaluating the dynamic and static properties of excited-state molecules is the Time-dependent density-functional theory (TD-DFT) method. For the compounds under study, the TD-DFT method was used to determine the peak absorption wavelength (λ_{max}) , excitation energies (E), oscillator strength (f), energy gap (eV), and molecular orbital

contribution. The spectral data of the selected solvents are being studied by UV-visible spectra, to assess the impact of solvents in transitions (N Elangovan et al., 2024). The correlated spectrum and data are explored in Fig. 4 and Table 2. The fact that the projected maximum values for different solvents are practically equal suggests that the solvent influences the compound's optical activity. The maximum absorption peak of the experimental UV-visible absorption in ethanol was measured at 282 nm.

Theoretical spectra have been performed in the gas phase and several polar solvents, such as DMSO (aprotic), C₂H₆O, and CH₃OH (protic). The highest possible absorption values are 288.17nm and 4.3019 eV with an oscillator strength of 0.052 in the gas phase, 302.50 nm and 4.0981eV in ethanol, 308.28 nm and 4.0213eV in methanol & DMSO, all with the same oscillator strength of 0.076. Moreover, the highest transition efficiency observed for HOMO-3>LUMO was 78% across all solvents (Rajimon et al., 2023). These outcomes and predictions lend evidence for selecting the molecule as a highly energetic and stable chemical for biological purposes.

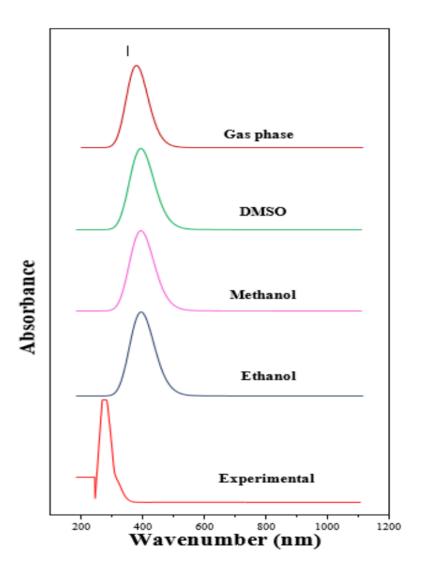


Fig. 4 Experimental and Theoretical UV- Vis spectral analysis of 24DCA
4.4 Molecular electrostatic potential

MEP investigation is an efficacious descriptor to forecast the nucleophilic and electrophilic attack for 2', 4'-dichloroacetophenone by the aforementioned approaches to scrutinize the chemical reactivity of the molecules, to explore the correlations among a chemical configuration along with its physiochemical characteristics, MEP projection is extremely helpful (Elangovan et al., 2022). The role

of molecular electrostatic potential surfaces in computer-aided drug design is noteworthy, by aiding in the optimization of the ligand-protein electrostatic interactions. Fig. 5 illustrates, a graphical depiction of the compound under investigation. With the use of color coding, this surface provides reaction point knowledge. Distinct color tones have denoted the surface electrostatic potential values of the MEP map. The strongest attraction is betrayed by blue color and red bespeaks strongest repulsion. Conventionally, negative potential sites are linked to the lone pair of electronegative atoms. The color scheme of the map in our compound is in the vicinity of 4.541e⁻² (deepest red) to 4.541e⁻² (deepest blue). The majority of the negative potential sites are centered on the electronegative oxygen atoms of the CO-CH3 group, according to the MEP sketch of the compound in issue (Priya et al., 2023b). Electrophilic attack sites are possible in the darkest red zone exposed around the O atom. In contrast, the H atoms inside the benzene rings are covered by positive potential. The blue color examines the available sites for nucleophilic attack on the benzene ring. The leftover spices are embraced by zero potential, which is highlighted in green color.

