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Molecular Optimization Study of Gentamicin Drug

Abbas A-Ali Drea* and Marwa Kamil Jalil

University of Babylon, College of Science, department of Chemistry, Hilla-Iraq

Abstract: Molecular optimization have been carried out in vacuumed to investigate the energetic properties of geometry optimized structure of Gentamicin drug. Theoretical methods like, Semi empirical methods and Density functional method have been performed to estimate the chemical reactivity through calculations of Energy gap, Binding energy, Heat of formation and Zero point energy. The reactivity of the molecule has been investigated using Surface potential energy calculations through PM3-Single point Configuration Interaction (3*3), Microstate method to examination chemical bound stability of main bounds in Gentamicin drug molecule through bond angle, bond length and torsion angle. The vibration spectrum and electronic spectrum have been calculated by semi empirical CNDO method and DFT (Minimal STO-3G level). The best conformations of Gentamicin molecule has been studied through rotation of most reactive bound torsion angle C_{19} - N_{26} - C_{27} - H_{62} .

They found, the energetics values of total energy, MP2 Correlation Energy, E_{gap} and Zero point according to DFT(Minimal STO-3G level of theory) are equal to -298643.889, -1553.812, 1.37284, and -3763.39691 respectively at kCal/mol units. The N₂₆-C₁₉ bond in Gentamicin represented the most probable to break down and shared in various reactions than other duo to the lowest bond dissociation energy value (75.734 kCal/mol). The conformations has most stable energy through energetic value equal to -6835.4145 kCal/mol.

Key words: Gentamicin, potential energy surface, Geometry optimization, Single point calculations, DFT, Semi-empirical.

Introduction

Gentamicin is the antibiotic of first choice for treatment serious bacterial infections in bone, respiratory, skin, urinary tract, stomach, soft tissue, blood and heart in form of injection, ointment, cream, suspension and also used in Veterinary Medicine in many developed and industrialized countries[1-3]. Gentamicin has been discovered in 1963 by Marvin Weinstein's set at Schering Plough through isolated from various species of the micro mono sporia echinospora [4, 5]. Gentamicin is a commonly used as aminoglycoside antibiotics [6]. Gentamicin has been designed from three rings purpose amine, 2-deoxy strep amine and gentos amine or garos amine ring [7], since the major compounds of

^{*}Corresponding author: aadreab22@yahoo.com

Gentamicin are differ in purpose amine component. The four major compounds of complex mixture for Gentamicin are C, C1a and C2a, a 6'-C-epimer of C2 [8, 9]. The other minor compounds that present include Gentamicin's A,A1,A2,A3,C2a,C2b,A4,B,B1and X2, and JI-20A,JI-20B,Sisomicin and G-418 have been described which are other antibiotics structurally related to Gentamicin [10,11]. It was one of few antibiotics thermally-stable that remains active until after sterilization process [12]. Gentamicin drug fundamentally acts by binding to the 30S subunit of the bacterial ribosome stopping the continuous progress of protein synthesis [13]. The two hydroxyl groups of deoxystreptamine(2-DOS) linked to Garosamine (3-methyl amino-3-deoxy-4-C-methyl-β-L-arabinose) and purpurosamine (2,6-diamino-2,3,4,6,7_pentadeoxyheptopyranose) amino sugars by glycosides' bonds, the amino groups in the two surges attached to (2-DOS) profoundly influence on the biological activity but presence or absence of hydroxyl groups not have a clear effect on the activity of aminoglycosides antibiotics. Figure 1. Show two dimensional view of the Gentamicin molecule [14-16].



Figure 1: Two dimensional view of the Gentamicin molecule [14-16]

They found several types of Gentamicin with different chemical activities due their structures, that's related to energetics properties and energy gaps for each of these stractures. The aim of the present work tend to find the molecular optimized structure towards the chemical activity of Gentamicin. The quantum mechanics calculation methods have been used, since different methods of semi-empirical and DFT- STO-3G at level of theory, to estimate the geometry optimization, heat formation, total energy, HOMO&LUMO, Energy gap, vibration spectrum, and electronic spectrum. Also Surface potential energy was used to investigate the reactivity of main chemical bonds that's consistent gentamicin molecule structure [17].

