



Molecular Geometry, Vibrational Spectroscopic, Molecular Orbital and Mulliken Charge Analysis of 4-(carboxyamino)-benzoic acid: Molecular Docking and DFT Calculations

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Abstract

Structure based biological and chemical properties of 4-(carboxyamino)-benzoic acid has been studied by quantum chemical methods. The revamped geometric structure and its quantum chemical parameters were obtained by DFT-B3LYP/6-311G method. Normal mode analysis is performed to assign the fundamental vibrational frequencies as per the potential energy distribution (PED) by using the VEDA program. Simulation of IR and Raman spectral patterns are observed by refinement of scale factors. TD-DFT approach is used to explore the excited states of molecule and prediction of electronic absorption spectra. NMR chemical shifts of the molecule are determined by the gauge independent atomic orbital method. The molecular docking is performed to recognize the binding energy of the ligand with the dynamic site of protein. In our docking analysis, the protein 5DT6 shows the best results than other three proteins which could be used for further analysis. Further inter and intra molecular interactions, electrophilic, nucleophilic and chemical reactivity sites are found by molecular electrostatic potential, HOMO-LUMO and Global chemical reactivity descriptors. Thermodynamic property of the title compound is also reported. The determined quantum chemical parameters show high reactivity and the dipole moment was sufficiently high enough to induce nonlinear characteristics which are required for applications in optoelectronic devices.

Keywords: DFT; HOMO; LUMO; MEP; FMO; RDG; ADMET

Introduction

p-Aminobenzoic acid (PABA) shows very important roles in a wide variety of metabolic processes. PABA, a benzene ring with para (carbon 1 and 4) places of amine and carboxylic acid gatherings, is a zwitterion. PABA is a flexible reagent for structure expansion through direct hydrogen holding relationship, through both the carboxylic and amine utilitarian gatherings. The interest in carboxylate builds all in all and a benzoic acid subsidiary specifically keeps on developing as a result of their substance and natural

properties. Benzoic acids and their subordinants are significant underlying components for some, regular items, being engaged with different physiological cycles in plants. Among these accumulates, PABA has shown organic, therapeutic and modern interest. PABA is a notable compound of high biological importance and it is present in plant and animal tissues and also sometimes referred to as bacterial vitamin H₁, B_x or B₁₀ [1]. PABA is a well-known forerunner of folic acid [2], and has as of late been recognized as a coenzyme Q precursor [3,4], which opens new possibilities for its applica-

tion in medicine. It is a structure block utilized in plan of medications and as often as possible found as a design moiety in drugs. PABA displays a wide scope of restorative uses as cell reinforcement, antibacterial, antimutagenic, anticoagulant, fibrinolytic and immunomodulating specialist, defensive medication against UV-illumination and in indicative tests for the condition of the gastrointestinal tract [5]. Late investigations in agribusiness report the part of PABA in plant thermotolerance and its activity as compound inducer of foundational obtained obstruction against plant microorganisms [6]. Viewed as in the B-complex nutrient family and described by security and cost adequacy, PABA is oftentimes added alongside plant chemicals in nutriment media as a trigger of seed germination [7].

Due to the interest in the field of new materials, the chemical structures of organic-inorganic hybrid materials have been extensively investigated. The knowledge of binding energy and energy band gap explains the intrinsic properties like chemical reactivity and electrophilic sites of the compound predicted from quantum chemical calculations. To the best of our knowledge, no detailed quantum chemistry study has been performed for 4-(carboxyamino)-benzoic acid. The optimized structure and its geometric parameters are also found. The vibrational studies for each functional group are studied in detail using VEDA4 program. The nonlinear optical materials have potential applications in areas like optical limiting, signal processing, optical logic gates, laser radiation protection, etc. [8]. In addition to these properties, UV spectra, and thermodynamic properties are also examined for the compound. The suitable computational calculation of DFT has been performed with the help of Gaussian 09W package and the results obtained in the theoretical technique were evaluated.

Materials and Methods

The three dimensional atomic design of the title compound for theoretical computational work was downloaded from PubChem site. Every one of the computations was performed on Gaussian 09W programming with DFT method. For direct molecular structure implementation to compute vibrational spectra, DFT-B3LYP/6-311G technique was utilized. The molecular geometry and vibrational frequencies values of the molecule for title compound in ground state have been determined.

Results and Discussion

Geometrical structure

The optimized molecular structure of 4-(carboxyamino)-benzoic acid is shown in figure 1. The molecular geometry of the title compound is explained in terms of bond lengths, bond angles and dihedral angles. In carboxyamino group, the C-N bond distance are C6-N5 = 1.395 Å and C13-N5 = 1.3898 Å. The bond length of N5-H18 and O1-H19 are found at 1.0173 Å and 0.9809 Å respectively. The shortest and medium C-O bond distances are observed at 1.2354 Å (C12-O2) and 1.3614 Å (C12-O1). Alcolea Palafox M., *et al.* [9] reported the corresponding values for phenyl ring atoms (C-C) in the monomer form varies from 1.392 Å to 1.402 Å. C-C bond lengths varies at B3LYP level in monomer form from 1.3948 Å (for C10-C8, C6-C8, C6-C9, C11-C9, C7-C11) to 1.4543 Å (for C7-C12). And also an endocyclic C-C-C bond angle varies from 119.9 degree to 120°. For the title compound, totally two C-N-H bonds are appeared. The DFT calculations give the bond angles C6-N5-H18 = 115.9° and C13-N5-H18 = 115.7°. The carboxyamino and phenyl ring are joined with C6-N5-C13 bond. It makes an angle of 128.37 degree. The dihedral angle of phenyl group attached to the carboxyamino group ranges from -179.9° to +179.9°.

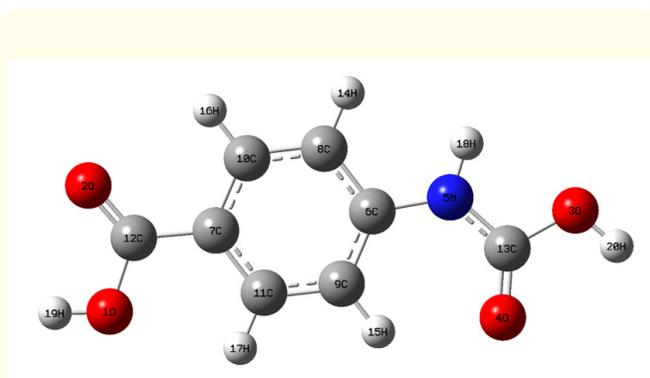


Figure 1: The optimized molecular structure of 4-(carboxyamino)-benzoic acid.

Spectral analysis

The molecule 4-(carboxyamino)-benzoic acid have 20 molecules with 54 normal modes of fundamental vibration. IR and Raman spectra of the title compound are shown in figure 2 and 3. All the normal modes of the vibrations are assigned based on the