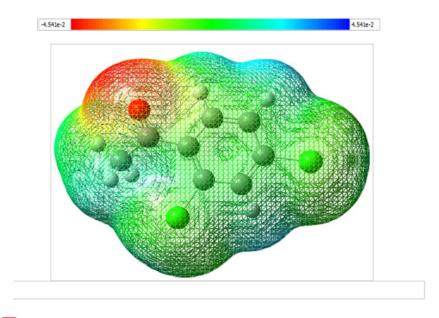


Fig. 5 Calculated 3D Molecular electrostatic potential contour map of 24DCA

4.5 Mulliken atomic charges

The charge distribution of the chemical in question is depicted in Table S3. Mulliken atomic charge distribution of 24DCA, computed by the above-mentioned methodologies is portrayed in Fig. S2. The Mulliken population provides information regarding Mulliken charges and makes it possible to assess partial atomic charges from the calculation accomplished by the computational chemistry method. The title compound under investigation has a total of 17 atoms in its chemical structure. Eight carbon atoms are present together with six hydrogen atoms, two chlorine atoms, and one oxygen atom. Carbon atoms together with negative values are dominant in C2, C3, C5, C12, and C14 owing to hydrogen and oxygen coupling. The residual carbon atoms C1, C4, and C6 are positive as a result of the re-distribution prompted by the attachment of CO-CH₃ and chlorine groups (N Elangovan et al., 2021). The utmost negative charge values of about -1.682e in the C2 atom and approximately 0.802e is the greatest positive charge of C1 in the molecule at the HF level. According to Mulliken's atomic charges, all of the hydrogen atoms connected with benzene rings have net positive charges, At the contrary, due to their proximity to the chlorine atom, H10 atoms possess a higher positive charge than the remaining hydrogen atoms. With an atomic charge value of 0.287e (HF). All electronegative (oxygen atoms maintain a negative charge), all electropositive (hydrogen atoms maintain a positive charge), and depending on position, carbon atoms can maintain either a negative (or positive) charge.

4.6 Donor-Acceptor interaction

Understanding inter- and intramolecular bonding interactions is critical to exploring charge transfer or hyperconjugation interactions inside a molecular system. The investigation of the Natural Bond Orbital (NBO) indicates strong charge-transfer interactions in the isolated gas-phase molecule of 24DCA. In this work, the investigation focused on a variety of donor-acceptor bond interactions, outlined in Table 3. Higher E (2) values propose stronger interactions between electron providers and

electron recipients, implying a greater degree of electron coupling delocalization across the system. The table lists the stabilization interaction energies of 2', 4'-dichloroacetophenone above 5 kcal/mol. The natural bond orbital computations were carried, out at the B3LYP6-311++G (d, p) level (Geethapriya et al., 2023). Regarding the head molecule, The BD*(2) electron's delocalization from distributed antibonding to anti-bonding demonstrates high stabilization Energy is 157.48 kcal/mol for BD*(2) C1-C2→ BD*(2) C5-C6 which could be a cause of the title molecule's biologically active behavior (Elangovan and Arumugam, 2024). A higher stabilization energy value is also achieved for the interaction between $BD^*(2) C3-C4 \rightarrow BD^*(2) C5-C6=106.72 \text{ kcal/mol}, BD^*(2) C12-O13 \rightarrow BD^*(2) C1-C2=48.27 \text{ kcal/mol},$ BD (2) C5-C6 \rightarrow BD*(2) C3-C4=23.27 kcal/mol and BD (2) C5-C6 \rightarrow BD*(2) C1-C2=21.11 kcal/mol. Besides that, the stabilization interaction occurs between lone pair LP (2) of O13→ BD*(1) C12-C14 and LP (2) of O13 →BD*(1) C1-C12, it is the order of 8.65 and 9.87kcal/mol respectively. Meanwhile, the stabilization interaction energy value of the corresponding interaction between the lone pair of chlorine, LP (3) Cl9, and the BD*(2) C3-C4 orbital was calculated to be 13.36 kcal/mol, and between LP (3) Cl (11) and the BD*(1) C14-H16 orbital to produce a stabilization interaction amounting to 18.06 kcal/mol and these values are spotted in Table 5. These interactions cause the lone pair orbital occupancy to significantly decrease compared to occupancy, and there is a chance of hyperconjugation between chlorine, oxygen atoms, and the aromatic ring.

