EDITOR P. GERSHOM JEBARAJ

CRYSTAL GROWTH AND CHARACTERIZATION

Indian Association for Crystal Growth

Crystal Growth and Characterization

Indian Association for Crystal Growth



Editor

P. Gershom Jebaraj

Life Member

Indian Association for Crystal Growth (IACG)



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CHAPTER 5

A THEORETICAL INVESTIGATION OF BUMETANIDE USING CHARGE DENSITY AND TOPOLOGICAL PROPERTIES.

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5.1. INTRODUCTION

Hypertension or High Blood Pressure (HBP) is called a "silent killer" and it is a major cause of premature death worldwide and it affects between 16-37% of the population globally. Besides, 80% of all adults are at the risk from their blood pressure. A survey from the literature says that there are 234 million Indian adults are affected with hypertension which is expected to be the large increase in cardiovascular diseases burden in near future. HBP can lead to many serious health problems, such as heart attack, heart failure, stroke, and kidney disease. Treating high blood pressure early is important in preventing these problems. Moreover dozens of different medications can help to treat high blood pressure. These drugs are called anti-hypertensives. One such potent anti-hypertensive drug candidate is Bumetanide. Bumetanide (BMT) also known as Bumex is highly useful in the treatment of swelling and high blood pressure. This is a loop diuretic and works by decreasing the reabsorption of sodium by the kidneys [1,2]. It works by blocking the cation-chloride co-transporter NKCC1 (membrane transport protein, Na–K–Cl), which further decreases chloride concentration in neurons and therefore intensifying the inhibitory effect of GABA (γ-aminobutryic acid). GABA is known as a chief inhibitory neurotransmitter in the central nervous system. It is directly accountable for muscle tone. This mechanism of action helps Bumetanide to act as a potential anti-seizure drug [3].

This study about Bumetanide in the view point of structural, topological, reactivity and interaction properties will give rich information which further enables pharmacologists and drug engineers to design new drugs with reduced side effects and improved efficiency

5.2. THEORETICAL CALCULATIONS

5.2.1. Geometry Optimization

In this work geometry optimization of Bumetanide (C₁₇H₂₀N₂O₅S) was done at the ground state level using Becke-3-Lee-Yang-Parr hybrid exchange functional (B3LYP) of the density functional theory with 6311G(d, p) basis sets using GAUSSIAN 09W software [4].

Generally, geometry optimization tries to locate minima on the potential energy surface, and is a mathematical relationship connecting molecular structure and the resultant energy. The target molecule has 45 atoms and it has 129 degrees of freedom.

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5.3. RESULTS AND DISCUSSION

5.3.1. Structural Aspects

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The structure of the molecule Bumetanide has two rings connected by an oxygen atom. The optimized

The structure of the molecule Bumetanide has two rings connected by an oxygen atom. The optimized The structure of the molecule Bulletanide of the molecule Bulletanide of the structure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure of Bulletanide was shown in figure 5.1. The bond length range (1.373Å-1.540Å) of the Constructure structure of Bumetanide was shown in the structure of Bumetanide was shown in the composition of the composi bonds suggests that they all have single bonds with neighboring atoms which also has the value is found to be 1.373Å. The C3 atom has three bonds with neighboring atoms which also has the C2-C3. found to be 1.373A. The C3 atom has the C2-C3.

C4, C2-C3-O12 and C3-C4-O12 plane angles 119.99°, 120.01° and 119.99° respectively which show the C4, C2-C3-O12 and C3-C4-O12 plants are geometry for the central carbon atom. The bond length and bond sp2 hybridization and trigonal planar geometry for the central carbon atom. The bond length and bond sp2 hybridization and trigonal planar geometry for the central carbon atom. The bond length and bond sp2 hybridization and trigonal planar geometry for the central carbon atom. The bond length and bond sp2 hybridization and trigonal planar geometry for the central carbon atom. The bond length and bond sp2 hybridization and trigonal planar geometry for the central carbon atom. The bond length and bond sp2 hybridization and trigonal planar geometry for the central carbon atom. angle values of the molecule BMT has been given in Table 5.1 and 5.2 respectively.