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Computational Details

Computations have been done using quantum methods that's implemented at Hyperchem program version 8.0.2 [18-20]. DFT, Minimal STO-3G at level of theory have been used to find out Geometrical properties, likes electrostatic potential, total charge density, HOMO and LUMO energy, Zero point energy, vibration spectrum and electronic spectrum [21]. Semiempirical calculation have been performed using several methods like MNDO, CNDO, INDO, MNDO3, AM1 and RM1 and to estimate energetic properties, that's followed by using DFT, Minimal STO-3G at level of theory for the same manner[22]. Potential energy stability of bond length, bond angle and torsion angle have been examined using PM3 method CI (3*3) Microstate to estimate the chemical reactivity of main bound in Gentamicin drug[23].

Results and Discussion

The Geometry optimization structure of gentamicin molecule is represented in figure 1. They found that properties such electrostatic potential and their molecular orbitals are distributed at different active sites. The chemical structure is consisted from three different sugar rings and chemical reactivity of the substituted functional groups on these ring are differ at different orientations due their different induced field effect, such amine groups, hydroxyl groups, and methyl amine . Table 1. Involved energetic values of optimized gentamicin, the total energy value with full MP2 -300197.701 kCal mol⁻¹ is comes due geometry optimization process in vacuumed. Zero point energy (ZPE) value is calculated as a result of vibrational spectrum data that's equal to -3763.39691 kCal mol⁻¹ is much stabilized structure and relax molecules due to this low value of ZPE. The cost time of calculus is 542 hours, 54 mints and 31 second. The red color regions of electrostatic potential that observed in figure 1&2 show high electron density on oxygen atoms due to the high electronegativity, while the green color reigns represented the low electron density of carbon atoms. The HUMO and LUMO energies have been appeared in red and green color, the negative part of wave function has been given by red color (subject attacked by an electrophile) and the green color represented the positive part of wave function (subject attacked by nucleophile), this behavior determine chemical effectiveness of Gentamicin molecule toward the substitution reaction [24-27].

| Table 1 |
|---|
| Energetic properties of Gentamicin molecule calculated at DFT Minimal STO-3G at |
| level of theory |

| Total energy (kCal/mol) | -298643.889 |
|-----------------------------------|-------------|
| Total energy with MP2 (kCal/mol) | -300197.701 |
| MP2 Correlation Energy (kCal/mol) | -1553.812 |
| LUMO (eV) | 55.92773 |
| HOMO (eV) | 54.55486 |
| Energy gap (eV) | 1.37284 |
| Zero point energy (kCal/mol) | -3763.39691 |
| Time of calculus (hr.min.sec) | 542:54:31 |



Figure 2: Geometrical optimization properties of Gentamicin molecule calculated by DFT, Minimal STO-3G at level of theory

Surface Potential energy calculation have been used to investigated the stability of bond length and bond angle for the bounds of Gentamicin molecule through PM3 Configuration Interaction microstate (3*3), as shown in Table. They found that's bonds of C_{14} - O_{28} and C_{18} - O_{30} were more stable than other bonds with 108.354 and 104.548 kCal/mol respectively of dissociation energy values than other bounds. From other view Figure 2. Shows the four bond with lowest chemical stability than other chemical bounds of Gentamicin molecule, since the red color on C_{14} - O_{28} and C_{18} - O_{30} bonds appears their higher

stability than other two bonds C_{19} - N_{26} and C_{10} - C_{21} bonds in yellow color. The lowest chemical stability due to the induced effect field of substituted functional groups that found in the molecule (N atom have electronegativity less than O atom, leads to weak hydrogen linkage in the amines group compared with a hydroxyl group and thus easily broken and shared in important reactions).