4.7 Non-covalent interaction (NCI)

Investigated reduced density gradient (RDG), a method that provides an approachable graphical representation of different forms of non-covalent interactions in three dimensions. Weak interactions known as non-covalent interactions are key, arising from the quantum electron density and its first derivatives in real space (Mallika et al., 2024). A basic dimensionless variable, the electron density value

in a decreased density gradient may be used to ascertain the intensity of an interaction; and it is transcribed by

$$RDG(r) = \frac{1}{2(3\pi^2)^{\frac{1}{2}}} \frac{\left|\nabla \rho(r)\right|}{\rho(r)^{\frac{4}{2}}}$$

Where the electron density has been betokened by the symbol $\rho(r)$, and the norm of the electron density vector is bespeak by $|\nabla \rho(r)|$. The highest value of the lessian matrix of electron density, sign (λ 2), aids in determining the type of interaction by color coding. The λ sign was deployed to differentiate between bonded (λ <0) and non-bonded (λ >0) interactions. Fig. 6 displays the predicted RDG plot for the aforementioned molecule, Through the use of multiwfn and the visual molecular dynamics (VMD) program (S. Sowrirajan N. Elangovan and Manoj, 2022). Areas of weak attractiveness are rendered in green on this graph, (λ <0) like van der Waals interactions, while the blue regions correspond to strongly attractive interactions like H bonds and C-Cl bonds. The steric effect is depicted by the color red. The left side of the figure's negative sign λ p values (green in color) shows attractive interactions with the other molecule. While the right side of the figure's (red) positive sign λ p-value indicates a strong repulsive interaction. Fig. 8 clearly demonstrate that the isosurface of RDG for the header compound is more affected by strong van der wall interactions when it is close to the methyl group, which is indicated by a greenish-brown isosurface, which denotes a low electron density in the area. A pronounced steric effect is manifested in red fusiform areas that existed in the middle zones of the phenyl ring, encircling the C=O bond, and within a ring-like configuration connecting Cl9 and Cl11.

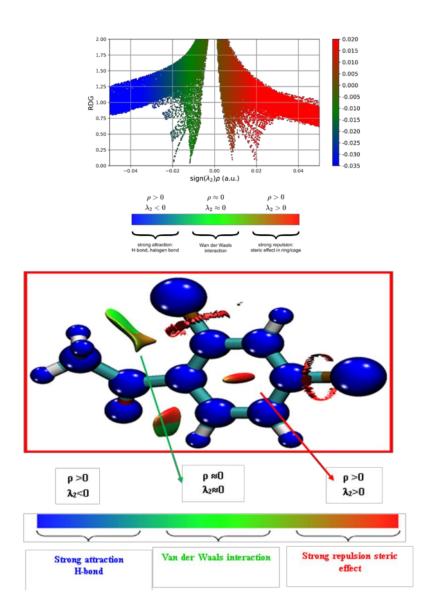


Fig. 6 NCI analysis of 24DCA

4.7.1 ELF and LOL Study

In chemical bonding, the electron localization function ELF, symbolized by $\tau(r)$, provides a density-based description, and the localized orbital locator symbolized by $\eta(r)$ are studies that primarily express the surface of a topological analysis, carried out using the multifrequency wave function