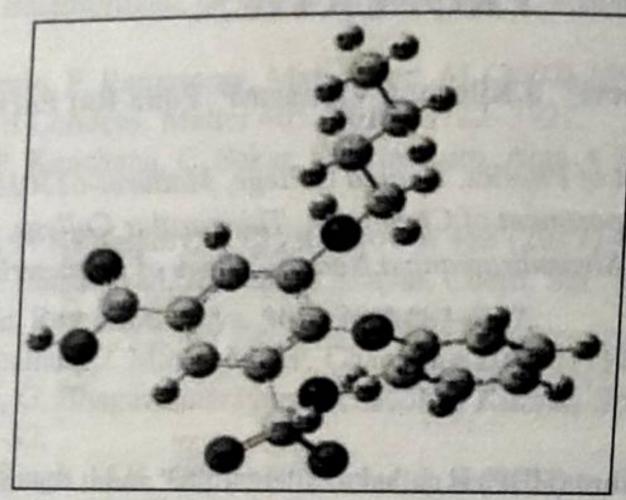


Figure 5.1 The Optimized structure of Bumetanide molecule.

Table 5.1 Bond Lengths of Bumetanide molecule

| Bonds | Bond Length (Å) | Bonds | Bond Length (Å) |
|---------|----------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------|
| C1-C2 | 1.395 | C16-H20 | 1.070 |
| C1-C6 | 1.395 | C17-H19 | 1.100 |
| C1-C7 | 1.100 | C33-H34 | |
| C2-C3 | 1.395 | C33-H35 | 1.070 |
| C2-C8 | 1.100 | C36-H37 | 1.070 |
| C3-C4 | 1.395 | C36-H38 | 1.070 |
| C4-C5 | 1.100 1.395 1.100 1.395 1.395 1.395 1.410 1.422 1.419 1.419 1.416 1.419 | The second secon | 1.070 |
| C5-C6 | 1.395 | C39-H40 | 1.070 |
| C13-C14 | 1.422 | C39-H41 | 1.070 |
| C13-C15 | 1.373 | C42-H43 | 1.070 |
| C14-C16 | 1.419 | C42-H44 | 1.070 |
| C15-C17 | | C42-H45 | 1.070 |
| C16-C18 | | C3-O12 | 1.430 |
| C17-C18 | 1.373 | C13-O12 | |
| C18-C21 | 1.540 | C21-O22 | 1.430 |
| C33-C36 | 1.540 | C21-O23 | 1.230 |
| C36-C39 | 1.540 | O23-H24 | 1.357 |
| C39-C42 | 1.540 | C14-S25 | 0.971 |
| | | S25-O26 | 1.780 |
| | | | 1.396 |

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| C15-N31 | 1.470 | S25-027 | |
|---------|-------|----------|-------|
| C33-N31 | 1.470 | S25-N28 | 1.396 |
| C4-H9 | 1.100 | N28-H29 | 1.710 |
| C5-H10 | 1.100 | N28-H30 | 1.000 |
| C6-H11 | 1.100 | N31-H32 | 1.000 |
| - | | 107-1132 | 1.000 |