| method CI (5 ⁻⁵) Microstate | | | | | | |
|---|------------------------|------------------------------|-------------------------|-------------------------------|-----------------------|--|
| Bonds | *Equilibrium energy | **Equilibrium bond length | *Dissociation energy | **Dissociation bond length | *Energy difference | |
| C ₂ -O ₁₂ | -6835.467 | 1.927 | -6751.963 | 2.326 | 83.504 | |
| $C_{7}^{2} - O_{12}^{13}$ | -6835.302 | 1.427 | -6751.080 | 2.826 | 84.222 | |
| $C_{5} - O_{20}^{15}$ | -6835.450 | 1.427 | -6748.511 | 2.627 | 86.939 | |
| $C_{17} - O_{20}^{20}$ | -6835.146 | 1.427 | -6737.786 | 2.327 | 97.36 | |
| $C_{21} - N_{23}^{20}$ | -6834.811 | 1.527 | -6753.967 | 2.627 | 80.844 | |
| $C_{4}^{2}-N_{22}^{2}$ | -6834.801 | 1.527 | -6754.999 | 2.627 | 79.802 | |
| C ₃ -N ₃₁ | -6834.985 | 1.527 | -6754.030 | 2.627 | 80.955 | |
| $C_{12} - N_{24}$ | -6834.914 | 1.527 | -6742.434 | 2.827 | 92.48 | |
| $C_{19}^{17} - N_{26}^{17}$ | -6834.991 | 1.527 | -6759.257 | 2.627 | 75.734 | |
| $C_{10} - C_{21}$ | -6835.109 | 1.527 | -6759.148 | 2.827 | 75.961 | |
| $C_{14}^{10} - O_{28}^{21}$ | -6835.449 | 1.427 | -6727.095 | 2.427 | 108.354 | |
| $C_{18}^{14} - O_{30}^{20}$ | -6835.181 | 1.427 | -6730.633 | 2.427 | 104.548 | |
| $C_{1}^{10} - O_{25}^{00}$ | -6835.192 | 1.427 | -6737.128 | 2.827 | 98.064 | |
| $C_{21} - C_{22}$ | -6835.439 | 1.527 | -6747.763 | 2.827 | 87.676 | |
| $C_{14}^{-1} - C_{29}^{-2}$ | -6835.394 | 1.527 | -6752.386 | 2.827 | 83.008 | |
| $N_{26} - C_{27}$ | -6834.412 | 1.527 | -6751.304 | 2.527 | 83.108 | |

 Table 2

 Surface Potential energy calculation of chemical bonds Semiemprical-PM3 method CI (3*3) Microstate

* kCal/mol units. ** Angstrom units



Figure 3: The lowest chemical stability bonds of Gentamicin molecule (active bonds)

Figure 4. Illustrates the potential energy stability of bonds at Gentamicin molecule. The C_2 - O_{13} bond is lowest stable and more active toward the reactions than other bond, since bond length equal to 1.927 ú. The difference in the stability of the bonds depends on the nature of substituted functional groups (tendency to donate or draw electrons) and also their steric effects of each bond in the bulk structure.



Figure 4: Potential energy stability curves of bond length in Gentamicin molecule calculated at semi emprical PM3 CI.Microstate (3*3)

Table 3. Show the calculation of bound angle stability toward the torsion energy stresses and their related equilibrium configuration as active site at Gentamicin structure. Comperes between all the chemical bounds described that, bond angle O_{13} - C_7 - C_{11} angle is more stable than other bond angles due their dissociation energy value of 615.711 kCal/mol [28].