(multiwfn) program. Their color shade maps are highlighted in Fig.S3 and are used to interpret the surface assessment that is dependent on the interactions between covalent bonds. Due to the kinetic energy density on which they depend, the ELF and LOL have homogeneous chemical compositions. Since there are more electron pairs, ELF is dredged up, and LOL can discern localized orbital gradients That become agitated when they overlap (interpreted by savin and silvi) (Sowrirajan et al., 2022). The ELF and LOL outcomes are constrained to the range of 0.0 to 1.0. Values between 0.5 and 1.0 indicate a comprehensive representation of both bonding and nonbonding localized electrons. The contours where electrons are intended to be delocalized are described by the diminutive value < 0.5. LOL exceeds >0.5 in contours where electron localization dominates the electron density. The high localization of electrons is attributed to the durability of the covalent bond within the contours that possess significant values. ELF and LOL color shades show the presence of bonding and non-bonding electrons. In the figure, across the hydrogen atoms H7, H8, H10 and H15, the red color implies high ELF and LOL values, which further spot the presence of highly localized electrons. The central region of a hydrogen atom is white on the LOL color-filled map, noting that the electron density is higher than the upper limit of the color shade (0.8au) of the color scale. Within the header 24DCA the delocalized electron cloud that surrounds the carbon atoms in the phenyl ring can be seen by the blue color. Whereas a small blue circle around the carbon atoms, chlorine atoms Cl₉ and Cl₁₁, and oxygen atom O₁₃ depict the region of electronic depletion between the valence layer and inner layer. As a consequence, the interpretation is consistent with the lone pair theory, which is derived from the Lewis structure.

4.8 Prediction of the ligand's (ADMET) profile and Drug scans approach

The most recent technique to pinpoint the compounds that are suggested to confirm the nature of a potentially active drug is called "drug-likeness." Drug similarity analysis is carried out based on several pharmacophoric characteristics of the sample, including those of bioavailability, reactivity, and

metabolic stability. The Swiss ADME web server generated bioavailability radar shown in Fig. 10 was used to confirm the "drug-likeness" requirements which use Lipinski's (Ro5) filter method descriptors, a chemical or drug compound is more likely to be a potential drug for human use if it meets specific requirements including MW< 500 Dalton, HBA< 10, HBD < 5, and lipophilicity laid out as log p <5 TPSA<140, and rotatable bonds <10) (Elangovan et al., 2023). Within that study, the title compound 24DCA shows a molecular weight of 189.09 g/mol within the acceptable range, and the number of hydrogen bond donors and acceptors is 0 and 5 for each. Besides that, the number of rotatable bonds for the current drug is 2, and Lipinski's rule count is less than 10. The molar refractivity of the drug currently being used, 24DCA, is 46.66, which is between the threshold ranges of 40 and 130. The capabilities of drug transportation, including blood-brain barrier crossing, intestinal absorption, and penetration, can be inferred from the topological polar surface area, In the current scenario, 17.072 Å.2 is the predicted value, which is below the permissible range of 140Å.². The lipophilicity values of header composition are as follows: log P = 2.09, XLOGP3=2.72 and an MLOG P (octanol partition co-efficient) value of 2.94 less than 5 pointing to the compound's balanced hydrophobic/lipophilic nature. Further examination of the Ghose filter elements, such as molar refractivity and atom count (17), reveals that the title compound also complies with the Goose rule. The aforementioned requirements are often not violated by more than one orally bioactive medication. With a bioavailability score of 0.55, the substances included in the current scenario demonstrate strong oral bioavailability since they adhere closely to Lipinski's criteria.

The qualities evaluated using the Swiss ADME server exhibited beneficial attributes, such as fantastic solubility (log s= -3.06) and a high likelihood of gastrointestinal absorption, leading to a promising potential for oral bioavailability. Pharmacokinetic studies examine the effects of drug-to-drug interactions on the activity of P-gp substrates and the suppression of CYP (Cytochrome enzymes) activities (Priya et al., 2023a). The protease-activated protein gp enables the movement of active substances across the membrane of living organisms. Furthermore, it plays a crucial function in the

process of repelling objects. This compound is devoid of any p-gp substrates. This medicine exerts an influence on the enzymes CYP1A2, CYP2C19, CYP2C9, CYP2D6, and CYP3A4, which are impacted by the substance. Pan-assay interference chemicals (PAINS) are toxicological compounds that frequently induce spurious positive outcomes in high-throughput screening assays in medicinal chemistry. As an alternative, 24DCA has exhibited a 0-alert characteristic, signifying that it generates the intended analeptic effect exclusively through interaction with a particular receptor protein. PAINS interacting with multiple biological targets is more common than with a single target. Lead likeness aids chemical optimization performance by choosing the ideal starting locations. This chemical has no "lead-like" prospects despite therapeutic research. We further assessed the compound's hepatotoxicity, carcinogenicity, immunotoxicity, mutagenicity, and toxicity (LD50 880 mg/kg) using the Protox II web server. Endpoint and toxicity model results are in Table S4. The chemical has two AMES mutagenicity and carcinogenicity alerts.