Table 5.2 Bond angles of Bumetanide molecule

| Bonds | Bond angle (°) | Bonds | Bond angle (°) | Bonds | Bond angles (") |
|-------------|-------------------|-------------|-------------------|-------------|-----------------------|
| C1-C2-C3 | 120.01 | C6-C1-H7 | 120.00 | C3-O12-C13 | 109.50 |
| C1-C6-C5 | 120.00 | C6-C5-H10 | 119.98 | C4-C3-O12 | 119.99 |
| C2-C1-C6 | 120.00 | C14-C16-H20 | 120.47 | C13-C14-S25 | 120.40 |
| C2-C3-C4 | 119.99 | C15-C17-H19 | 118.98 | C14-S25-O26 | 110.10 |
| C3-C4-C5 | 119.99 | C18-C17-H19 | 120.65 | C14-S25-O27 | 110.10 |
| C4-C5-C6 | 120.00 | C18-C16-H20 | 120.47 | C14-S25-N28 | 93.80 |
| C13-C14-C16 | 119.20 | C33-C36-H37 | 109.47 | C15-N31-H32 | 109.47 |
| C13-C15-C17 | 120.32 | C33-C36-H38 | 109.47 | C15-N31-C33 | 109.47 |
| C14-C13-C15 | 120.50 | C36-C39-H40 | 109.47 | C16-C14-S25 | 120.40 |
| C14-C16-C18 | 119.06 | C36-C39-H41 | 109.47 | C18-C21-O22 | 130.07 |
| C15-C17-C18 | 120.37 | C39-C42-H43 | 109.47 | C18-C21-O23 | 112.29 |
| C16-C18-C17 | 120.55 | C39-C42-H44 | 109.47 | C21-O23-H24 | 110.61 |
| C16-C18-C21 | 118.59 | C39-C42-H45 | 109.47 | O12-C13-C14 | 118.64 |
| C17-C18-C21 | 120.86 | H34-C33-C36 | 109.47 | O12-C13-C15 | 120.86 |
| C33-C36-C39 | 109.47 | H35-C33-C36 | 109.47 | O22-C21-O23 | 117.63 |
| C36-C39-C42 | 109.47 | H37-C36-C39 | 109.47 | O26-S25-O27 | 119.60 |
| C13-C15-N31 | 120.69 | H38-C36-C39 | 109.47 | O26-S25-N28 | 110.10 |
| C17-C15-N31 | 118.99 | H40-C39-C42 | 109.47 | O27-S25-N28 | 110.10 |
| C1-C2-H8 | 119.98 | H41-C39-C42 | 109.47 | S25-N28-H29 | 109.47 |
| C1-C6-H11 | 120.01 | H34-C33-H35 | 109.47 | S25-N28-H30 | 109.47 |
| C2-C1-H7 | 120.00 | H37-C36-H38 | 109.47 | H29-N28-H30 | 109.47 |
| C3-C2-H8 | 120.01 | H40-C39-H41 | 109.47 | H32-N31-C33 | 109.47 |
| C3-C4-H9 | 119.98 | H43-C42-H44 | 109.47 | N31-C33-H34 | 109.47 |
| C4-C5-H10 | 120.01 | H43-C42-H45 | 109.47 | N31-C33-H35 | 109.47 |
| C5-C4-H9 | 120.01 | H44-C42-H45 | 109.47 | N31-C33-C36 | 109.47 |
| C5-C6-H11 | 119.99 | C2-C3-O12 | 120.01 | | |

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5.3.2. Intra and Inter Molecular Interactions

5.3.2. Intra and Inter Molecular interactions of the molecule with the neighbouring molecules can be explained. The intermolecular interactions of the molecule with the neighbouring molecules can be explained and intermolecular interactions of the molecule with the neighbouring molecules can be explained and intermolecular interactions of the molecule with the neighbouring molecules can be explained and intermolecular interactions of the molecule with the neighbouring molecules can be explained and intermolecular interactions of the molecule with the neighbouring molecules can be explained and intermolecular interactions of the molecule with the neighbouring molecules can be explained and intermolecular interactions of the molecule with the neighbouring molecules can be explained and intermolecular interactions of the molecule with the neighbouring molecules can be explained and intermolecular interactions of the molecule with the neighbouring molecules can be explained and intermolecular interactions. The intermolecular interactions of the distance from the Hirshfeld analysis and finger print plot [5, 6]. The distance from the Hirshfeld analysis and finger print plot [5, 6]. The distance from the Hirshfeld analysis and picturized by the Hirshfeld analysis and inside the surface are characterized by the quantities de and distance from the Hirshfeld analysis and picturized by the quantities de and distance from the Hirshfeld analysis and finger print plot [5, 6]. picturized by the Hirshfeld analysis and inside the surface are characterized by the quantities de and de a the nearest atoms outside and histor the 3D Hirshfeld plot and 2D finger print plot have been given in Figure normalized contact distance d_{norm}. The 3D Hirshfeld plot and 2D finger print plot have been given in Figure normalized contact distance d_{norm}. The 32 normalized contact distance d_{norm} in Figure 22 normalized contact d_{normalized} in Figure 22 normalized contact d_{normalized} in Figure 22 normalized co 2a and 2b. The large and deep red chedral points out the close contact interactions responsible for hydrogen bond contacts (O-H...O) and the light red spots on the surfaces. contact interactions responsible for higher strength of the O-H...O interaction is seen by the red colour spots. The represent C-H...π interactions. The strength of the O-H...O interaction is seen by the red colour spots. The represent C-H...π interactions. The state of the contacts and they play a major role which has the O...H/H...O inter molecular interest of the contribution of points between the spikes are from H...H contacts and they play a major role which has the contribution of 53.7% in the packing of the molecule in the crystal.