| Bond angle | *Equilibrium | **Equilibrium | *Dissociation | **Broking | *Energy |
|---|--------------|---------------|---------------|------------|------------|
| | bound energy | bond angle | bond energy | bond angle | difference |
| C_{21} - C_{10} - C_{9} | -6834.4438 | 110 | -6807.3588 | 140 | 27.085 |
| C_{21} - C_{10} - C_{11} | -6834.2270 | 120 | -6806.6005 | 140 | 27.627 |
| N_{23} - C_{21} - C_{10} | -6835.4672 | 110 | -6800.2763 | 140 | 35.191 |
| C ₂₉ -C ₁₄ -C ₂₈ | -6835.3413 | 110 | -6790.5341 | 140 | 44.807 |
| N ₂₆ -C ₁₉ -C ₁₈ | -6834.5932 | 110 | -6803.9326 | 140 | 30.661 |
| N_{26} - C_{19} - C_{14} | -6834.7910 | 110 | -6805.7739 | 140 | 29.017 |
| C ₂₇ -N ₂₆ -C ₁₉ | -6834.9833 | 120 | -6822.2983 | 140 | 12.685 |
| O ₃₀ -C ₁₈ -C ₁₉ | -6835.3989 | 110 | -6803.3569 | 140 | 32.042 |
| O ₃₀ -C ₁₈ -C ₁₇ | -6834.8481 | 110 | -6810.2363 | 140 | 24.612 |
| N_{23} - C_{21} - C_{22} | -6834.7915 | 110 | -6789.6723 | 140 | 45.119 |
| C_{19} - C_{14} - C_{28} | -6835.1088 | 110 | -6803.7934 | 140 | 31.315 |
| C_{29} - C_{14} - C_{15} | -6835.4667 | 110 | -6801.0942 | 140 | 34.373 |
| N ₃₂ -C ₆ -C ₄ | -6834.6279 | 110 | -6813.6079 | 140 | 21.02 |
| N ₃₂ -C ₆ -C ₅ | -6835.4580 | 110 | -6802.7153 | 140 | 32.743 |
| $O_{25}-C_1-C_5$ | -6835.4526 | 110 | -6794.6425 | 140 | 40.810 |
| $O_{25}-C_1-C_2$ | -6833.9047 | 120 | -6810.5952 | 140 | 23.309 |
| N ₃₁ -C ₃ -C ₄ | -6835.3852 | 110 | -6796.5888 | 140 | 38.796 |
| N ₃₁ -C ₃ -C ₂ | -6835.4658 | 110 | -6804.6333 | 140 | 30.833 |
| N ₂₄ -C ₁₂ -C ₇ | -6835.4399 | 110 | -6804.6567 | 140 | 30.783 |
| C ₂ -O ₁₃ -C ₇ | -6834.901 | 120 | -6820.601 | 140 | 14.300 |
| C ₁₇ -O ₂₀ -C ₅ | -6834.779 | 110 | -6819.388 | 140 | 15.392 |
| C_{18} - O_{17} - C_{20} | -6835.323 | 110 | - 6806.133 | 140 | 29.19 |
| O_{20} - C_{17} - O_{16} | -6835.0009 | 100 | -6775.118 | 140 | 59.882 |
| O ₂₀ -C ₅ -C ₆ | - 6835.164 | 110 | -6787.224 | 140 | 47.94 |
| C ₃ -C ₂ -O ₁₃ | -6835.416 | 110 | -6730.334 | 140 | 105.082 |
| O ₁₃ -C ₂ -C ₁ | -6834.943 | 110 | -6783.069 | 140 | 51.875 |
| O ₁₃ -C ₇ -C ₁₁ | -6835.456 | 100 | -6219.745 | 130 | 615.711 |
| O ₁₃ -C ₇ -C ₁₂ | -6833.747 | 120 | -6811.620 | 140 | 22.127 |
| O_{20} - C_5 - C_1 | -6835466 | 110 | -6803 631 | 140 | 31 835 |

 Table 3

 Surface Potential energy investigation of Gentamicin bond angle calculated at semi empirical -PM3, CI (3*3) Microstate

Table 4 shows the Potential energy stability of the mean torsion angles in Gentamicin molecule. The bound angle C_2 - O_{13} - C_7 - C_{12} is represented more stable bound than other bounds due to their requirements for 3037.6617 kCal/mol to break down, also the dissociation energy value of the bound angle C_{19} - N_{26} - C_{27} - H_{62} is equal to 3.231 kCal/mol,

so that it's represented the lowest stable than other angles of gentamicin [29]. Figure 5 Show comparison between these two bonds, the high stable bond defined in red color and the lowest stable in yellow color.

