

FT-IR and FT-Raman Spectral Investigation and DFT Computations of Pharmaceutical Important Molecule: Ethyl 2-(4-Benzoyl-2,5-Dimethylphenoxy) Acetate

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Abstract

The vibrational contribution studies of Pharmaceutical activity of Ethyl 2-(4-benzoyl-2,5-dimethylphenoxy)acetate (EBDA) have been performed using FTIR, FT-Raman analysis. More support on the experimental findings were added from the quantum chemical studies performed with DFT (B3LYP) method using 6-311++G (d, p) basis sets. The observed FT-IR and FT-Raman spectra have been compared with the calculated theoretical data. The calculated vibrational data have also been found in good agreement with the experimental results. Natural bond orbital analysis revealed important details of the electronic structure and dominant intramolecular interactions in Ethyl 2-(4-benzoyl-2,5-dimethylphenoxy)acetate. The HOMO and LUMO analysis reveals the possibility of charge potential (MEP), charge analysis also were investigated using theoretical calculations.

Keywords: Vibrational spectra; DFT; NBO; PED; HOMO-LUMO; Pharmaceutical important

Introduction

Phenoxyacetic acid, a well-known aryloxyacetic acid, which is useful in the treatment of insulin resistance, and hyperglycaemia, has been investigated by various researchers [1-3]. Phenoxyacetic acid and substituted phenoxyacetic acids have potential biological properties and so these acids are widely used in herbicides [4] and pesticides [5] formulations. Antimicro-bioactivity [6], anticancer, antitumor, analgesic, anti-inflammatory, plant growth regulation and inhibition of tillage are also some of its reported properties. Hydroxy benzophenones can be attained from natural products [7,8] and also by synthetic methods [9-11]. The enormous importance of these substances is essentially due to the diverse biological and chemical properties they acquire. Benzophenone analogues with nitro substituent exhibit significant in vivo antitumor activity and they have been reported to show activity such as immune modulators [12]. The title compound Ethyl 2-(4-benzoyl-2, 5-dimethylphenoxy) acetate (EBDA), has been synthesized via Fries rearrangement of hydroxy benzophenone [13].

In recent years, there has been an increasing interest in the application of DFT calculations to pharmaceutical drugs as calculations provide additional interpretation of the vibrational spectroscopic data collected [14-17]. During the past decade, DFT [18] has been accepted by the ab initio quantum chemistry community as a popular approach for the computation of molecular structure, vibrational frequencies and energies of chemical reactions. Calculation of vibrational frequencies using DFT provides a promising cost effective approach for calculating vibrational spectra of large molecules. In recent theoretical studies, the harmonic vibrational frequencies for a larger number of small and well studied organic molecules have been computed with HF, MP2 and DFT methods [19-21].

Vibrational spectroscopy is highly sensitive to the structural changes and is useful for the study of the pharmacological properties [22,23]. Accordingly, in order to understand the relationship between molecular structure and biological activity, the knowledge of the electronic structure and complete vibrational spectra are particularly important. Information about the geometry and structure of the

molecule and its electrostatic potential surfaces, together with a complete analysis of the vibrational spectra using Raman and infrared techniques, based on frequency, intensity and potential energy distribution over the internal coordinates helps in understanding the structural and spectral characteristics, by allowing us to obtain a quantitative as well as qualitative understanding of energy distribution. *Ab initio* HF and DFT calculations have been independently performed as they form the basis of the assignment of the vibrational spectrum.

The present work deals with the optimization of molecular structure and analysing the vibrational spectra using FT-Raman and FT-IR techniques for the title compound and comparing them with the theoretical Raman and IR spectra computed by DFT method. Calculated Potential energy distribution enables to make a detailed assignment of the vibrational spectra. Natural bond orbital analysis has been performed to identify the possible intra- and inter- molecular interactions present in the compound. HOMO-LUMO energy gap calculated confirms the occurrence of charge transfer interaction. Electrostatic potential analysis has also been made to identify the mapping surface of the molecule.

Materials and Method

Computational

Molecular structure, vibrational frequencies and energies of optimized geometries have been computed employing DFT [24] with

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Gaussian '09 program package [25] using Becke's three parameter hybrid exchange functional with Lee-Yang-Parr correlation functional (B3LYP) [26-28] as a cost effective approach. The basis set 6-311++G (d, p) augmented by 'd' polarization functions on heavy atoms and 'p' polarization functions on hydrogen atoms have been used [29,30]. Absolute Raman intensities and infrared absorption intensities have been calculated with harmonic approximation at the same level of theory as used for optimized geometries from the derivatives of polarizability and dipole moment associated with each normal mode. Owing to the complexity of the molecule, Potential Energy Distribution has been carried out to obtain complete information of the molecular motions involved in the normal modes. The experimentally observed spectral data of the compound are found to be in good agreement with the spectral data obtained by quantum chemical calculations.

The vibrational modes have been assigned on the basis of PED analysis using VEDA program [31]. Vibrational wave numbers calculated have been scaled [32] with the scale factor in order to figure out how the calculated data are in agreement with those of the experimental ones. Calculated vibrational frequencies have been scaled down by using the scaling factor [32] to offset the systematic error caused by neglecting anharmonicity and electron density. Raman activities (S_i) calculated have been converted to relative Raman intensities (I_i) using the following relationship derived from the basic theory of Raman scattering [33,34].

$$I_{i} = \frac{f(v_{o} - v_{i})^{4}S_{i}}{v_{i} \left[1 - \exp\left(\frac{-hcv_{i}}{kT}\right)\right]}$$
(1)

where v_o is the exciting frequency (in cm⁻¹ units), v_i is the vibrational wave number of the ith normal mode, h, c and k are universal constants, and S is the suitably chosen common scaling factor for all the peak intensities.