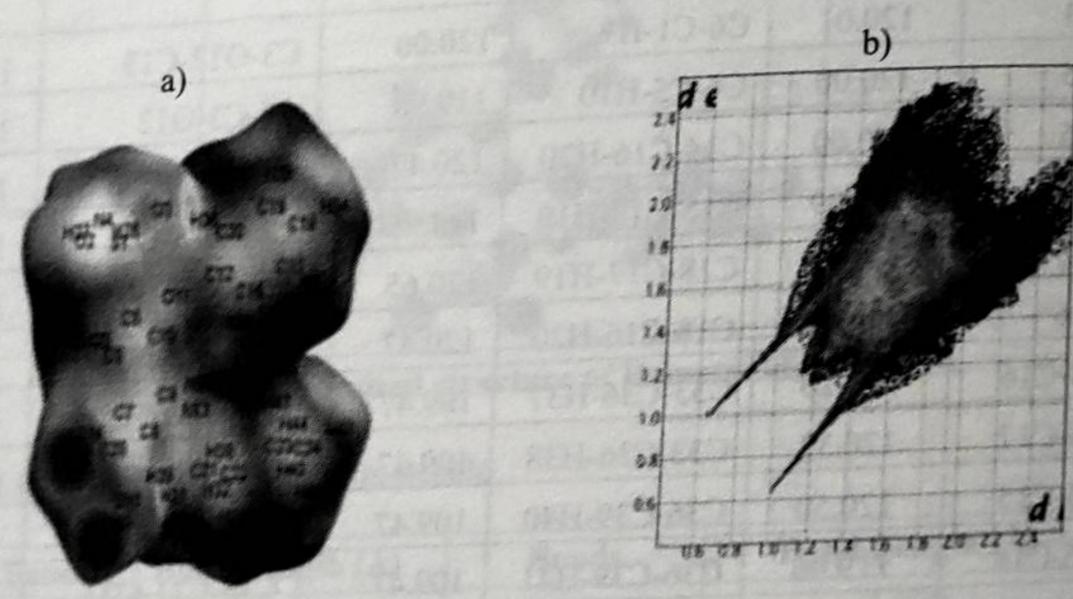
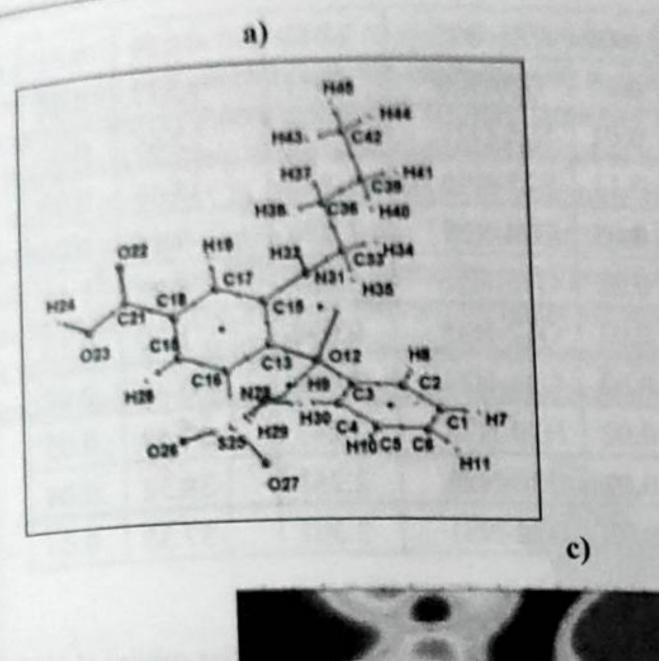