Moreover, SwissADME encompasses a "BOILED-egg evaluation" feature delivering information on the human gastrointestinal absorption and blood-brain barrier permeability based on lipophilicity (WLOGP) and polarity (topological polar surface area, TPSA) as exemplified in Fig. S4 (top right). At certain points in the egg, the yolk and white align with the molecules that were expected to pass through the BBB and be absorbed by the gastrointestinal tract. Blue and red spots indicate compounds expected to be effluxed and not effluxed from the central nervous system by p-glycoprotein, respectively. The 24DCA molecule was positioned just inside the yolk region of the boiled egg model, as evidenced by the red dot position in Figure 10. This indicates that the chemical portrays a rapid absorption rate in the gastrointestinal tract and readily traverses the blood-brain barrier. In addition to reflecting that the molecule cannot traverse the blood-brain barrier, the red dot's presence in the white area signifies a rapid rate of absorption in the gastrointestinal system. The chemical compound exhibits medicinal properties due to its high gastrointestinal absorption and skin permeability values (Log kp = -

5.52 cm/s), together with its capacity to pass through the blood-brain barrier, emphasizing the potential for therapeutic use of the tested compound 24DCA.

4.9 Molecular Docking

In the design and discovery of new drugs, molecular docking has demonstrated itself to be an effective sequential asset. Molecular docking allows for a comprehensive understanding of the interactions between potential therapeutic compounds and the binding site of the targeted protein, providing new perspectives on underlying biological systems. Furthermore, molecular docking is an economical drug-designing approach that might offer a compact molecule, with more accurate and reliable binding positions when compared to experimental results (N. Elangovan et al., 2021a). It determines the types of interactions and binding conformations seen between the ligand and the active site protein, as well as their bonding distances and residue group proximity target protein to be used for docking with the ligand, is chosen based on the binding energy value; spices with a lower binding energy value are more likely to be able to dock with the target protein. This research looked at the DFT/B3LYP theory, applied to the 6-311++G (d, p) method, yielded the optimal ligand structure. from Gaussian 09W was used to dock into the active sites of receptors 1B1U, 1PCV, 2V32, 5ZEC and 6UEI of the insulin inhibitor, antifungal, antioxidant, antiseptic and antiviral proteins downloaded from the protein data bank PASS (prediction of atomic spectra) is a web-based tool that makes predictions about various types of activities. Concerning binding affinity with all of the targeted proteins, the ligand molecule was docked appropriately for its position and orientation (N. Elangovan et al., 2024b). Fig. 7 highlights optimized ligand-target protein orientations by the least energy of binding. All computations were done exploiting Auto dock to predict the ligand-protein active binding site interaction. Leveraging the Kollaman and Gasteiger phase process, the co-factors and polar hydrogen were appended to the suitable intended protein in pdb style, while the co-crystalline ligand water was eliminated. For the first two