Experimental

The infrared spectrum of the sample was recorded between 4000 and 400 cm⁻¹ on a Mattson 1000 FT-IR spectrometer which was calibrated using polystyrene bands. The sample was prepared as a KBr disc. FT-Raman spectrum of the sample was recorded between the region 3500 and 50 cm⁻¹ on a Bruker FRA 106/S FT-Raman instrument using 1064 nm excitation from an Nd: YAG laser. The detector used was a liquid nitrogen cooled Ge detector.

Results and Discussion

Optimized geometry

Initial geometry generated from standard geometrical parameters was optimized and equilibrium geometry has been determined by the energy minimization. The ground state optimized structure of the molecule is presented in Figure 1. The optimized structure of EBDA and experimental structure of EBDA [13] were compared. The agreement between the optimized and experimental crystal structure is quite good showing that the geometry optimization almost exactly reproduces the experimental conformation (Figure 1) (Table 1).

The relative energies of the molecule have been calculated employing *ab initio* functions and DFT functional (B3LYP). The optimized structural parameters (bond lengths, bond angles, dihedral angle) have been compared with those obtained experimentally for the EBDA as shown in Table 1. The table reveals that, all the theoretical values agree well with the experimental values. Calculated C-H bond length is around 1.08 Å, while the experimental value from the neutron

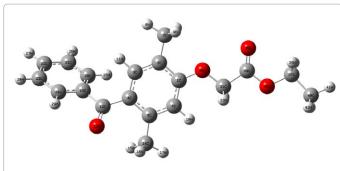


Figure 1: Optimized molecular structure of EBDA.