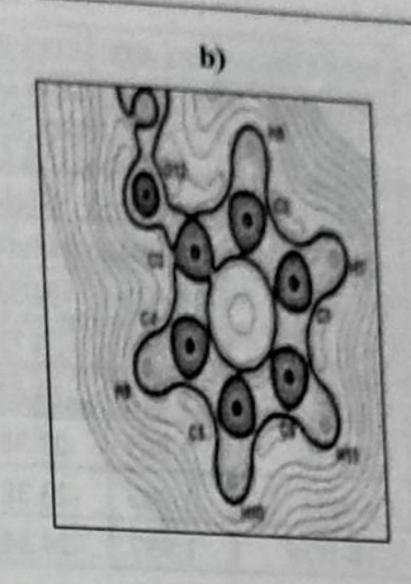
Figure 5.2 a) Hirshfeld plot and b) Finger print plot of the molecule.

5.3.3. Topological Properties

AIM topological analysis was carried out with the help of Multiwfn 3.3.9 [7]. Bader's AIM theory [8] is used as a great tool for the qualitative analysis of the electron density $\rho(r)$ and Laplacian $\nabla^2 \rho(r)$ of the electron density. Within the AIM theory, the topology of the electron density is exposed by its gradient vector field. The nucleus is known as an attractor and the paths converging to a nucleus represents a region of space known as atomic basin. Some gradient paths reaching a nucleus do not start from infinity but from a point in between two nuclei which is called a Bond Critical Point (BCP) where the gradient of p(r) vanishes.

The topological properties of the electron density are listed in table 5.3. The range of electron density for the C-C bonds 1.613 e/Å3 - 2.085e/Å3 in which the C1-C2, C1-C6, C5-C6, C14-C16 and C17-C18 bonds has the same electron density value of 2.085 e/Å3. The molecular graph showing the critical point between the atoms is shown in figure 5.3 a. The regions of charge concentration and charge depletion can be viewed through the positive and negative Laplacian of the electron density values. Charges are highly concentrated in the bonding region (Figure. 3b). Among all the bonds, the O12-N28 and O12-H35 bonds are having positive Laplacian values (0.96 e/Å5 and 1.20 e/Å5) which show the closed shell interaction of the bonds. The plot of electron localization function [9] clearly shows the sharing of electrons between the atoms in the bond path and also the lone pair details of the O12 atom (Figure.3c).





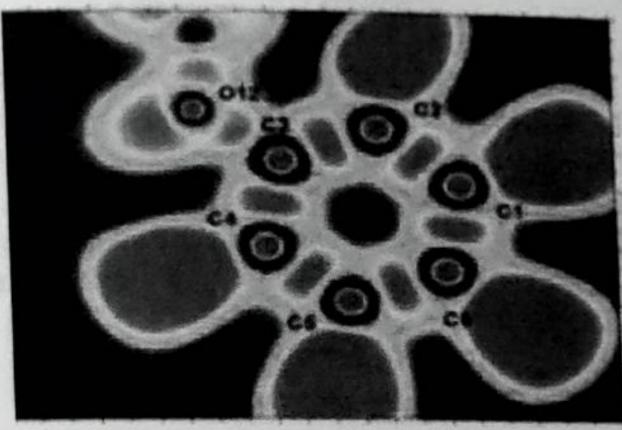


Figure 5.3: a) Molecular graph b) Laplacian of the electron density c) Electron localization function of the Bumetanide molecule.