potential energy distribution (PED) values computed using VEDA4 program and also in comparison with the literature. The theoretical frequencies are presented in table 1. The absorption through DFT method at 3491 cm^{-1} is assigned to N-H symmetry stretching mode and 3499 cm^{-1} is obtained in HF method. Wang and Ma [10] suggested N-H stretching bands in the region, $3365 - 3374\text{ cm}^{-1}$. Generally the aromatic C-H stretching vibration appears weakly between $3120 - 3000\text{ cm}^{-1}$. But in B3LYP calculations, it gives band in the range $3058-3133\text{ cm}^{-1}$. It shows that symmetry stretching mode is active. These four modes are pure stretching modes as is evident from PED column; they almost contribute 95%. The in-plane C-H bending vibrations normally appear in the range $1000 - 1300\text{ cm}^{-1}$ in the substituted benzenes and the out-of-plane bending vibrations occur in the region $750 - 1000\text{ cm}^{-1}$ region [11,12]. In aromatic compounds, the C-N stretching vibration usually appears in the region $1400 - 1200\text{ cm}^{-1}$. The absorption band located at 1331 cm^{-1} is attributed to C-N stretching vibration. An asymmetric stretching mode of C-N band appears at 1246 cm^{-1} . Usually, C=N stretching bands appears in the region $1598-1638\text{ cm}^{-1}$ [10]. In the present study, the C=N stretching mode is observed at 1623 cm^{-1} for HF method. The absorption is touchy for both the carbon and oxygen particles of the carbonyl gathering. Both have the same while it vibrates. Normally, the C-O stretching vibrations occur in the region $1260 - 1000\text{ cm}^{-1}$ [13]. In the present study, the C-O stretching vibration is assigned at 1314 cm^{-1} at B3LYP/6-311G level. The assignments of the COO group vibrations are closely agree with the literature values [14,15]. The in-plane COO bending vibration is localized at 839 cm^{-1} . A very weak COO out-of bending mode is found at 684 cm^{-1} . The band observed at $960, 871$ and 839 cm^{-1} was assigned to C-C-C stretching vibrations in DFT method. The O-H stretching vibrations are touchy to hydrogen bonding. The hydrogen bonding alters the frequencies of the stretching and bending vibration. The hydrogen bonding if present in five or six member ring system would reduce the O-H stretching band to $3200-3550\text{ cm}^{-1}$ region [16]. In the computed IR spectrum, a weak band observed at 3573 and 3538 cm^{-1} are assigned to O-H stretching mode of vibration. The potential energy distribution contribution to these modes is 100%.

UV spectral analysis

The UV spectral analysis represents that the electron absorption corresponds to the transition from the ground state to the first

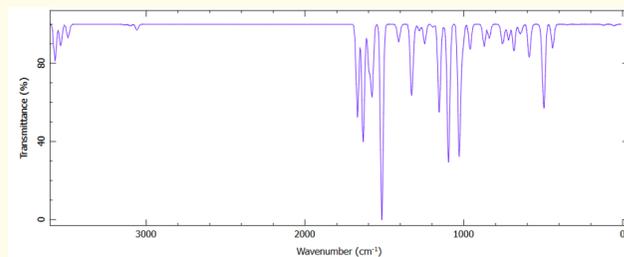


Figure 2: Computed FT-IR spectra for 4-(carboxyamino)-benzoic acid using DFT-B3LYP/6-311G method.

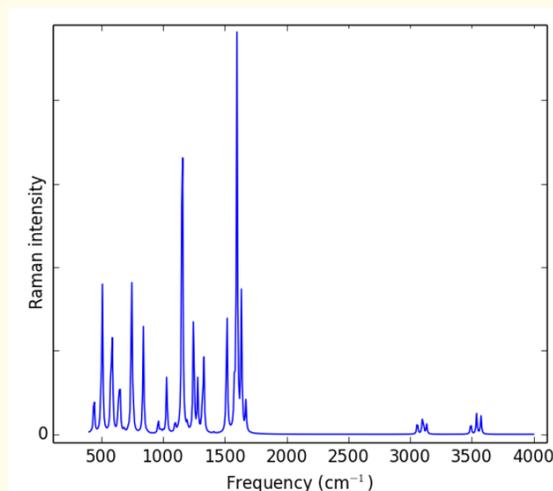


Figure 3: Computed Raman spectra for 4-(carboxyamino)-benzoic acid using DFT-B3LYP/6-311G method.

excited state. The band gap energy was calculated using the formula $E=hc/\lambda$, where h and c are constants, λ is the cut-off wavelength. The theoretical excitation energies, absorption wavelength and oscillator strength are summarized in table 2. The UV spectrum of the title compound is shown in figure 4. Energy gap were determined at the B3LYP/6-311G level utilizing TD-DFT approach in gas phase. These transitions are based totally on the groundwork of fundamental contribution of molecular orbitals. The orbitals con-

Mode	Symmetry Species Cs	Theoretical Wavenumbers (cm ⁻¹)				Vibrational Assignments (%PED)
		DFT B3LYP/6-311G		HF/6-31G		
		Un-scaled	Scaled	Un-scaled	Scaled	
1	A	50.65	48.93	50.72	45.79	τHOCN(81)
2	A	62.31	60.19	69.52	62.77	OUT CCCC(53)
3	A	83.23	80.39	88.16	79.60	τHNCO(80)
4	A	124.06	119.84	135.65	122.49	τCCCC(65)
5	A	148.77	143.70	167.80	151.52	OUT CCCC(68)
6	A	216.87	209.50	236.41	213.47	τCCNC(70)
7	A	288.11	278.31	310.88	280.72	τCCCO(81)
8	A	311.06	300.48	345.46	311.94	τHOCN(80)
9	A	361.02	348.74	391.27	353.31	τCCCC(62)
10	A	432.32	417.61	476.33	430.12	τCCCC(64)
11	A	455.56	440.06	496.82	448.62	τHCCC(77)
12	A	511.48	494.09	556.71	502.70	τHCCC(63)
13	A	512.57	495.14	573.25	517.64	τHCCC(73)
14	A	524.97	507.12	584.90	528.16	τHCCC(75)
15	A	597.12	576.81	641.01	578.83	τHNCO(81)
16	A	609.32	588.60	660.89	596.78	OUT CCCC(76)
17	A	658.54	636.15	716.09	646.63	τHOCC(85)
18	A	668.59	645.85	730.15	659.32	βCNC(77)
19	A	672.21	649.35	739.17	667.47	βCNC(35)
20	A	708.58	684.48	789.00	712.46	γCCO(61)
21	A	743.06	717.79	842.89	761.13	βCNC(77)
22	A	770.52	744.32	861.31	777.75	βCNC(69)
23	A	783.96	757.30	892.13	805.59	βCCN (71)
24	A	868.99	839.44	943.99	852.42	βCCC(61)
25	A	869.10	839.54	982.60	887.29	βOCO(68)
26	A	902.28	871.60	1024.39	925.02	γCCC(51)
27	A	993.92	960.12	1110.49	1002.7	βCCC (38)
28	A	1015.09	980.57	1135.91	1025.7	βHCC (57)
29	A	1033.43	998.29	1168.26	1054.9	γHCC(49)

30	A	1042.89	1007.4	1187.20	1072.0	βHCC (28)
31	A	1065.09	1028.8	1201.83	1085.2	γHCC(50)
32	A	1134.98	1096.3	1256.78	1134.8	βHNC (27)
33	A	1163.23	1123.6	1267.00	1144.1	βHOC (40)
34	A	1195.04	1154.4	1297.94	1172.0	βHOC (33)
35	A	1236.99	1194.9	1341.73	1211.5	βCCC (73)
36	A	1290.10	1246.2	1353.80	1222.4	αNC(36)
37	A	1325.19	1280.1	1422.95	1284.9	νCC(44)
38	A	1361.12	1314.8	1465.49	1323.3	νOC(70)
39	A	1375.35	1328.5	1502.96	1357.1	νCC(39)
40	A	1378.52	1331.6	1511.58	1364.9	νNC(63)
41	A	1459.59	1409.9	1591.84	1437.4	νCC(59)
42	A	1558.47	1505.4	1702.83	1537.6	νCC(47)
43	A	1569.61	1516.2	1729.04	1561.3	νCC(28)
44	A	1632.62	1577.1	1797.91	1623.5	νNC(30)
45	A	1653.52	1597.2	1817.38	1641.1	αCC(33)
46	A	1691.09	1633.5	1887.95	1704.8	νOC(73)
47	A	1728.13	1669.3	1898.80	1714.6	νCC(29)
48	A	3165.96	3058.3	3361.97	3035.8	νCH(91)
49	A	3207.52	3098.4	3405.66	3075.3	νCH(95)
50	A	3217.69	3108.2	3414.41	3083.2	νCH(91)
51	A	3244.09	3133.7	3454.93	3119.8	νCH(95)
52	A	3614.53	3491.6	3875.66	3499.7	νNH(100)
53	A	3662.73	3538.1	4003.83	3615.4	νOH(100)
54	A	3699.58	3573.7	4025.91	3635.4	νOH(100)

Table 1: Calculated vibrational frequencies of 4-(carboxyamino)-benzoic acid.