Bond	Value		Value		Value
length	(Å)	Angle	(/°)	Dihedral angles	(/°)
C ₁ -C ₂	1.407	C ₁ -C ₂ -C ₃	117.0	C ₁ -C ₂ -C ₃ -C ₄	1.37
C ₂ -C ₃	1.387	C ₂ -C ₃ -C ₄	123.2	C ₂ -C ₃ -C ₄ -C ₅	-0.42
$C_3 - C_4$	1.407	C ₃ -C ₄ -C ₅	119.1	C ₃ -C ₄ -C ₅ -C ₆	-1.14
$C_4 - C_5$	1.409	C ₄ -C ₅ -C ₆	118.1	$C_{6}-C_{1}-C_{2}-C_{7}$	179.33
C ₅ -C ₆	1.399	C ₁ -C ₂ -C ₇	120.6	C ₃ -C ₂ -C ₇ -H ₈	-0.66
C ₂ -C ₇	1.506	C ₂ -C ₇ -H ₈	110.6	C ₃ -C ₂ -C ₇ -H ₉	120.02
C ₇ -H ₈	1.091	C ₂ -C ₇ -H ₉	111.2	C ₃ -C ₂ -C ₇ -H ₁₀	-121.56
C ₇ -H ₉	1.094	C ₂ -C ₇ H ₁₀	111.3	C ₁ -C ₂ -C ₃ -H ₁₁	179.73
C ₇ -H ₁₀	1.094	C ₂ -C ₃ -H ₁₁	118.0	C ₂ -C ₃ -C ₄ -C ₁₂	175.6
C ₃ -H ₁₁	1.084	C ₃ -C ₄ -C ₁₂	119.5	C ₃ -C ₄ -C ₁₂ -O ₁₃	-142.36
C ₄ -C ₁₂	1.494	C ₄ -C ₁₂ -O ₁₃	121.2	C ₃ -C ₄ -C ₅ -C ₁₄	179.41
C ₁₂ -O ₁₃	1.223	C ₄ -C ₅ -C ₁₄	123.4	C ₄ -C ₅ -C ₁₄ -H ₁₅	42.79
C ₅ -C ₁₄	1.511	C ₅ -C ₁₄ -H ₁₅	112.1	C ₂ -C ₅ -C ₁₄ -H ₁₆	-75.76
C ₁₄ -H ₁₅	1.091	C ₅ -C ₁₄ -H ₁₆	110.9	C ₄ - C ₅ -C ₁₄ -H ₁₇	163.87
C ₁₄ -H ₁₆	1.092	C5-C14-H17	110.3	C ₄ -C ₅ -C ₆ -H ₁₈	-178.64
C ₁₄ -H ₁₇	1.092	C5-C6-H18	118.1	C ₃ -C ₄ -C ₁₂ -C ₁₉	37.37
C ₆ -H ₁₈	1.082	C ₄ -C ₁₂ -C ₁₉	119.6	C ₄ -C ₁₂ -C ₁₉ -C ₂₀	32.08
C ₁₂ -C ₁₉	1.505	C ₁₂ -C ₁₉ -C ₂₀	122.6	C ₁₂ -C ₁₉ -C ₂₀ -C ₂₁	176.14
C ₁₉ -C ₂₀	1.401	C ₁₉ -C ₂₀ -C ₂₁	120.4	C ₁₉ -C ₂₀ -C ₂₁ -C ₂₂	0.9
C ₂₀ -C ₂₁	1.394	C ₂₀ -C ₂₁ -C ₂₂	120.1	$C_{20} - C_{21} - C_{22} - C_{23}$	-0.89
C ₂₁ -C ₂₂	1.394	C ₂₁ -C ₂₂ -C ₂₃	119.9	C4-C12-C19-C24	-151.92
C ₂₂ -C ₂₃	1.396	C ₁₂ -C ₁₉ -C ₂₄	118.2	C ₁₂ -C ₁₉ -C ₂₀ -H ₂₅	-2.63
C ₁₉ -C ₂₄	1.402	C ₁₉ -C ₂₀ -H ₂₅	119.9	C ₁₉ -C ₂₀ -C ₂₁ -H ₂₆	-179.03
C ₂₀ -H ₂₅	1.083	C20-C21-H26	119.8	C ₂₀ - C ₂₁ -C ₂₂ -H ₂₇	179.4
C ₂₁ -H ₂₆	1.084	$C_{21} - C_{22} - H_{27}$	120.0	C ₂₁ -C ₂₂ -C ₂₃ -H ₂₈	-179.81
C ₂₂ -H ₂₇	1.084	C22-C23-H28	120.0	C ₂₂ -C ₂₃ -C ₂₄ -H ₂₉	-178.49
C ₂₃ -H ₂₈	1.084	C23-C24-H29	120.9	$C_{3}-C_{2}-C_{1}-O_{30}$	179.15
C ₂₄ -H ₂₉	1.083	C ₂ -C ₁ -O ₃₀	115.1	$C_2 - C_1 - C_{30} - C_{31}$	179.02
C ₁ -O ₃₀	1.366	$C_{1} - C_{30} - C_{31}$	119.1	C ₁ -C ₃₀ -C ₃₁ -H ₃₂	60.08
C ₃₀ -C ₃₁	1.408	C ₃₀ -C ₃₁ -H ₃₂	111.7	C ₁ -C ₃₀ -C ₃₁ -H33	-60.77
C ₃₁ -H ₃₂	1.097	$C_{30} - C_{31} - H_{33}$	111.7	$C_{1}-C_{30}-C_{31}-C_{34}$	179.4
C ₃₁ -H ₃₃	1.096	C ₃₀ -C ₃₁ -C ₃₄	108.7	C ₃₀ -C ₃₁ -C ₃₄ -O ₃₅	-3.75
C ₃₁ -C ₃₄	1.522	$C_{31} - C_{34} - O_{35}$	126.4	C ₃₀ -C ₃₁ -C ₃₄ -O ₃₆	176.59
C ₃₄ -O ₃₅	1.2	C ₃₁ -C ₃₄ -O ₃₆	108.6	$C_{31} - C_{34} - C_{36} - C_{37}$	179.3
C ₃₄ -O ₃₆	1.347	$C_{34} - C_{36} - C_{37}$	116.6	C ₃₄ -C ₃₆ -C ₃₇ -H ₃₈	-60.27
C ₃₆ -C ₃₇	1.453	C ₃₆ -C ₃₇ -H ₃₈	108.4	C ₃₄ -C ₃₆ -C ₃₇ -H ₃₉	56.65
C ₃₇ -H ₃₈	1.092	C ₃₆ -C ₃₇ -H ₃₉	108.4	$C_{34} - C_{36} - C_{37} - C_{40}$	178.22
C ₃₇ -H ₃₉	1.092	C ₃₆ -C ₃₇₋ C ₄₀	107.6	C ₃₆ -C ₃₇ -C ₄₀ -H ₄₁	179.88
C ₃₇ -C ₄₀	1.514	C ₃₇ -C ₄₀ -H ₄₁	109.5	C ₃₆ -C ₃₇ -C ₄₀ -H ₄₂	-60.6
C ₄₀ -H ₄₁	1.093	C ₃₇ -C ₄₀ -H ₄₂	111.1	C ₃₆ -C ₃₇ -C ₄₀ -H ₄₃	60.32
C ₄₀ -H ₄₂	1.092	C ₃₇ -C ₄₀ -H ₄₃	111.1		
C ₄₀ -H ₄₃	1.092				

 Table 1: Optimized parameters of EBDA on B3LYP/6-31++G(d,p) level.

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diffraction method is 0.93 Å. The difference in C-H bond is due to experimental values taken from X-ray diffraction method that is solid phase, but the theoretical method is the gas phase. From the above figure it is seen that the EBDA molecule consists of mono substituted phenyl ring (Ph1), Tetra substituted phenyl ring (Ph2), three methyl groups and two methylene groups. Table shows that the bond length of C_{12} - O_{13} and C_{34} - O_{35} varies from 1.20 Å-1.22 Å whereas the bond length of C_1 - O_{30} and C_{34} - O_{36} varies from 1.34 Å-1.36 Å, which substantiate the single bond and double bond nature of carbonyl group. The C-C bond length of C_5-C_{14} and $C_{20}-C_7$ is around 1.50 Å while the normal C-C bond length is 1.40 Å. The increase in bond length from the normal value is due to electron transfer from a ring (Ph2) to the methyl group. EBDBA molecular structure is shown as a non planer surface due to the twisting between Ph1 to Ph2 and this twisting takes place due to the steric repulsion between the ring hydrogen H_{25} and H_{11} . From the table, it is seen that all the experimental bond angles are in good agreement with the theoretical values.