Table 5.3 Topological properties of the Bumetanide molecule.

| Bonds | ρ (e/ų) | $\nabla^2 \rho$ (e/Å ⁵) | 8 | Bonds | ρ (e/ų) | $\nabla^2 \rho$ (e/Å ⁵) | 3 |
|---------|------------|-------------------------------------|------|---------|------------|-------------------------------------|------|
| C1-C2 | 2.085 | -20.72 | 0.21 | C17-H19 | 1.910 | -23.62 | 0.02 |
| C1-C6 | 2.085 | -20.72 | 0.2 | C33-H34 | 1.890 | -22.89 | 0.03 |
| C2-C3 | 2.112 | -21.45 | 0.25 | C33-H35 | 1.917 | -23.62 | 0.03 |
| C3-C4 | 2.112 | -21.21 | 0.24 | C36-H37 | 1.836 | -21.45 | 0 |
| C4-C5 | 2.078 | -20.48 | 0.21 | C36-H38 | 1.849 | -21.93 | 0 |
| C5-C6 | 2.085 | -20.72 | 0.2 | C39-H40 | 1.849 | -21.93 | 0.01 |
| C13-C14 | 2.078 | -20.48 | 0.28 | C39-H41 | 1.849 | -21.69 | 0.01 |
| C13-C15 | 2.031 | -19.76 | 0.27 | C42-H44 | 1.842 | -21.93 | 0.0 |
| C14-C16 | 2.085 | -20.48 | 0.21 | C42-H43 | 1.842 | -21.69 | 0.0 |
| C15-C17 | 2.065 | -20.24 | 0.22 | C42-H45 | 1.842 | -21.69 | 0.0 |
| C16-C18 | 2.072 | -20.48 | 0.2 | C3-O12 | 1.728 | -5.99 | 0.0 |
| C17-C18 | 2.085 | | 0.21 | C13-O12 | 1.856 | -9.16 | 0.0 |
| C17-C18 | 1.802 | | 0.13 | | 2.814 | -5.06 | 0.0 |

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| | The second second second second second | | | | | The second | |
|---------|----------------------------------------|--------|------|---------|-------|------------|-----|
| | 1.633 | -13.25 | 0.04 | C21-O23 | 2.018 | -12.29 | - |
| C33-C36 | 1.620 | -12.77 | 0.01 | S25-O26 | 1.971 | 22.17 | 0.0 |
| C36-C39 | And the second second | -12.77 | 0.01 | S25-O27 | 1.971 | 22.65 | 0.0 |
| C39-C42 | 1.613 | -21.45 | 0.11 | S25-N28 | 1.545 | -14.46 | 0.0 |
| C15-N31 | 2.038 | -15.18 | 0.03 | C14-S25 | 1.356 | -9.16 | 0.0 |
| C33-N31 | 1.701 | -23.38 | 0.02 | O12-N28 | 0.061 | 0.96 | 0.0 |
| C1-H7 | 1.903 | | 0.02 | O12-H35 | 0.094 | 1.20 | 1.4 |
| C2-H8 | 1.890 | -23.13 | 0.02 | O23-H24 | 2.429 | -60.49 | 0.1 |
| C4-H9 | 1.910 | -23.62 | 0.02 | H29-N28 | 2.247 | -37.59 | 0.0 |
| C5-H10 | 1.903 | -23.38 | | H30-N28 | 2.261 | -38.32 | 0.0 |
| C6-H11 | 1.903 | -23.38 | 0.02 | H32-N31 | 2.301 | -37.35 | 0.0 |
| C16-H20 | 1.937 | -24.34 | 0.02 | H32-N31 | 2.501 | -51.55 | 0.0 |