| Bond Torsion angles | *Equilibrium energy | **Equilibrium Torsion angle | *Dissociation energy | **Broking Torsion angle | *Dissociation energy of angle |
|--|------------------------|--------------------------------|-------------------------|----------------------------|----------------------------------|
| C ₆ -C ₅ -O ₂₀ -C ₁₇ | -6834.7261 | 120 | -6736.3330 | 180 | 98.3931 |
| C ₁₈ -C ₁₇ -O ₂₀ -C ₅ | -6834.9556 | 130 | -6829.4888 | 180 | 5.4668 |
| $C_2 - O_{13} - C_7 - C_{12}$ | -6834.5577 | 90 | -3796.8960 | 180 | 3,037.6617 |
| C ₁ -C ₅ -O ₂₀ -C ₁₇ | -6831.0410 | 180 | -6752.6841 | 60 | 78.3569 |
| C ₇ -O ₁₃ -C ₂ -C ₁ | -6803.3691 | 180 | -6118.7456 | 100 | 684.6235 |
| C ₂₂ -C ₂₁ -C ₁₀ -C ₉ | -6834.4434 | 170 | -6829.2099 | 110 | 5.2335 |
| C ₃ -C ₂ -O ₁₃ -C ₇ | -6834.0347 | 110 | -6444.9834 | 180 | 389.0513 |
| C ₁₁ -C ₇ -O ₁₃ -C ₂ | -6821.1719 | 180 | -5192.0469 | 70 | 1,629.125 |
| O ₁₆ -C ₁₇ -O ₂₀ -C ₅ | -6826.2759 | 180 | -6761.5801 | 90 | 64.6958 |
| C ₂₇ -N ₂₆ -C ₁₉ -C ₁₈ | -6832.7802 | 140 | -6820.0845 | 180 | 71.2001 |
| N ₂₃ -C ₂₁ -C ₁₀ -C ₉ | -6834.4331 | 130 | -6824.4809 | 100 | 9.9522 |
| N ₂₃ -C ₂₁ -C ₁₀ -C ₁₁ | -6834.3843 | 160 | -6828.9585 | 110 | 5.4258 |
| C ₂₇ -N ₂₆ -C ₁₉ -C ₁₄ | -6834.3779 | 60 | -6828.5972 | 150 | 5.7807 |
| C ₂₂ -C ₂₁ -C ₁₀ -C ₁₁ | -6832.6406 | 60 | -6828.61377 | 180 | 4.02683 |
| C ₁₉ -N ₂₆ -C ₂₇ -H ₆₂ | -6835.4145 | 180 | -6832.1835 | 140 | 3.231 |
| C ₁₉ -N ₂₆ -C ₂₇ -H ₆₃ | -6833.7505 | 60 | -6829.4551 | 140 | 4.2954 |
| C ₁₉ -N ₂₆ -C ₂₇ -H ₆₄ | -6834.4927 | 60 | -6830.1133 | 140 | 4.3794 |
| 1 | 1 | 1 | 1 | 1 | 1 |

Table 4Potential energy stability and dissociation of Gentamicin Torsion angle by
Semiemprical-PM3 method CI(3*3)

The C_{19} - N_{26} - C_{27} - H_{62} torsion angle represented the more stable from other in Potential energy stability value by -6835.4145 kCal/mol in 180° due to the functional groups that found in the molecule. Different optimized structures (configurations) can be occurs at Gentamicin through different transitions torsion angle states into C_{19} - N_{26} - C_{27} - H_{62} angle. Figure 5 Show that the torsion angle of C_{19} - N_{26} - C_{27} - H_{62} in Gentamicin molecule includes the presence of several cases of branches transition to vibrate bond chemical angle in different energies for getting the best situation stabilizing steric optimum , it's have several stationary point such as A, C and E are minimal . Points such as B and D are maximum energetic states. Only structures at points A, C and E represented the stable conformations for this torsion angle. the lowest energy conformation was the anti-conformation in structure E, it was the most stable from other by -6835.4145 kCal/mol in 180° due to the molecule arranged its groups to adopt the alleviated torsional strain and reduced the electron repulsion in the torsion angle [31].

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Figure 5: Surface potential energy curve of torsion angle C₁₉-N₂₆-C₂₇-H₆₂ of Gentmicine

CONCLUSIONS

- The Energetic properties of Gentamicin molecule have been found by using several semi empirical methods and DFT calculations.
- The chemical reactivity of the molecule appeared the N₂₆-C₁₉ bond length the most probable to break down by using S.P.E calculations duo to the less value of bond dissociation energy (75.734kCal/mol)of functional group (methyl amine) at Garosamine.
- The bond angle O₁₃-C₇-C₁₁ angle is more stable than other bond angles due their dissociation energy value of 615.711 kCal/mol.
- The bound angle C₂-O₁₃-C₇-C₁₂ is more stable bound than other bounds due to their requirements for 3037.6617 kCal/mol to break down.
- The C₁₉-N₂₆-C₂₇-H₆₂ is the most stable angle than others in Gentamicin with Potential energy stability value of -6835.4145 kCal/mol in 180^ú.

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