The decrease in endocyclic angles C_{23} - C_{22} - C_{21} and C_{20} - C_{19} - C_{24} from the normal value shows the electron with drawing nature of ring. The endocyclic angle of Ph2, C_4 - C_5 - C_6 (118.13°) is decreased and the exocyclic angle, C_4 - C_5 - C_{14} (123.384°) is increased from the normal value which confirms the heavy substitution of the ring. Similarly the endocyclic angle C_3 - C_2 - C_1 (116.99°) and exocyclic angle C_3 - C_2 - C_7 (122.42°) shows that the methyl substitution of the ring Ph2 is at the para position.

NBO Analysis

NBO analysis has been proved to be an effective tool for chemical interpretation of hyper conjugative interaction and electron density transfer (EDT) from filled lone electron pairs of the n(Y) of the "Lewis base" Y into the unfilled anti bond $\sigma^*(X-H)$ of the "Lewis acid" X-H in X-H.....Y hydrogen bonding systems [35]. Also, in order to obtain structure of molecule, the main natural orbital interactions have been analyzed with the NBO 5.0 program [36]. Lowering of the orbital energy due to the interaction between the doubly occupied orbital and the unoccupied ones is a very convenient guide to interpret the molecular structure. In energy terms, hyper conjugation is an important effect [37,38] in which an occupied Lewis-type natural bond orbital is stabilized by overlapping with a non Lewis-type orbital. This electron delocalization can be described as a charge transfer from a Lewis valence orbital (donor), with a decrease of its occupancy, to a non-Lewis orbital (acceptor). Several other types of valuable data, such as directionality, hybridization and partial charges, were analyzed in the output of NBO analysis.

Table 2 gives the second-order perturbation energy values, Eij(2) corresponding to the interactions and the overlap integral of each orbital pair. From the analysis, it is seen that the intermolecular C-H....O Hydrogen bonding is formed due to the overlap between the LPO₁₃ and σ^* (C₁₄-H₁₇) which result in ICT causing stabilization of the Hydrogen bond system. The NBO analysis of EBDA clearly shows the evidence of the formation H-bonded interaction between oxygen lone pairs and σ^* (C₁₄-H₁₇) antibonding orbital having stabilization energy 99.84 KJ/mol (Table 2).

The larger the E(2) values, the more intensive is the interaction between the electron donors and the electron acceptors. Among the most energetic donor acceptor NBO interactions, those involving p-type lone pair of oxygen atom, $\sigma(C_{40}-H_{43})$ with σ^* ($C_{14}-H_{17}$) antibonding have energy contributions 4661.89KJ-mol⁻¹ which shows the intermolecular dislocations in EBDA. The intermolecular hyper

conjugative interactions are formed by the orbital overlap between $\pi(C-C)$ and $\pi^*(C-C)$ bond orbital's, which results in intermolecular charge transfer (ICT) and causes stabilization of system. Table shows the most important interactions between Lewis and non-Lewis orbital with oxygen lone pairs and the second order perturbation energy values E (2) corresponding to this interaction. From the Table 2. it is seen that a very strong interaction between the p-type orbital containing the lone electron pair of O_{35} and neighbour $\sigma^*(C_{31}-C_{34})$ anti bonding orbital exists.

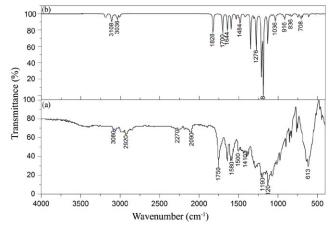
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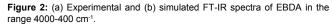
Vibrational analysis

The vibrational spectral analysis was performed on the basis of the characteristic vibrations of the amino group, hydroxyl group, carbonyl group and methyl group. The computed wave numbers, their IR intensities and Raman activity corresponding to different modes are listed in Table 3 along with detailed assignments. The observed and simulated FT-IR and FT-Raman spectra and selected vibrational

Demon (i)	A	E(2) ^a	E(j)-E(i) ^b	F(ij)⁰
Donar (i)	Acceptor (j)	(KJ/mol)	a.u.	a.u
π(C ₁ -C ₆)	π*(C2-C3)	23.38	0.22	0.065
π(C ₁ -C ₆)	$\pi^{*}(C_{4}-C_{5})$	22.95	0.30	0.075
σ(C ₁ -O ₃₀)	σ*(C ₇ -H ₉)	1657.20	0.03	0.192
σ(C ₁ -O ₃₀)	σ*(C ₁₂ -C ₁₉)	1422.18	0.03	0.192
$\sigma(C_3-C_4)$	π*(C ₁₉ -C ₂₀)	3731.85	o.59	1.451
π(C ₄ -C ₅)	π*(C ₂ -C ₃)	29.89	0.20	0.070
π(C ₄ -C ₅)	π*(C ₁₂ -O ₁₃₎	33.40	0.10	0.055
π(C ₁₉ -C ₂₀)	$\sigma^{*}(C_{5}-C_{14})$	269.28	0.57	0.385
π(C ₁₉ -C ₂₀)	σ*(C ₇ -H ₁₀)	84.75	0.82	0.259
π(C ₁₉ -C ₂₀)	π*(C ₁₂ -O ₁₃)	213.41	0.46	0.293
π(C ₁₉ -C ₂₀))	σ*(C ₁₄ -H ₁₅)	231.89	0.65	0.382
π(C ₁₉ -C ₂₀))	σ*(C ₁₄ -H ₁₇)	1148.22	0.24	0.512
π(C ₁₉ -C ₂₀))	σ*(C ₃₁ -H ₃₂)	108.98	0.72	0.275
π(C ₁₉ -C ₂₀))	σ*(C ₃₁ -H ₃₃)	103.63	0.74	0.271
π(C ₁₉ -C ₂₀))	σ*(C ₃₁ -C ₃₄)	518.18	0.37	0.422
LP(2)O ₃₅	σ*(C ₃₁ -C ₃₄)	296.83	0.05	0.111
LP(2)O ₃₅	σ*(C ₃₄ -O ₃₆)	30.98	0.65	0.129
LP(1)O ₁₃	σ*(C ₁₄ -H ₁₆)	99.84	0.23	0.136
LP(2)O ₃₆	π*(C ₃₄ -O ₃₅)	43.91	0.36	0.112