Abbreviations: ν - Stretching Vibrations; α - Asymmetric Stretching; τ - Torsion; β - in Plane Bending Vibrations; γ - Out of Plane Bending.

tributions ≤ 10% are neglected [17]. In this case, the strong transition was observed at 281 nm with maximum oscillator strength f=0.6462 with 4.41 eV energy gap and from HOMO-LUMO diagram is 5 eV. The other transitions are found at 258, 272 nm. UV visible spectra show that entire transitions belong to the quartz ultraviolet region (240-340 nm). So the compound can be used for quartz

optics. The peak was observed at 281 nm which was due to the presence of two π conjugated system of bonds (C=N, C=O) in the compound. It was evidence that the present compound is able to produce the optical energy with second and third fold frequency. The group contributions to the molecular orbital and the density of state (DOS) are found using Gauss-sum 2.2 program. The calculated TDOS diagram of the title molecule is shown in figure 5. The DOS spectra were created by convoluting the molecular orbital information with Gaussian curves of unit height [18].

NMR assessment

The DFT shielding calculations are fast, accurate and significant technique for structural determination of the carbon-hydrogen

λ_{max} (nm)	Band gap (eV)	Energy (cm^{-1})	f
258	4.81	38682.61	0.543
272	4.56	36722.67	0.612
281	4.41	35538.64	0.6462

Table 2: The theoretical excitation energies, absorption wavelength and oscillator strength.

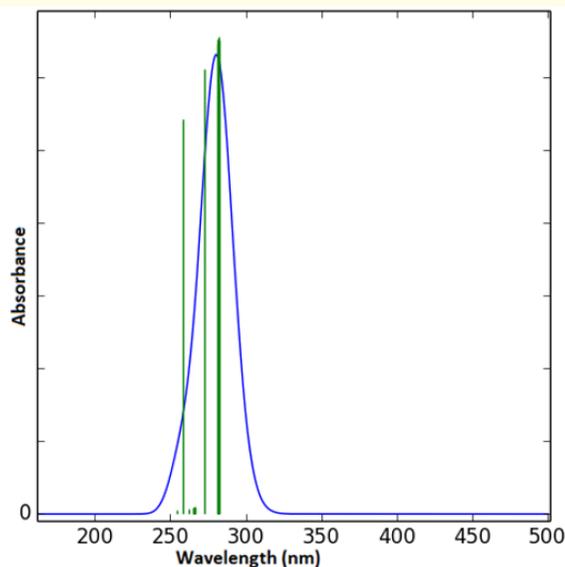


Figure 4: The UV spectrum of the title compound.

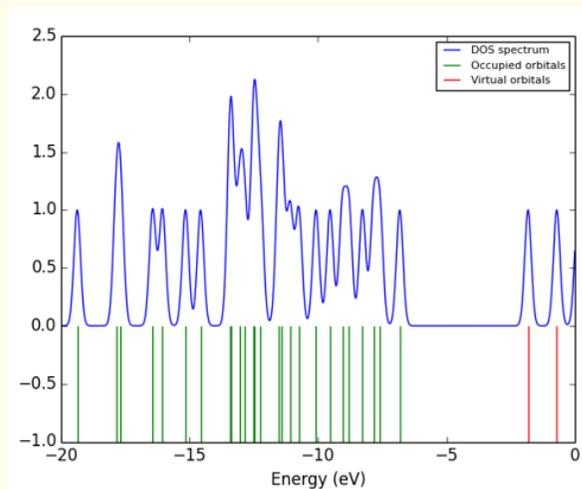


Figure 5: The calculated TDOS diagram.

framework of organic compounds [19]. The chemical shifts for ^1H and ^{13}C atoms of the compound are calculated for optimized structure, supported by gauge including atomic orbital (GIAO) method using B3LYP functional with 6-311G basis set. The graphical representations of computed values are proven in figure 6 and 7. Because of the distinctive idea of adjoining molecules and encompassing electrons, every core in the compound is synthetically inconsistent to another. So, NMR signals resonate for chemically unequal nuclei at various ppm values. The carbonyl carbons C12 and C13 are attached to two oxygen atoms and hence it is low deshielded than all other carbon atoms. Its chemical shift values for C15 and C17 were observed at 14.34 and 29.42 ppm respectively. C12 is much deshielded than all the other carbon atoms. The C7 carbon atom is more deshielded due to its connection with carbonyl carbon atom. Their computed values occurred at 64.41 ppm. The large values of C13 and C7 indicate that the large magnitude of the charges associated with these atoms, through the first one is highly positive and latter is highly negative. The chemical shifts of the carbon atoms, namely C8, C9, C10 and C11, almost lie in the same range, from 48.25 ppm to 71.5 ppm. Figure 7 clearly indicates that the chemical shift values of the hydrogen atoms are observed almost equal around 30 ppm, which shows that the chemical environment of hydrogen atoms is affected by oxygen, nitrogen or any carbon atoms. This is because of the theoretical calculations carried out in

the isolated gaseous phase of the theoretical chemical conversion assay. The signals of aromatic protons were observed at 24.07 - 27.26 ppm. The obtained results are found to be in good agreement with other investigators [20]. The theoretically calculated chemical shift of N5 was observed at 135.39 ppm. In addition, four oxygen peaks of O1, O2, O3 and O4 are observed at 121.6, -125.8, 137.3 and -36.35 ppm.

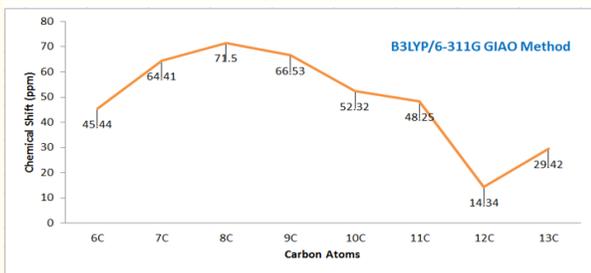


Figure 6: The graphical representation of chemical shift value (ppm) of carbon atoms.

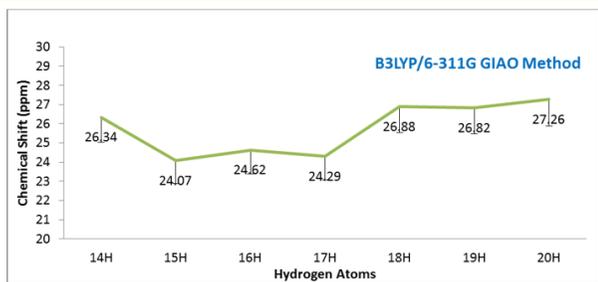


Figure 7: The graphical representation of chemical shift value (ppm) of hydrogen atoms.