The highest occupied molecular orbital (HOMO) and lowest unoccupied molecular orbital (LUMO) are to the highest occupied molecular orbital (HOMO) represents the ability to donate an electrical control of the HOMO represents the ability to donate an electrical control of the HOMO represents the ability to donate an electrical control of the highest occupied molecular orbital (LUMO) are to the highest The highest occupied molecular orbitals. The HOMO represents the ability to donate an electron which has two important molecular orbitals. The HOMO represents the ability to accept an electron. The House two important molecular orbitals. The homo and Line character and the LUMO with π^* character has the capability to accept an electron. The HOMO and Line character and the LUMO with π^* character has the capability to accept an electron. The HOMO and Line character and the LUMO with π^* character has the capability to accept an electron. The HOMO and Line character and the LUMO with π^* character has the capability to accept an electron. The HOMO and Line character and the LUMO with π^* character has the capability to accept an electron. energies were calculated by the standard basis set B3LYP/6-311G++ (d, p) (5D, 7F) where H0M0 see found at -0.231eV and LUMO is found at -0.069eV. The DFT method predicts that the HOMO-LUMO energy gap of BMT is 0.087eV which is found to be very low and it leads to less stability of the molecule and pronounces the softness of the molecule.

Table 5.4: Reactivity properties of the Bumetanide molecule.

| Molecular descriptor | Energy (a.u.) HCTZ |
|------------------------------------------------|--------------------|
| Electron affinity | 0.069 |
| A=[-E _{LUMO}] Ionization potential | 0.231 |
| I=[-E _{HOMO}] Global hardness | 0.081 |
| η=(I-A)/2 softness | 12.392 |
| S=1/η Electronegativity | 0.150 |
| e=(I+A)/2 Electrophilicity index 0=μ²/2η | 0.139 |

On the basis of the density functional theory the ionization Potential, electron affinity, hardness softness, electronegativity and the electrophilicity index are calculated and tabulated in Table 5.4. The values help us to characterize the molecule that this molecule is chemically reactive, less toxic and selfmolecule.

The molecular electrostatic potential helps to identify the site and chemical reactivity of the molecule [10].

Nucleophilic regions (δ-)(prope to electrostic) Nucleophilic regions (δ-)(prone to electrophilic attack) are found near O atoms and electrophilic regions(δ-) (prone to nucleophilic attack) are found near H atoms and remaining other atoms. The electrostatic potential map of the molecule BMT is shown in Figure 4 map of the molecule BMT is shown in Figure.4.

This map evidently shows that all the oxygen atoms such as O12, O22, O23, O26 and O27 are electron rich sites where they all can interact with the electron poor regions in the target protein. It can be a good polar molecule as the value of dipole moment is slightly large (µ= 4.86Debye).

Population analysis is the unembellished model in order to describe charge distribution and also for deriving atomic charges. In the application of quantum chemical calculation to molecular systems, the Mulliken atomic charges play a vital role. The AIM charges are based upon the electron density values. The Among all the atoms, the S25 atoms bears the most positive charge (AIM: 2.806e, MPA: 1.081e) and the O26 atom through AIM (-1.270e) MPA(-0.47e) bears the most negative charge. The MPA and AIM charges are listed in table 5.5. All the N and O atoms bears negative charges in both the schemes and the schemes clearly predict that the O26 atom possess the more electron rich region among all the O atoms.

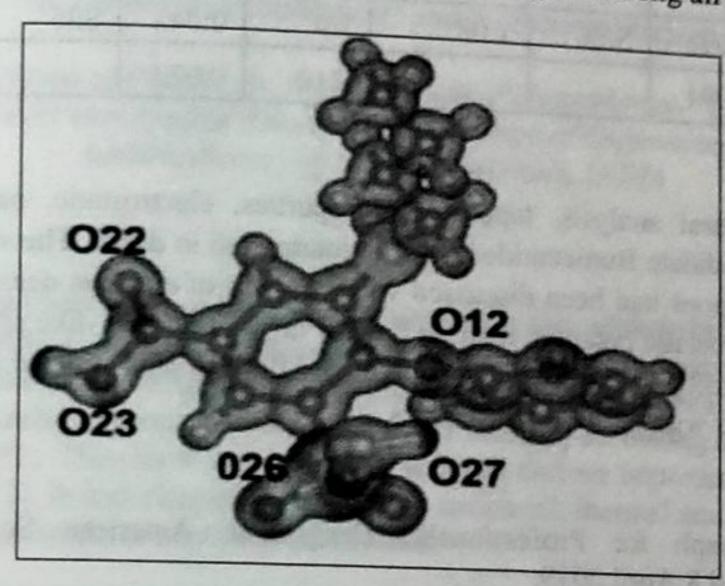


Figure 5.4: Molecular electrostatic potential map of Bumetanide molecule.