 Table 2: Second order perturbation theory analysis of Fock matrix in NBO basis for EBDA.





normal modes are shown in Figures 2 and 3.

Phenyl ring vibrations: The hetero aromatic structure shows the presence of C-H stretching vibration in the region of 3100-3000 cm⁻¹ which is the characteristic region for the ready identification of C-H stretching vibration [39]. In this region, the bands are not affected appreciably by the nature of the substituent. The C-H stretching modes usually appear with strong Raman intensity and are highly polarized. Owing to this high polarization, Raman bands have not been observed in the experimental spectra. The observed weak band in IR at 3086 cm⁻¹ and in Raman at 3106 and 3083 cm⁻¹ are assigned to the C-H stretching mode of ring 1. The same mode observed in IR at 3060 and 3045 cm⁻¹ and in Raman at 3072 and 3052 cm⁻¹ is assigned for ring 2 (Figures 2 and 3) (Table 3).

PED corresponding to this vibration is a pure mode as it is evident from the contribution of ~ 100%. The C-H stretching anharmonic wave number lies between 3100 and 3000 cm⁻¹. The C-H in-plane bending frequencies appear in the range of 1000-1300 cm⁻¹ and are very useful for characterization purposes [39]. The observed series of strong bands in IR at 1310, 1280, 1258, 1200, 1157 cm⁻¹ is assigned to the C-H inplane bending vibration of phenyl ring 2. The observed strong bands at 1070, 1014 cm⁻¹ in IR are assigned to the C-H in plane bending vibration of phenyl ring 1. The same mode of vibration of ring 2 is observed in Raman at 1152 cm⁻¹ as a medium band. PED confirms that these vibrations are of mixed mode as is evident from the table.

The C-H out of plane bending vibrations is strongly coupled vibrations and occurs in the region of 1000-750 cm⁻¹ [40,41]. The aromatic C-H out-of-plane bending vibrations of EBDA are assigned to the bands observed at 898 and 803 cm⁻¹ in the FT-IR spectrum which is well correlated with the B3LYP method as shown in table with a PED contribution of ~ 70-90%. Also the bands in the FT-IR spectrum, at 899, 764 cm⁻¹ and in FT-Raman spectrum at 881 and 805 cm⁻¹ are assigned to C-H stretching vibrations. Benzene ring possesses six stretching vibrations, of which four occur near the highest wave number 1600, 1580, 1490 and 1440 cm⁻¹. In the absence of ring conjugation, the band near 1580 cm⁻¹ is usually weaker than that near 1600 cm⁻¹. The fifth ring stretching vibration is active near 1335 ± 35 cm⁻¹, the region which overlaps strongly with that of the C-H in-plane deformation and the intensity is in general, low, medium or high [41].

The sixth ring stretching vibration or ring breathing mode appears as a weak band near 1000 cm⁻¹ in mono, 1,3-di and 1,3,5-trisubstituted benzenes. For substituted benzene, however, this vibration is substituent sensitive and is difficult to distinguish from the ring inplane deformation. In EBDA the C-C stretching vibrations are observed in the FT-IR and FT-Raman spectra at 1550, 1558, 1500 cm⁻¹ as a medium band and at 1555, 1501 cm⁻¹ as a weak band, which correlate well with the predicted theoretical data. The PED corresponding to all C-C vibrations lies between 40% and 50% as shown in Table 3 with a combination of C-H in-plane bending in this region. The in-plane deformation vibration is at higher wave numbers than the out-ofplane vibrations [41] gave the frequency data for these vibrations for different benzene derivatives as a result of normal coordinate analysis. The theoretically computed C-C-C in-plane and out-of-plane bending vibration by the B3LYP/6-31G (d, p) method shows good agreement with the recorded spectral data.

Carbonyl group vibrations: The carbonyl group stretching vibrations give rise to the characteristic bands in IR and Raman. The intensity of these bands can increase because of the formation of hydrogen bonds. The carbonyl group vibration is observed in

the region 1760-1730 cm⁻¹ [42,43]. The strong band at 1750 cm⁻¹ is assigned to carbonyl C_{34} - O_{35} stretching mode. The C_{12} - O_{13} stretching mode is observed in IR at 1640 cm⁻¹ as a medium band and the C_{12} - O_{13} stretching vibrations are lower from the normal value. The red shifting of carbonyl stretching mode is attributed to the fact that the carbonyl group chelate with the other nucleophilic group, thereby forming both intra- and inter-molecular hydrogen bonding in the crystal. The C_{12} - O_{13} stretching is lowered due to the formation of C-H...O hydrogen bonding in the molecule.