Parameters of NLO properties

When electromagnetic fields interact with various media, it produces new fields with changes in frequency, phase, amplitude, or other characteristics of the incident fields resulting nonlinear optical (NLO) properties [21]. The total hyperpolarizability (β_{tot}) is the significant boundary to assess the NLO susceptibility. It means that the compound with large value of β_{tot} is predicted to be a potential

NLO active one and vice versa [22]. NLO behavior of compounds can be easily perceived and understood using theoretical calculations. Therefore, a theoretical correlation between the structure of the compound and NLO property may suggest the possibilities of design and synthesis of new NLO materials. First order hyperpolarizability (β) can be described by $3 \times 3 \times 3$ matrices. The second order hyperpolarizability is a fourth rank tensor. It is known as third-order NLO coefficient. The nonlinear parameters of the title compound are calculated by using DFT theory based on the finite field approach and HF method. The nonlinear parameters are reported in table 3. The vector component (β_{μ}) and average second order hyperpolarizability ($\langle \gamma \rangle$) are normally calculated from electronic field induced second harmonic generation experiments and are used to compare nonlinear optical response of different compounds. The total static dipole moment (μ_{tot}), isotropic molecular polarizability (α^{iso}), asymmetry parameter (η), Polarizability anisotropy ($\Delta\alpha$), first-order hyperpolarizability (β_{tot}) and second-order hyperpolarizabilities ($\langle \gamma \rangle$) are defined using the x, y, z components as [23-25].

$$\mu = (\mu_x^2 + \mu_y^2 + \mu_z^2)^{1/2} \text{----- (1)}$$

$$\alpha^{iso} = (\alpha_{xx} + \alpha_{yy} + \alpha_{zz})/3 \text{----- (2)}$$

$$\Delta\alpha = [\alpha_{xx} - (\alpha_{yy} + \alpha_{zz})/2] \text{----- (3)}$$

$$\eta_{\alpha} = [(\alpha_{yy} - \alpha_{zz})/(\alpha_{xx} - \alpha^{iso})] \text{----- (4)}$$

$$\beta_{tot} = [(\beta_{xxx} + \beta_{yyy} + \beta_{zzz})^2 + (\beta_{yyy} + \beta_{xxx} + \beta_{zzz})^2 + (\beta_{zzz} + \beta_{xxx} + \beta_{yyy})^2]^{1/2} \text{----- (5)}$$

$$\beta_{\mu} = \frac{\sum_{i=x,y,z} \mu_i (\beta_{ixx} + \beta_{iyy} + \beta_{izz})}{\sqrt{\mu_x^2 + \mu_y^2 + \mu_z^2}} \text{----- (6)}$$

$$\langle \gamma \rangle = \frac{1}{5} (\gamma_{xxxx} + \gamma_{yyyy} + \gamma_{zzzz} + 2\gamma_{xxyy} + 2\gamma_{xxzz} + 2\gamma_{yyzz}) \text{----- (7)}$$

For the (hyper)polarizabilities, the calculated values are in atomic units throughout the work. The atomic units have been converted into SI unit system. Conversion factors are 1 a.u. of $\alpha = 1.648778 \times 10^{-41} \text{ C}^2\text{m}^2\text{J}^{-2}$; 1 a.u. of $\beta = 3.206361 \times 10^{-53} \text{ C}^3\text{m}^3\text{J}^{-2}$; 1 a.u. of $\gamma = 6.235377 \times 10^{-65} \text{ C}^4\text{m}^4\text{J}^{-3}$. For B3LYP/6-311G, the calculated first order hyperpolarizability (β_{tot}) is $281.63 \times 10^{-53} \text{ C}^3\text{m}^3\text{J}^{-2}$ and average second order hyperpolarizability is $-0.7693 \times 10^{-61} \text{ C}^4\text{m}^4\text{J}^{-3}$. Urea is the prototype molecule generally considered as a reference for the comparison of good nonlinear optical properties.

The first-order static hyperpolarizability of title molecule is two times greater than that of urea ($\beta_{tot} = 138.365 \times 10^{-53} \text{ C}^3\text{m}^3\text{J}^{-2}$). It shows that title molecule is good candidate as materials for applications in nonlinear optics.

	B3LYP/6-311G Debye		B3LYP/6-311G (a.u.)		B3LYP/6-311G (a.u.)
μ_x	2.4071	β_{xxx}	78.9910	γ_{xxxx}	-2750.5895
μ_y	0.9477	β_{xxy}	-53.1922	γ_{yyyy}	-512.1349
μ_z	0.0007	β_{xyy}	-4.6846	γ_{zzzz}	-74.3812
μ_{total}	2.5869	β_{yyy}	13.1973	γ_{xyxy}	-721.5517
α_{xx}	-66.9074	β_{xxz}	-0.0071	γ_{xxzz}	-585.4700
α_{yy}	-74.2979	β_{xyz}	0.0029	γ_{yyzz}	-109.1318
α_{zz}	-76.7142	β_{yyz}	0.0056	γ_{yyzz}	-1233.88
α_{xy}	12.5758	β_{xzz}	2.2503		
α_{xz}	-0.0017	β_{yzz}	-3.0654		
α_{yz}	-0.0004	β_{zzz}	-0.0024		
α^{iso}	-72.6398	β_{tot}	87.835		
$\Delta\alpha$	8.5986	β_{μ}	225.07		
η_{α}	0.4215				

Table 3: Dipole moments, static electronic polarizabilities, first-order hyperpolarizabilities and second-order hyperpolarizabilities.

FMO analysis

According to FMO theory, HOMO and LUMO are important terms in quantum chemistry that reflect the bioactivity of compounds. The HOMO particularly donates electrons, while LUMO represents the ability to gain an electron. Thus, studies on FMO provide some useful information for physicists and chemists about the active mechanism [26]. The graphical representation of the FMOs for B3LYP/6-311G basis set is shown in figure 8. The global reactivity descriptors like chemical potential, electronegativity, hardness, softness and electrophilicity index can be calculated using DFT. The electronic chemical potential describes the escaping tendency of electron from a stable system. According to Parr and Pearson [27], it can be calculated as $\mu = (E_{HOMO} + E_{LUMO})/2$. The negative value of the electronic chemical potential is called electronegativity

(χ). Chemical hardness demonstrates the resistance to alteration in electron distribution and also it is correlated with the stability and reactivity of the chemical system. It is given by $\eta = \text{Energy gap}/2$. The global electrophilicity index (ω) is measured in terms of chemical potential and hardness. It assesses the lowering of energy due to maximal electron flow between donor and acceptor [28]. Table 4 represents the calculated quantum molecular descriptors of title compound for B3LYP/6-311G basis set. The energy value of HOMO and LUMO are -6.82 eV and -1.82 eV respectively. The band gap energy value of the title compound was calculated as 5.0 eV, which confirms that the molecule has a stable structure. The ionization energy value indicates that the energy value of 6.82 eV is required to remove an electron from HOMO. The lower value of electron affinity (1.82 eV) indicates that the compound readily accepts electrons to form bonds. All these values demonstrate the biological activity of the title compound with different molecules. The calculated value of electrophilicity index (3.73) describes the biological activity of the compound. The values of electronegativity, global hardness, global softness and chemical potential of the title compound are -4.32, 2.5, 0.4 and 4.32 respectively.

Molecular electrostatic potential analysis

Molecular electrostatic potential (MEP) is important to characterise inter and intra molecular electrostatic interactions [29].

S. No.	Parameters (B3LYP/6-311G)	Values
1	E_{HOMO} (a.u)	-6.82
2	E_{LUMO} (a.u)	-1.82
3	IA: Ionization energy = $-E_{HOMO}$ (eV)	6.82
4	EA: Electron affinity = $-E_{LUMO}$ (eV)	1.82
5	E_g : Energy gap = $E_{HOMO} - E_{LUMO}$ (eV)	5.0
6	η : Hardness = Energy gap/2 (eV)	2.5
7	μ : Electronic chemical potential = $(E_{HOMO} + E_{LUMO})/2$ (eV)	-4.32
8	ω : Electrophilicity index = $\mu^2/2\eta$	3.73
9	S: softness = $(1/\eta)$	0.4
10	χ : Electro negativity = $-\mu$	-4.32
11	$\Delta N_{max} = -\mu/\eta$	-1.72

Table 4: The calculated quantum molecular descriptors.