Table 5.5 MPA and AIM charges of Bumetanide molecule.

| Atoms | Q(e) MPA | Atoms | Q(e) MPA | Atoms | Q(e) AIM | Atoms | Q(e) AIM |
|-------|-------------|-------|-------------|-------|-------------|-------|-------------|
| C1 | -0.093 | H11 | 0.092 | C1 | -0.003 | H11 | 0.018 |
| C2 | -0.088 | H19 | 0.142 | C2 | 0.001 | H19 | 0.060 |
| C3 | 0.165 | H20 | 0.132 | C3 | 0.426 | H20 | 0.111 |
| C4 | -0.067 | H24 | 0.267 | C4 | 0.018 | H24 | 0.584 |
| C5 | -0.08 | H29 | 0.261 | C5 | 0.003 | H29 | 0.399 |
| C6 | -0.102 | H30 | 0.261 | C6 | -0.010 | H30 | 0.401 |
| C13 | 0.274 | H32 | 0.221 | C13 | 0.504 | H32 | 0.347 |
| C14 | -0.323 | H34 | 0.131 | C14 | -0.122 | H34 | 0.005 |
| C15 | 0.051 | H35 | 0.119 | C15 | 0.402 | H35 | 0.030 |
| C16 | 0.096 | H37 | 0.109 | C16 | 0.019 | H37 | -0.028 |
| C17 | 0.054 | H38 | 0.136 | C17 | -0.006 | H38 | -0.012 |
| C18 | -0.237 | H40 | 0.111 | C18 | 0.003 | H40 | -0.021 |
| C21 | 0.414 | H41 | 0.11 | C21 | 1.515 | H41 | -0.022 |

| C33 | -0.038 | H43 | 0.108 | C33 | 0.353 | H43 | 1 |
|-----|--------|-----|--------|-----|--------|-----|-------|
| C36 | -0.293 | H44 | 0.106 | C36 | 0.043 | H44 | -0.0 |
| C39 | -0.232 | H45 | 0.106 | C39 | 0.061 | H45 | -0.0 |
| C42 | -0.284 | 012 | -0.36 | C42 | 0.031 | 012 | -0.0 |
| N28 | -0.606 | O22 | -0.347 | N28 | -1.130 | 022 | 1-1.0 |
| N31 | -0.497 | O23 | -0.341 | N31 | -1.017 | 023 | 1-1.1 |
| H7 | 0.095 | 026 | -0.47 | H7 | 0.020 | 026 | -1.0 |
| H8 | 0.099 | O27 | -0.47 | H8 | 0.027 | 027 | -1.2 |
| Н9 | 0.093 | S25 | 1.081 | Н9 | 0.042 | S25 | -1.2 |
| H10 | 0.094 | | | H10 | 0.023 | | 2.80 |

5.4. CONCLUSION

In this study, the structural analysis, topological properties, electrostatic, nature, reactivity and interaction of the drug candidate Bumetanide had been interpreted in detail. The strength of the bonds, electron rich and poor regions had been discussed with the help of electron density, Laplacian of the electron density and electrostatic potential map. It has the favourable drug-like properties such as high chemical reactivity, very soft and less toxicity. This study will pave path way for the drug engineers to design more efficacy drugs with reduced side effects.

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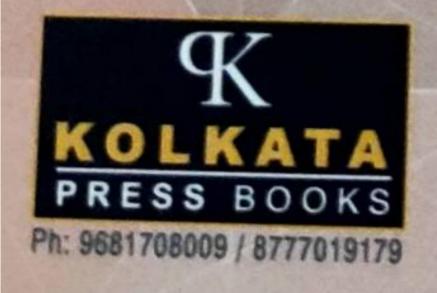


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