Methyl group vibration: Methyl group vibrations are generally referred to as electron donating substituent in the aromatic ring system, the asymmetric and symmetric C-H stretching mode of CH, are expected to occur around 2980cm⁻¹ and 2870 cm⁻¹ [44]. The medium band observed in IR at 3005 cm⁻¹ is assigned to the symmetric stretching of Me1. The same vibration is observed in IR and in Raman at 2986 cm⁻¹ as a weak band and is assigned for Me2. The Me3 asymmetric stretching vibration is observed in IR at 2960 cm⁻¹ and in Raman at 2968 cm⁻¹ as a weak band. The shifting of the methyl (Me1 and Me2) asymmetric stretching wave number is due to the influence of the electronic effect resulting from the hyper conjugation and induction of methyl group in the aromatic ring [45]. Hyper conjugation causes the interaction of the orbital of the methyl group with the π orbital of an aromatic ring system [46]. The Methyl Symmetric bending of Me1 is observed at 1380 cm⁻¹ in IR as a medium band. This characteristic wave number is in close agreement with those reported for the similar compounds [46]. The band observed at 854 cm⁻¹ in IR is CH₃ out of plane bending of Me1. The observed band in IR at 764 cm⁻¹ and in Raman at 769 cm⁻¹ is assigned to the CH₃ out of plane bending modes of Me3.

Methylene group vibration: The methylene group vibrations are assigned on the basis of the spectral similarity to the related amino acid compounds. In amino acids, the CH₂ asymmetric and symmetric vibrations are expected to occur in the regions 3100-3000 and 3000-2900 cm⁻¹ respectively [47-49]. The observed weak band in Raman at 3023 cm⁻¹ is assigned to asymmetric stretching of C_{37} -H₂. The C_{31} -H₂ symmetric stretching is observed as a weak band in Raman at 2985 cm⁻¹. Electronic effects including back-donation, mainly caused by the presence of nitrogen atom adjacent to methylene groups, which can shift the position and alter the intensity of C-H stretching and bending modes. The scissoring mode of the CH, group gives rise to characteristic band near 1420-1412 cm⁻¹. The scissoring mode of the C₃₁-H₂ group is observed in IR at 1440 cm⁻¹ as a medium band. The rocking and wagging vibrations appear in the region of 1309-1288 cm⁻¹ and 910-905 cm⁻¹ respectively. The observed CH₂ deformations are given in Table 3.

HOMO-LUMO energy gap

Spatial distribution of molecular orbital's, especially those of highest occupied molecular orbital and lowest available unoccupied molecular orbital, are excellent indicators of electron transport in molecular systems. Conjugated molecules are generally characterized by a small HOMO-LUMO separation, which is the result of a significant degree of intra molecular change transfer (ICT) from the end-capping electron donor groups to the efficient electron acceptor groups through conjugated path. In EBDA, the HOMO-LUMO energy gap is 4.4913 eV and the lowering of this gap is essentially a consequence of the large stabilization of the LUMO due to the strong electron-accepting ability of the electron-acceptor group. The HOMO and LUMO orbital's are shown in Figure 4. In EBDA the atomic π -orbital's point towards each other and have better overlap, an increase in π -character points the fact that sigma bonds are stronger as evidenced by NBO analysis. A highly delocalized LUMO indicates that the electrons can more readily move