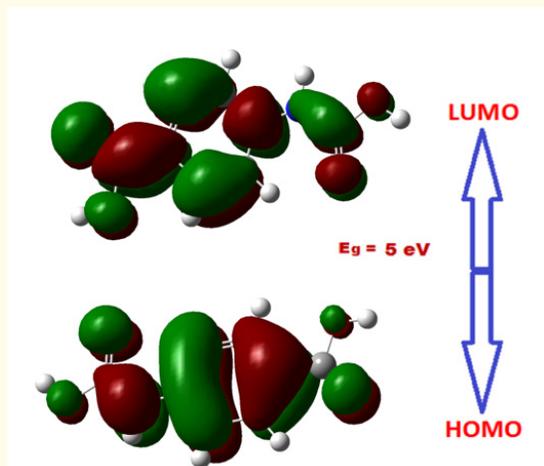


Figure 8: The graphical representation of the FMOs.

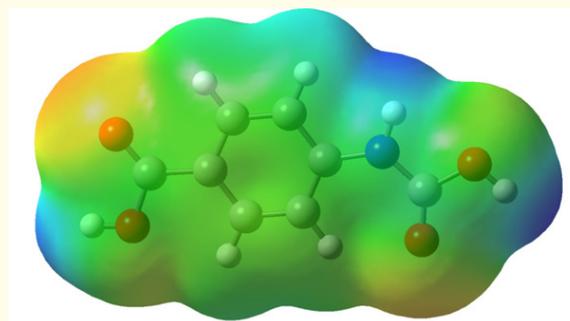


Figure 9: 3D map of the molecular electrostatic potential (MEP).

The charge spreads over the surface can be used to determine the interaction of nonlinear active molecules as well as to determine the type of chemical bond [30]. MEP is essential in structural biology to determine ligand-substrate interactions, drug-receptor and enzyme-substrate interactions [31]. To predict reactive molecular sites, the 3D map of the molecular electrostatic potential (MEP) surface was plotted for the total electron density of the title compound and is shown in figure 9. The electrostatic potential contour surface of the title molecule is presented in figure 10. The different values of the electrostatic potential at 3D map are mentioned by different colours. In Gauss View, the colour code in 3D map varies between -7.113 a.u. and +7.113 a.u. The red colour region (negative electrostatic potential) is related to electrophilic attack while the blue colour region (positive electrostatic potential) represents nucleophilic attack [32]. It has been revealed through figure 10 that there are four possible sites for electrophilic attack. As can be seen from the Figure, the negative region is mainly localised over oxygen atom. It is the most propitious site for electrophilic attack having the values -0.02 a.u. to -0.05 a.u. The nucleophilic attacks are around the hydrogen atoms with values in the range 0.05 to 0.0634 a.u. The maximum positive value 0.0634 a.u. corresponds to the hydrogen atom (H_2O) of carboxyl group attached with Oxygen (O_3).

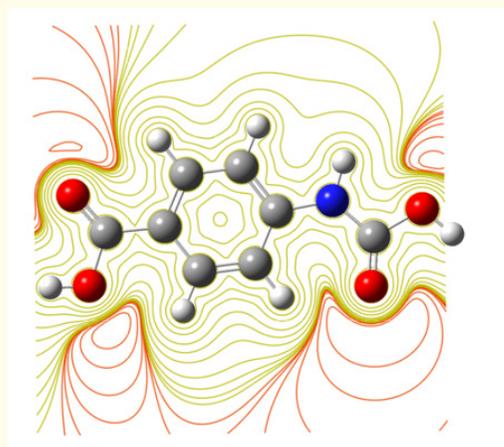


Figure 10: The electrostatic potential contour surface of the title molecule.

Mulliken atomic charges

The Mulliken atomic charge is directly related to the vibrational properties of the molecule and chemical bonds present in it. And also it shows that improved numerical stability and portrays the distribution of electrons in a better way [33]. Mulliken atomic charges affect the dipole moment, polarizability, electronic struc-

ture and more properties of molecular system [34]. Figure 11 represents the atomic charge distribution of the title compound for B3LYP/6-311G basis set. Atomic charges of all carbon atoms are negative except C6, C12 and C13; those are bonded with nitro and carboxylic acid groups. Due to the reason of electronegative atoms attached with the carbon atoms pull out the partial charges from these atoms. Hence these carbon atoms become positive charges. The extremely maximum protonated and deprotonated atoms are C13 (0.655 e) and N5 (-0.795 e) respectively. All the hydrogen atoms acquire positive Mulliken atomic charge. The charge distribution shows that O1, O2, O3 and O4 are negatively charged. The biggest value of charge (0.402 e) is noticed for H₂₀ may be due to hydrogen bonding.

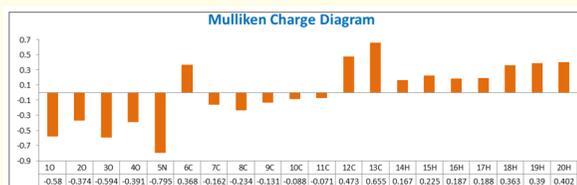


Figure 11: The atomic charge distribution of the title compound for B3LYP/6-311G basis set.

Force constant and reduced masses

The force constant and reduced mass of 4-(carboxyamino)-benzoic acid molecule calculated using DFT-B3LYP/6-311G and are shown in figure 12 and 13. In figure 12, Force constant can be divided into two regions; the first part range is from 0 to 1600 cm⁻¹. The second part range is from 1600 cm⁻¹ to near 3700 cm⁻¹. The highest force constant value is 13.92 mDyne/Å. In figure 13, the title molecule has highest reduced mass at 9.33 a.u. According to the formula $\nu = \frac{1}{2\pi} \sqrt{\frac{k}{\mu}}$, the reduced mass is inversely proportional to the frequency and force constant is directly proportional to the frequency. Figure 12 and 13 shows that the values of reduced mass are decreasing and force constant are increasing.

Donor-acceptor interactions

NBO is an efficient method to understand the intra and inter-molecular non-bonded interactions for bio-molecular compounds. Table 5 shows the occupancy and energies of i and j orbitals with

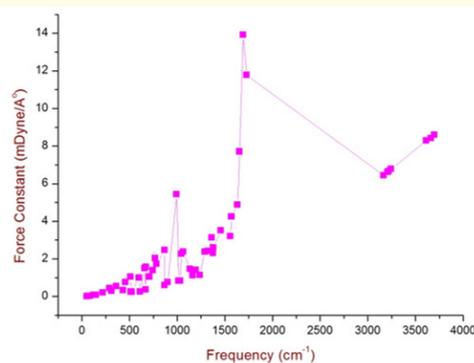


Figure 12: The force constant for title molecule.

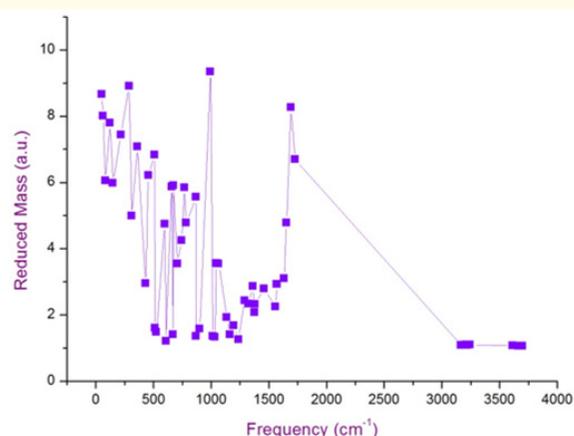


Figure 13: The reduced mass for title molecule.