ranks (1 and 2), the ranking capacity of the ligand with the targeted protein is pictured by Lamarckian docking binding energies (Kcal/mol), estimated inhibition constants (M), and reference RMSD values. The best pose was ascertained by having a higher binding energy value and the existence of a significant number of conventional hydrogen bonds. Considering the minimum energy of binding of interactions among ligands and proteins, the optimized ligand-target protein orientations are shown in Fig. 7. As illustrated in Fig. 7 the titled molecule contains a carbonyl group (C = O) that has hydrogen bond interaction with residues LEU A: 17(2.6Å), ARG A: 44 (2.23Å), ASN A: 200 (2.08Å), VAL A: 198(1.80Å) LEU A: 87, and CYS A: 73 (1.99Å) (3.05Å). The binding energies of proteins1B1U,1PCV, 2V32, 5ZEC and 6UEI are -4.55kcal/mol, -4.72kcal/mol, -5.69kcal/mol, -5.95 kcal/mol -6.18kcal/mol, respectively, with estimated inhibition constant values of 213.10 μ M, 214, 50 μ M, 67.58 μ M,43.15 μ M and 29.67 µM has been reported in Table S5. Amongst these four proteins, the header molecule reacted favorably with antifungal protein (6UEI) has the good binding energy-6.18kcal/mol, inhibition constant 29.67 µM, total intermolecular energy of -6.48 kcal/mol and RMSD value 69.88 Å with the higher number of hydrogen bonds to the amino acid residues LEU A: 87, and CYS A: 73. It is noteworthy that this molecule interacts with active sites containing oxygen atoms through hydrogen bonding. The preceding arguments promote the possibility of using 24DCA as potent antiviral medications.

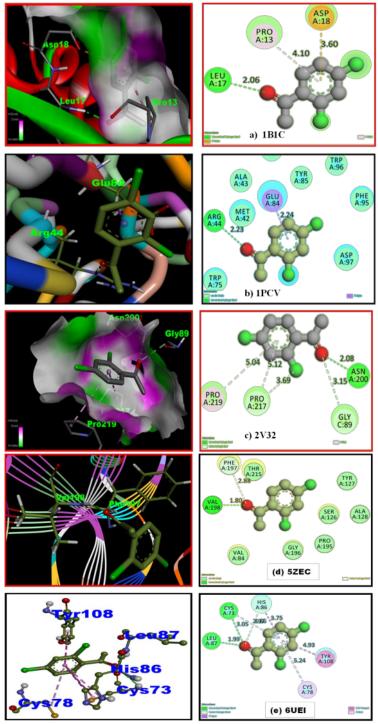


Fig. 7. Ligand 24DCA embedded in active sites of Insulyin inhibitor (a) 1B1U, Antifungal (b) 1PCV, Antioxidant (c) 2V32 Anticeptic (d) 5ZEC, Antiviral (e) 6UEI targeted proteins.

5.0 Conclusion

Ultimately, we successfully analyzed a 24DCA molecule in the DFT add method. The TD-DFT method was used to analyze the titled molecule solvent interaction. The structural, physicochemical, and biological features were analyzed using molecular spectroscopy methods. The Gaussian 09 software has been utilized for figuring out the optimum geometrical variables (bond lengths and bond angles) of the current material utilizing three distinct functional levels of theory and a single basis set. It is worth mentioning that the bond distance values obtained from the HF and CAM-B3LYP function levels are slightly shorter than those from the B3LYP technique. However, they are slightly longer than the B3LYP approach when compared to experimental data. The comparison of the scaled and experimental FT-IR and Raman wave numbers is supportive of one another. The allocations given for the different functional groups that exist in the molecule are further supported by the measured frequencies and PED output. These outcomes supported the validity of the methodology used to interpret the vibrational spectra. The NBO assessment method was adopted to scrutinize the stability of the molecule resulting from charge delocalization and hyper-conjugative interaction, Moreover, the energy assessments of the prominent molecular orbital theories of the headline compound were conducted relying on the B3LYP method. The molecular electrostatic contour reveals that the electronegative oxygen atoms possess the negative potential, while the positive potential is attributed to hydrogen atoms, which serve as the reactive sites in the molecule. The HOMO-LUMO energy gap of the 24DCA molecule is reduced by 4.253 eV using the B3LYP/6-311++G (d, p) method. This reduction has a significant effect on the molecule's intramolecular charge transfer and biological impact. The weak and strong attractive interactions, as well as the steric repulsion, that existed in the title compound, are visualized and identified using RDG analysis and VMD software. The outcome of analyzing the drug-likeness criteria by using Lipinski's (RO5) confirmed that the subject molecule had acceptable drug-likeness characteristics. The minimum energy

of binding for the 6UEI receptor was -6.18 kcal/mol, suggesting that the header compound seems to have an anti-viral-potential to be utilized in biomedical applications.

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