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Cal	Evn ID	Eyn Domon	TED (%)
Freq.	Exp IR	Exp. Raman	TED (%)
3091	-	3106 w	u _{с-н} ar in ring 1 (88)
3087	3086 w	3083 w	u _{с-н} ar in ring 1 (88)
3086	-	-	U _{C-H} ar in ring 2 (45)
3078	-	-	u _{с-н} ar in rin(88)g 1(83)
3073	-	3072 w	U _{CH} ar in ring 2 (96)
3068	3060 w	_	U_{C-H} ar in ring 1 (82)
3058	3045 w	3052 w	v_{CH} ar in ring 1 (87)
3015	-	3023 w	C_{37} -H ₂ Asym Stretching (96)
3006		3005 w	Me1 Asym. Stretching (90)
3006		0000 W	Me1 Asym. Stretching (96)
	-	-	, , ,
3002	-	-	Me3 Asym. Stretching (95)
2997	-	-	$C_{31}H_2$ Sym. Stretching (97)
2989	-	2986 w	Me2 Asym. Stretching (95)
2979	2960 w	2968 w	Me3 Asym. Stretching (99)
2955	-	2952 w	C ₃₁₋ H ₂ Sym. Stretching (98)
2946	-	-	C ₃₁ H ₂ Asym Stretching C31H2 Sym. Stretching (97)
2939	-	-	Me2 Sym. Stretching C31H2 Sym. Stretching (98)
2938	-		Me3 Sym. Stretching C31H2 Sym. Stretching (99)
2931	2930 w	2930 w	Me1 Sym. Stretching (84)
2909	-	2895 w	C ₃₁₋ H ₂ Sym Stretching (98)
1764	1750 s	-	C ₃₄ -O ₃₅ Stretching (90)
1647	1640 m	-	C ₁₂ -O ₁₃ Stretching (85)
1589	1600 m	-	v_{c-c} ar in ring 2 (45)
1582	1580 m	_	v_{c-c} ar in ring 1 (67)
1563	1558 m	1555 w	v_{c-c} ar in ring 1 (47)
1541	-	-	u_{c-c} ar in ring 2 (57)
1483	1500 m	1501 w	v_{c-c} ar in ring 2 (49)
	1300 11	1301 W	
1469	-	-	$C_{37}H_2$ Sissoring (86)
1467	-	-	u_{c-c} ar in ring 2 (71)
1449	-	-	Me1, Me2 Asym.Bending (67)
1448	-	-	Me3 Asym.Bending (63)
1439	1440 m	-	C ₃₁₋ H ₂ Sissoring (72)
1438	-	-	Me3 Asym. Bending (61)
1435	-	-	C ₃₁₋ H ₂ Sissoring (66)
1428	-	-	Me1, Me2 Asym. Bending (77)
1428	-	-	Me1, Me2 Asym. Bending (64)
1426	-	-	$\delta_{_{C-H}}$ ar in ring 1 (20)
1385	-	-	C ₃₇₋ H ₂ Wagging (28)
1377	1380 m	-	Me1 Sym. Bending (60)
1373	-	-	Me2 Sym. Bending (75)
1369	-	-	C ₃₁ H ₂ Wagging (33)
1359	-	-	Me1 Sym. Bending (23)
1348	-	-	δ_{c-H} ar in ring 2 (65)
1306	1310 s	_	δ_{c-H} ar in ring 2 (75)
1303	-		u_{c-c} ar in ring 1 (26)
1288	- 1280 s	-	δ_{c-H} ar in ring 2 (63)
	1200 5	-	
1265	-		$C_{37}H_2$ Twisting (49)
1256	1258 s	-	δ _{c-H} ar in ring 2 (76)
1243	-	-	δ _{c-H} in ring 1 (69)
1236	-	-	C ₃₁₋ H ₂ Twisting (42)
1218	1200 vs	-	δ _{c-H} ar in ring 2 (88)
1171	-	-	$C_{31}H_2$ Rocking (38)
1170			δ_{c-H} ar in ring 1 (51)

1162	-	-	δ _{c-H} ar in ring 1 (46)
1149	1157 vs	1152 m	$\delta_{_{C-H}}$ ar in ring 2 (41)
1143	-	-	δ _{c-н} ar in ring 2 (79)
1137	1120 vvs	1132 m	C ₃₇₋ H ₂ Twisting (76)
1098	-	1100 m	C ₃₇₋ H ₂ Rocking (70)
1092	-	-	δ _{c-H} ar in ring 1 (40)
1067	1070 s	-	δ _{c-H} ar in ring 1 (71)
1041	_	_	
10-11			C ₅ -C ₁₄ Stretch (28)
1028	-	-	Me1 Bending (77)
1019	-	-	Me2 Bending (61)
1013	1014 s		δ _{c-H} ar in ring 1 (42)
1011	-	-	Me2 Bending (41)
1001	-	-	Me3 Bending (65)
1000		991 vvs	C ₃₇₋ H ₂ Rocking (66)
983	-	-	Radial skel. in ring 1 (36)
982	-	-	Me2 Bending (44)
975	977 m	-	δ _{c-H} ar in ring 1 (79)
962	-	-	δ _{c-H} ar in ring 1 (86)
923	-	-	$\delta_{C,H}$ ar in ring 2(52)
922	-	-	δ_{c-H} ar in ring 1 δ_{c-H} ar in ring 1 (66)
901	-	-	δ_{c-H} ar in ring 2 δ_{c-H} ar in ring 1 (50)
888	899 m	881 w	δ_{c-H} in ring 2 δ_{c-H} ar in ring 1 (36)
852 m	854 m	_	Me1 out of plane δ_{c-H} ar in ring 1 (76)
834 m	834 m	-	Me2 out of plane δ_{CH} ar in ring 1 (94)
828 w	_	_	$\delta_{C,H}$ ar in ring 2 (82)
807 w	_	805 w	δ_{c-H} in ring 1 (17)
791 w	_	-	δ_{c-H} ar in ring 2 (27)
787 w	764 w	769	Me3 out of plane Bending (88)
741 w	751 w	755 w	Radial skel. in ring 2 (29)
741 W	-		
699 w	_	-	б _{с.н} ar in ring 2 (37) Radial skel. in ring 2 (34)
695 w			
685 w	-	-	δ _{c-H} ar in ring 2 (38) OP skel. in ring 1 (54)
	686 vvw	684 w	
667 w 630	-	647 w 632 w	Radial skel. in ring 1 (42)
	637 w		OP skel. in ring 2 (25)
611	615 s	619 w	6b Radial skel. in ring 1
590	-	-	OP skel. in ring 2 (29)
586	-	-	OP skel. in ring 2 (25)
563	-	575 w	OP skel. in ring 2 (23)
508	516 w	543 w	OP skel.in ring 1 (26)
457	470 vw	466 w	C_2 - C_7 stretching (40)
450	445 vw	450 w	OP skel.in ring 1 (21)
436	-	-	OP skel.in ring 2 (53)
409	416 vw	-	OP skel.in ring 1 (58)
401	-	-	OP skel.in ring 1 (78)
384	-	-	OP skel.in ring 2 (30)
367	-	377 w	OP skel.in ring 1 (29)
340	-	347 w	C-C Bending (29)
318	-	-	Me1 Torsion (39)
267	-	-	Me2 Torsion (61)
255	-	-	Me3 Torsion (59)
252	-	248 w	Me2 Torsion (18)
247	-	-	OP skel.in ring 1 (43)
209	-	-	OP skel.in ring 1 (46)
186	-	186 w	OP skel.in ring 1 (48)
171	-	-	OP skel.in ring 2 (28)
150	_	158 w	C-O Bending (61)