the ΔE_{ij} of the most significant lone pair (LP) and bonding (σ/π) to antibonding (σ^*/π^*) transitions. Some significant orbital interactions and corresponding second order perturbation energies derived from the NBO computation for the compound 4-(carboxyamino)-benzoic acid. The Donor-acceptor interactions are happened between filled Lewis and empty non-Lewis NBOs after estimating their energies through second order perturbation theory. These interactions are termed as 'delocalization' corrections to the zeroth-order natural Lewis structure. The energy of these

delocalization interactions E(2) is proportional to the NBO interacting intensities and energies which gives the interactions among various parts of the molecules [35-37]. For each donor (i) and acceptor (j), the interaction energy E(2) was obtained from the standard second order perturbation approach $E_2 = \Delta E_{ij} = q_i \frac{F(i,j)^2}{E_i - E_j}$. Where the term q_i refers the occupancy of donor orbital, E_i and E_j denote the energies of donating and accepting orbitals and $F(i,j)^2$ is the KS matrix (Fock matrix) between i and j NBO orbitals. Table 8 shows the most significant interactions between Lewis and non-

Lewis orbital with O and N lone pairs. The N5 lone pair to the adjacent anti-bonding orbital of the ring [N5→π*(O4-C13)] was the significantly important. The stabilization energy of this interaction is 71.78 kcal/mol. The O4-C13 ties well and shows the biological activity because of the oxidation property. The stabilization energy E(2) associated with the interaction [O1→π*(O2-C12)] is obtained as 43.52 kcal/mol. The high intra molecular hyper conjugative interaction of π*(O2-C12) distributes to π*(C7-C11) with energy of 51.36 kcal/mol. Similarly, large energy is obtained by the transition π(O2-C12) to π*(C7-C11); π(C6-C9) to π*(C7-C11); π(C7-C11) to π*(C8-C10); π(C8-C10) to π*(C6-C9) with energies 28.53, 24.33,

Donor NBO(i)	Type	ED/e	Acceptor NBO(j)	Type	ED/e	E(2) ^a Kcal/mol	E(j)-E(i) ^b a.u.	F(i,j) ^c a.u.
O1 - C12	σ	1.99514	C7 - C10	σ*	0.02329	1.89	1.46	0.047
O1 - H19	σ	1.98916	O2 - C12	σ*	0.01875	1.07	1.27	0.033
O1 - H19	σ	1.98916	C7 - C12	σ*	0.05974	3.29	1.18	0.056
O2 - C12	σ	1.99522	C7 - C11	σ*	0.02420	1.37	1.60	0.042
O2 - C12	σ	1.99522	C7 - C12	σ*	0.05974	2.50	1.48	0.055
O2 - C12	π	1.98238	O2 - C12	π*	0.27326	1.07	0.36	0.019
O2 - C12	π	1.98238	C7 - C11	π*	0.38314	4.81	0.40	0.043
O3 - C13	σ	1.99251	N5 - C6	σ*	0.02891	3.08	1.30	0.057
O3 - H20	σ	1.98888	O4 - C13	σ*	0.02014	0.77	1.28	0.028
O3 - H20	σ	1.98888	N5 - C13	σ*	0.06783	3.24	1.17	0.056
O4 - C13	σ	1.99540	N5 - C13	σ*	0.06783	2.08	1.47	0.050
O4 - C13	σ	1.99540	N5 - H18	σ*	0.01388	1.04	1.44	0.035
O4 - C13	π	1.99491	O4 - C13	π*	0.36480	2.83	0.35	0.031
N5 - C6	σ	1.98488	O3 - C13	σ*	0.10377	1.71	1.08	0.039
N5 - C6	σ	1.98488	N5 - C13	σ*	0.06783	1.32	1.22	0.036
N5 - C6	σ	1.98488	N5 - H18	σ*	0.01388	0.58	1.20	0.024
N5 - C6	σ	1.98488	C6 - C8	σ*	0.02281	1.26	1.35	0.037
N5 - C6	σ	1.98488	C6 - C9	σ*	0.02467	1.63	1.36	0.042
N5 - C6	σ	1.98488	C8 - C10	σ*	0.01326	1.52	1.39	0.041
N5 - C6	σ	1.98488	C9 - C11	σ*	0.01487	1.33	1.39	0.038
N5 - C13	σ	1.98908	O3 - H20	σ*	0.00632	1.30	1.21	0.036
N5 - C13	σ	1.98908	O4 - C13	σ*	0.02014	1.44	1.38	0.040
N5 - C13	σ	1.98908	N5 - C6	σ*	0.02891	1.51	1.25	0.039
N5 - C13	σ	1.98908	C6 - C8	σ*	0.02281	1.82	1.39	0.045
N5 - H18	σ	1.98313	O3 - C13	σ*	0.10377	1.37	0.94	0.033
N5 - H18	σ	1.98313	O4 - C13	σ*	0.02014	4.72	1.19	0.067
N5 - H18	σ	1.98313	C6 - C9	σ*	0.02467	3.90	1.21	0.062
C6 - C8	σ	1.97208	N5 - C6	σ*	0.02891	1.28	1.10	0.034

C6 - C8	σ	1.97208	N5 - C13	σ^*	0.06783	3.54	1.11	0.057
C6 - C8	σ	1.97208	C6 - C9	σ^*	0.02467	4.16	1.25	0.064
C6 - C8	σ	1.97208	C8 - C10	σ^*	0.01326	3.04	1.28	0.056
C6 - C9	σ	1.97343	C6 - C8	σ^*	0.02281	4.37	1.23	0.066
C6 - C9	π	1.62122	C7 - C11	π^*	0.38314	24.33	0.29	0.075
C6 - C9	π	1.62122	C8 - C10	π^*	0.29521	16.43	0.28	0.062
C7 - C10	σ	1.97153	C7 - C11	σ^*	0.02420	4.47	1.24	0.067
C7 - C10	σ	1.97153	C8 - C10	σ^*	0.01326	2.82	1.26	0.053
C7 - C10	σ	1.97153	C8 - H14	σ^*	0.01432	2.57	1.11	0.048
C7 - C11	σ	1.97488	C7 - C10	σ^*	0.02329	4.59	1.24	0.067
C7 - C11	σ	1.97488	C9 - C11	σ^*	0.01487	2.65	1.27	0.052
C7 - C11	π	1.63175	O2 - C12	π^*	0.27326	28.53	0.23	0.075
C7 - C11	π	1.63175	C6 - C9	π^*	0.38058	18.23	0.27	0.063
C7 - C11	π	1.63175	C8 - C10	π^*	0.29521	23.27	0.27	0.073
C7 - C12	σ	1.97588	C7 - C11	σ^*	0.02420	2.73	1.23	0.052
C7 - C12	σ	1.97588	C8 - C10	σ^*	0.01326	2.51	1.25	0.050
C8 - C10	π	1.69335	C6 - C9	π^*	0.38058	22.64	0.28	0.072
C8 - C10	π	1.69335	C7 - C11	π^*	0.38314	16.26	0.29	0.062
C8 - H14	σ	1.97824	C6 - C9	σ^*	0.02467	4.37	1.06	0.061
C8 - H14	σ	1.97824	C7 - C10	σ^*	0.02329	4.01	1.07	0.059
C9 - C11	σ	1.97508	N5 - C6	σ^*	0.02891	4.66	1.09	0.064
C9 - C11	σ	1.97508	C7 - C12	σ^*	0.05974	3.65	1.13	0.058
C9 - H15	σ	1.97608	C6 - C8	σ^*	0.02281	4.57	1.04	0.061
C9 - H15	σ	1.97608	C7 - C11	σ^*	0.02420	4.24	1.06	0.060
C10 - H16	σ	1.97740	C6 - C8	σ^*	0.02281	4.40	1.04	0.060
C11 - H17	σ	1.97730	C6 - C9	σ^*	0.02467	4.57	1.05	0.062
C11 - H17	σ	1.97730	C7 - C10	σ^*	0.02329	4.83	1.06	0.064
O1	LP(1)	1.97975	O2 - C12	σ^*	0.01875	5.83	1.12	0.072
O1	LP(2)	1.83397	O2 - C12	π^*	0.27326	43.52	0.31	0.106
O2	LP(2)	1.85772	O1 - C12	σ^*	0.10158	33.96	0.54	0.123
O2	LP(2)	1.85772	C7 - C12	σ^*	0.05974	15.25	0.69	0.094
O3	LP(1)	1.97894	O4 - C13	σ^*	0.02014	6.39	1.13	0.076
O3	LP(2)	1.86072	O4 - C13	π^*	0.36480	40.92	0.30	0.105
O4	LP(2)	1.84065	O3 - C13	σ^*	0.10377	33.42	0.53	0.121
O4	LP(2)	1.84065	N5 - C13	σ^*	0.06783	21.45	0.67	0.110
N5	LP(1)	1.67344	O4 - C13	π^*	0.36480	71.78	0.24	0.119
N5	LP(1)	1.67344	C6 - C9	π^*	0.38058	34.85	0.30	0.092
O2 - C12	π^*	0.27326	C7 - C11	π^*	0.38314	51.36	0.04	0.074
O3 - C13	σ^*	0.10377	N5 - C6	σ^*	0.02891	1.77	0.13	0.052