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147	-	-	C-O Bending (66)
143	-	-	C-O out of plane Bending (72)
132		134 w	OP skel.in ring 2 (47)
120		119 w	OP skel.in ring 2 (59)
103			OP skel.in ring 1(71)
90		83 w	OP skel.in ring 2 (21)
64			C-O out of plane Bending (51)
59			C-O out of plane Bending (36)
53			OP skel.in ring 2 (51)
44			OP skel.in ring 2 (55)
32			OP skel.in ring 2 (50)
22			C-O out of plane Bending (76)
15			OP skel.in ring 2 (73)

 Table 3: Calculated and observed vibrational frequencies for EBDA and their tentative assignment.

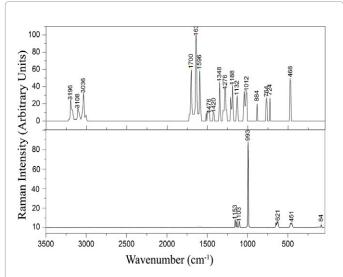
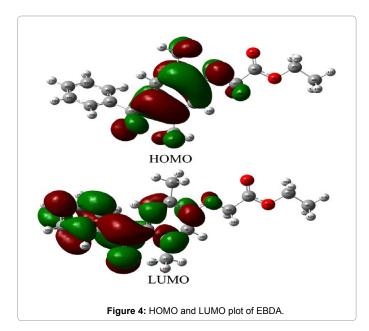
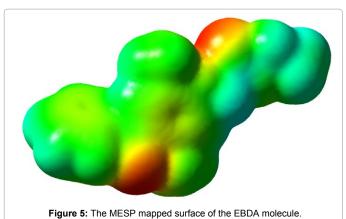


Figure 3: (a) Experimental and (b) simulated FT-Raman spectra of EBDA in the range 3500-50 $\mbox{cm}^{-1}.$





around the molecule from HOMO-LUMO and hence an improved ICT [50] (Figure 4).

In EBDA the LUMO strongly localizes on the phenyl ring indicating the presence of favourable atomic centres within EBDA for possible nucleophile attacks revealing its bioactivity and pharmaceutical activity. The Lowering of HOMO-LUMO energy gap confirms the possibility of ICT and HOMO-LUMO delocalization showing its possibility of nucleophile attacks which serves as an evidence for the pharmaceutical activity of the EBDA molecule. The HOMO-LUMO plot shows that both the HOMO and LUMO orbital's predominantly localize on the phenyl ring. The HOMO, LUMO and HOMO-LUMO energy gap of EBDA in the DFT level at 6-311++G (d, p) basis set has been calculated.

Molecular electrostatic potential

Molecular electrostatic potential is used primarily for predicting sites and relative reactivity's towards electrophilic attack, in studies of biological recognition and hydrogen bonding interactions [51]. To predict the reactive sites for electrophilic and nucleophilic attack for EBDA, the MEP at the B3LYP/6-311++G (d, p) method was calculated as shown in Figure 5. The different values of the electrostatic potential at the surface are represented by different colours. Electrostatic potential increases in the order red<orange<yellow<green
blue. Colour code of the maps is in the range between 0.0142 a.u. (deepest red) and 0.0142 a.u. (deepest blue) in title molecule, where blue colour indicates the strongest attractions and red indicates the strongest repulsion. The region of negative V(r) is associated with the lone pair of electrons (Figure 5).

As seen from Figure 5, in EBDA, the region having negative potential are over carbonyl group, the region having the positive potential are over the Phenyl ring and methyl group, the predominance of the light green region the MEPs surface corresponding to a potential halfway between the two extremes red and dark blue colour. The positive (blue) regions of MEP are related to electrophilic reactivity and the negative (red) regions to nucleophilic reactivity as shown in Figure 5. It can be seen from the MEP map of the present molecule that the negative regions are mainly localized on the O₁₃ and O₃₆ atoms. A maximum positive region is localized on the carbonyl carbon atom indicating a possible site for nucleophilic attack. MEP map shows that the negative potential sites are on electronegative atoms as well as the positive potential sites are around the carbonyl atom.

Conclusion

The detailed interpretation of the vibrational spectra has been

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carried out for the pharmaceutically important molecule Ethyl 2-(4-benzoyl-2, 5-imethylphenoxy) acetate. Molecular geometry, HOMO and LUMO energy and vibrational wave numbers of EBDA in the ground state have been calculated by using density functional theory. FT-IR and FT-Raman spectra of the EBDA have been recorded and analyzed. Observed and calculated wave numbers are found to be in good agreement. The lowering of carbonyl stretching wave number is due to the p-electron being localized. Lowering of the HOMO-LUMO energy gap value has substantial influence on the intra molecular charge transfer and pharmaceutical nature of the molecule.

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