Table 5: Stabilization energies obtained from the most significant lone pair to antibonding transition of the compound 4-(carboxyamino)-benzoic acid.

23.27 and 22.64 kcal/mol respectively.

Reduced gradient of density (RDG)

By employing NCI analysis, it is possible to determine the nature of the interactions based on electron density and the sign of the second derivative in the perpendicular direction of the bond (λ_2). RDG density is a dimensionless quantity and is written in the following form [38]:

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

We have represented, in figure 13, a 2D plot of RDG, as a characteristic of the electron density accelerated by means of the signal of the second eigenvalue of the matrix of Hessian. λ_2 is the greatest eigenvalue of the Hessian matrix which characterizes the fluctuation of the density in the vicinity of the critical point. From the value of the sign (λ_2)* ρ , we can distinguish between the one kinds of non-covalent interactions. The $-\lambda_2$ confirms a bonding interaction, such as hydrogen bonds, whereas $+\lambda_2$ shows nonbonding interactions and Vander Waals interactions can be determined by negligible values of λ_2 . According to figure 14, the red, green and blue circles indicate repulsive interactions (steric effect), VDW and H-bond respectively. Generally, VDW interactions have very low electron densities. The regions of interactions corresponding to the hydrogen bond and the repulsive interactions have a high density. The blue

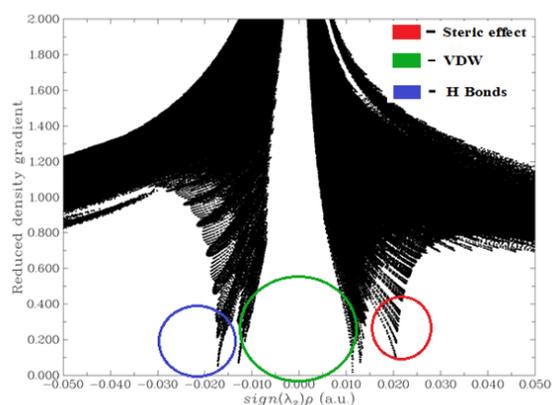


Figure 14: Graphical representation of the reduced density gradient versus the electron density.

circle confirms H-bond formation between the title molecules.

Thermodynamic properties

Thermodynamic data obtained from theoretical methods are

very important for understanding the chemical processes. DFT is efficient tool to predict various statistical thermodynamic properties of chemical compounds. On the basis of vibrational analysis at DFT-B3LYP and HF methods with basis set 6-311G and 6-31G level, the standard thermodynamic functions heat capacity, entropy, enthalpy, zero point vibrational energy, thermal energy and dipole moment for the title compound were found from the theoretical harmonic frequencies and presented in table 6. These parameters obtained here give available information for the further studies on the title compound and also estimate directions of chemical reactions according to the second law of thermodynamics. The ZPVE is lower in DFT/6-311G than HF/6-31G. Variations of the ZPVE seems to be important [39-41]. The biggest value of ZPVE of the title compound is 100.78414 Kcal mol⁻¹. As a result of HF method calculation the highest dipole moment was 2.6123 Debye. The

Thermodynamic Parameters (298.15 K)	DFT/6-311G	HF/6-31G
SCF energy	-664.73434	-660.6995
Total energy (Thermal), E_{total} (Kcal mol ⁻¹)	99.348	107.211
Translational energy, E_{tran} (Kcal mol ⁻¹)	0.889	0.889
Rotational energy, E_{rot} (Kcal mol ⁻¹)	0.889	0.889
Vibrational energy, E_{vib} (Kcal mol ⁻¹)	97.570	105.433
Zero point vibrational energy, E_0 (Kcal mol ⁻¹)	92.42369	100.78414
Heat capacity, C_v (Cal/Mol-Kelvin)	41.678	38.141
Entropy, S (Cal/Mol-Kelvin)	105.214	102.121
Rotational constants (GHZ)		
A	2.72089	2.79879
B	0.33939	0.34400
C	0.30175	0.30635
Dipole moment (Debye)	2.5869	2.6123

Table 6: Calculated thermodynamic parameters using HF/6-31G and DFT-B3LYP/6-311G methods.

highest values of specific heat capacity and entropy were gotten for B3LYP/6-311G method.

ADMET and drug like nature assessment

The significant rule or filter of drug-likeness is "Rule of Five" recommended by Lipinski, which describes the five principles of orally active compounds for physicochemical properties [42]. According to Lipinski's standard of five, that the molecule is discovered to be orally bioactive with the following factors, Molecular weight < 500, $Mi \log P < 5$, HBD is not exactly or equivalent to 5, HBA is not exactly or equivalent to 10 and Polar surface area is less than 120 \AA^2 [43]. These molecular properties were determined by using Molinspiration software. This web tool is freely accessible (<https://www.molinspiration.com>). The assessed drug-likeness properties of the title molecule are recorded in table 7. The bioavailability score is obtained to be 0.56. This shows that the title molecule is a better candidate for biological properties and hence applied for docking studies. Medication applicants ought to have ideal ADMET prediction. The ADMET for title molecule is obtained utilizing the pkCSM-ADME program which is freely accessible through web tool (<http://biosig.unimelb.edu.au/pkcsm/>). This is an important tool to enable medicinal scientists to discover the balance between pharmacokinetic, potency, and toxicity evaluation. The title compound was assessed by their Absorption, Distribution, Metabolism, Excretion and Toxicity were analyzed and classified in table 8. The CaCo_2 permeability and low permeability for skin is resolved to assess the absorption of title compound were obtained to be 0.0643 ns m^{-1} and $-2.735 \log \text{ kp}$. The Human Intestinal Absorption is the most important factor for a drug molecule which usually lies between 70 to 100%. It determines good intestinal absorption for

drug molecules. The HIA value of title molecule was found to be 97.63%, which is well absorbed for the human body [44,45]. The limitation of a drug to enter the brain is a critical limit that ought to be considered to decrease side effects and toxicity and the measured value for the title molecule is -0.824 [46]. The value of skin permeability is -2.735 which specifies that the compound cannot be absorbed by human skin. The pure water solubility was found to be low value (-2.513). This 4-(carboxyamino)benzoic acid clearly less harmful accordingly. The 4-(carboxyamino)benzoic acid is a non-inhibitor of P-glycoprotein and it doesn't interfere with the P-glycoprotein's in drug absorption and excretion. Here GI absorption (gastro intestinal) is high. It is a non-inhibitor of CYP1A2, CYP2C19, CYP2C9, CYP2D6, CYP3A4 enzymes. These things specify that this molecule can be safeguarded as an imminent aspirant for bioactive cases. Excretion also predicts to be non-substrate of Organic Cation Transporter-2 (OCT2). The toxicity of the AMES exhibiting negative shows that its non-cancer and non-mutagenic molecule. It does not inhibit hERG-I and hERG-II and also does not cause skin sensitivity. *T. pyriformis* Toxicity and minnow toxicity of the title compound was determined to be $0.282 \log \mu\text{g/L}$ and $2.525 \log \text{ mM}$ [47-49]. Every one of the parameters result mentions that the title molecule has been the biological activity of the drug in future.

Molecular docking

The Protein Data Bank (PDB) is the single, global archive for information about the 3D structure of biomacromolecules and their complexes, as determined by X-ray crystallography, NMR spectroscopy and cryoelectron microscopy. And also it includes more than a few Nobel Prize winning structures. Molecular docking is carried out for 4-(carboxyamino)-benzoic acid against active sites of

Descriptors	Values	Expected range
Hydrogen bond donor (HBD)	3	<5
Hydrogen bond acceptor (HBA)	4	<10
Partition coefficient ($Mi \log P$)	0.94	<5
Molecular weight (dalton)	181.15	<500
Polar surface area (PSA) (\AA^2)	86.62	<120
Number of rotatable bonds	2	<10
Molar refractivity	44.87	40-130
Bioavailability score	0.56	0.55

Table 7: Molinspiration property values of 4-(carboxyamino) benzoic acid.

Property	Model Name	Predicted Value	Unit
Absorption	Water solubility	-2.513	Numeric (log mol/L)
Absorption	Caco2 permeability	0.643	Numeric (log Papp in 10 ⁻⁶ cm/s)
Absorption	Human Intestinal absorption	97.63	Numeric (% Absorbed)
Absorption	Skin Permeability	-2.735	Numeric (log Kp)
Absorption	P-glycoprotein substrate	No	Categorical (Yes/No)
Absorption	P-glycoprotein I inhibitor	No	Categorical (Yes/No)
Absorption	P-glycoprotein II inhibitor	No	Categorical (Yes/No)
Distribution	VDss (human)	-2.129	Numeric (log L/kg)
Distribution	Fraction unbound (human)	0.574	Numeric (Fu)
Distribution	BBB permeability	-0.824	Numeric (log BB)
Distribution	CNS permeability	-3.349	Numeric (log PS)
Metabolism	CYP2D6 substrate	No	Categorical (Yes/No)
Metabolism	CYP3A4 substrate	No	Categorical (Yes/No)
Metabolism	CYP1A2 inhibitor	No	Categorical (Yes/No)
Metabolism	CYP2C19 inhibitor	No	Categorical (Yes/No)
Metabolism	CYP2C9 inhibitor	No	Categorical (Yes/No)
Metabolism	CYP2D6 inhibitor	No	Categorical (Yes/No)
Metabolism	CYP3A4 inhibitor	No	Categorical (Yes/No)
Excretion	Total Clearance	0.407	Numeric (log ml/min/kg)
Excretion	Renal OCT2 substrate	No	Categorical (Yes/No)
Toxicity	AMES toxicity	No	Categorical (Yes/No)
Toxicity	Max. tolerated dose (human)	0.689	Numeric (log mg/kg/day)
Toxicity	hERG I inhibitor	No	Categorical (Yes/No)
Toxicity	hERG II inhibitor	No	Categorical (Yes/No)
Toxicity	Oral Rat Acute Toxicity (LD50)	1.649	Numeric (mol/kg)
Toxicity	Oral Rat Chronic Toxicity (LOAEL)	1.697	Numeric (log mg/kg bw/day)
Toxicity	Hepatotoxicity	No	Categorical (Yes/No)
Toxicity	Skin Sensitisation	No	Categorical (Yes/No)
Toxicity	Pyriminism toxicity	0.282	Numeric (log ug/L)
Toxicity	Minnnow toxicity	2.525	Numeric (log mM)

Table 8: Pharmacokinetic descriptors of pkCSM-ADMET properties for 4-(carboxyamino)benzoic acid.

four proteins. Regarding this issue the protein structures (PDB ID: 3UOW, 4WIN, 4WIO, 5DT6) were obtained from <http://www.rcsb.org>. Docking studies were performed by iGEMDOCK 2.1 program. The numbers of runs taken are 70 with population size 800 and 10 numbers of solutions. iGEMDOCK provides target protein and compound interaction profiles of electrostatic (E), hydrogen bond-

ing (H), and Vander Waal's (V) interactions. The empirical scoring function of iGEMDOCK was calculated by $\text{Fitness} = \text{VDW} + \text{Hbond} + \text{Elec}$. Whereas, VDW is vander Waal energy, Hbond and Elec are hydrogen bonding energy and electro static energy respectively [50]. Table 9 described the docking result of 4-(carboxyamino)-benzoic acid with four target proteins using iGEMDOCK. The docked poses

of the compound with four target proteins were represented in figure 15, 16, 17 and 18. For molecular docking, the protein 3UOW, 4WIN and 4WIO are GMP Synthetase binding protein from Plasmodium Falciparum which is essential to causative agent of the fatal form of malaria. GMPS is a vital protein in the purine biosynthetic pathway. Protein 5DT6 is glutamine receptor. The binding energy

between compound and proteins 5DT6, 4WIO receptors are found to be -99.0424 and -95.151 respectively. In the interaction state of title molecule with 4WIN receptor, the binding energy of corresponding receptor is found to be -87.7694. The highest binding energy of 4-(carboxyamino)-benzoic acid is observed against 3UOW

S. No.	PDB ID	Energy	VDW	H-Bond	Elec	Interacting residues
1	3UOW	-69.6664	-46.1628	-22.7832	-0.720351	PHE-433, PRO-434, GLN-476, LYS-547, ILE-552, GLU-553, GLN-476, ILE-552, ILE-552, GLU-553, GLU-553
2	4WIN	-87.7694	-60.5374	-28.5852	1.35322	GLN-93, ASN-171, ASP-172, GLN-206, PRO-59, TYR-60, SER-61, HIS-170, ASN-171, ASN-171, ASP-172, ASP-172 SER-16, GLN-93,
3	4WIO	-95.151	-62.8058	-30.2687	-2.07647	ASN-169, HIS-170, ASN-171, ASP-172, HIS-208, GLY-58, ASN-169, HIS-170, HIS-208
4	5DT6	-99.0424	-60.1934	-36.6199	-2.2291	ARG-102, THR-97, ARG-102, THR-143, SER-148, THR-149, THR-149, TRP-150, TYR-66, SER-148, TYR-179, GLU-198

Table 9: Docking result of 4-(carboxyamino)-benzoic acid using iGEMDOCK.

and chemical reactivity predictions using quantum chemical computations. The theoretical spectral properties of title compound are successfully studied by IR, Raman, UV and NMR spectroscopic techniques. The various modes of vibrations are assigned on the results of PED value obtained from the normal coordinate analysis. TD-DFT calculations predict a peak at 281 nm which was due to the presence of two π conjugated system of bonds (C=N, C=O) in the compound. The calculated HOMO-LUMO band gap energy value is 5.0 eV. It decides the significant degree of charge transfer interactions taking place in the compound and also it is responsible for the enhanced chemical as well as biological activity of the molecule. Under reactivity predictions molecular electrostatic potential map clearly demonstrates that the possible sites for electrophilic attack are mainly localised over oxygen atom, while the nucleophilic sites are around the hydrogen atoms (H_2O). Normally, the electrophilic ability strengthens and nucleophilic ability weakens. It means that one move outwards from the inner core of the molecule. The title compound has negative chemical potential (-4.32 eV). This infers that it has a weaker tendency to change into the components it is created from and does not decompose. It very well may be delivered intentionally from the components. On the basis of RDG, It confirms H-bond formation between the title molecules. Thus, the present investigation provides a complete and reliable structural, spectral and biological activity of the compound. This investigation

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