

A Study on the Electronic and Structural Properties of C₁₂X₈ (X = C, B) and Their Interaction with Glycine with Potentially Drug Delivery Vessels

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ABSTRACT

In this paper, the structural properties of C₂₀ and C₁₂B₈ fullerene interacting with glycine based on three active sites of glycine and one C atom or one B atom in C₁₂B₈ were analyzed through the density functional theory. It was found out that the binding of glycine to C₁₂B₈ generated a complex. Our results were extremely relevant in order to identify the potential applications of functionalized C₁₂B₈ as drug delivery systems. Glycine preferred to interact with the C₁₂B₈ cage via its carbonyl oxygen (B=O) active site. B atoms were relatively favored in energy over the C atoms in the C₁₂B₈ – glycine while the stable ordering of three active sites on glycine molecule was =O site > –O site > –N site.

Keywords: C₁₂X₈; glycine active site; DFT; HOMO-LUMO gap

INTRODUCTION

Long after the initial proposal by Osawa in 1970 [1], and the mass spectroscopic detection by Curl, Kroto, and Smalley in 1985 [2], the synthesis of a macroscopic quantity of fullerenes by Krätschmer and Huffman in 1990 suddenly fueled the interest of experimentalists in fullerenes [3]. One of the future applications that came immediately to mind was their potential use in biology.

Then a variety of biocompatible fullerene-related materials have been widely considered in nanotechnology [4] and biomedical fields [5, 6].

Up to now much interest in their applications in electronics, materials science, chemistry and biochemistry as

well as their unique physical properties such as electrical conductance, ferroelectricity, nonlinear optical properties, and so forth had been noted [7, 8] that could revolutionize industries. Substitutional doping fullerenes in which one or more carbon atoms replaced by heteroatoms like nitrogen, boron, germanium, and silicon, have been the subject of numerous experimental and theoretical investigations [9 - 13] opening a new window into the field of multicomponent systems.

Nitrogen- and boron-doped fullerenes were special important. Since B-doped fullerenes behave as positive hole carriers and N-doped fullerenes act as electron

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carriers. For example, Chen *et al.* further studied heterofullerene structures $C_{60-n}N_n$ for $n = 2, 4, 6$ and 8 using MNDO, AM1, PM3, and the Hartee-Fock method.

Manaa [10] and Bryant *et al.* [11] have studied the structural stabilities of $C_{48}N_{12}$. Jindal *et al.* [12] reported the results of *ab initio* calculations of structural, electronic and, vibrational properties for nitrogen-doped fullerenes $C_{60-n}N_n$, $n = 1-12$.

Beside C_{60} , the smallest possible fullerene cage, i.e., C_{20} with 12 pentagons and no hexagons [14], can be doped with these methods. C_{20} is somehow different from C_{60} because of its extreme curvature and reactivity. This curvature leads to a higher ratio of sp^3 to sp^2 bonding. Theoretical analysis shows that there are two kinds of carbon atoms in C_{20} . Atoms with the coordination number 4 (12 atoms) are sp^3 hybridized, but is very distorted compared to the ideal sp^3 hybridization in diamond. Atoms with coordination number 3 (8 atoms) are sp^2 hybridized.

The ideal goal of substitutional doping of fullerene cage is to improve its characteristics for special purposes while keeping its structural stability. In one of our previous work we were focusing our calculations on $C_{12}X_8$ heterofullerenes where $X = B, Al, Ga, C, Si, Ge, N, P$, and As .

Results from molecular dynamics simulation, showed that it is dynamically stable up to 1000 K [15]. Tian *et al.* [16] theoretically found that $C_{12}N_8$ is dynamically stable at ambient pressure.

Experimental studies demonstrate that the functionalization of C_{60} amino acids has the potential to provide greater interaction between the fullerene and a biological environment which is leading to novel medical applications [17, 18]. This has had further implications in the synthesis and biological activity of fullerene-containing bio molecules [19-22].

There are some theoretical studies using DFT and semi-empirical (AM1) methods about C_{60} -glycine [23-25] and $C_{59}B$ -glycine [26] and the applications in their biological affinity have been experimentally observed.

However the need for further study was crucial and relevant. The focus of the current research investigation was on the interaction of $C_{12}X_8$ ($X = C$ and B) with amino acid species. Fullerene-functionalized derivatives were used in pharmacology. For example in neuroprotective effect, HIV-protease inhibitors [27] and antioxidant [28].

Bio molecules delivery introduces several chemical and biochemical problems concerning the interactions between the bio molecules and the fullerene or nanostructure. There were numerous studies on this problem, either experimental [29] or theoretical [30] Jiang and Zheng [31], in 2005, synthesized the fullerene-glycine derivative which displays better antitumor activity *in vitro* against bone tumor cells.

These previous works allow us to make the assertion that fullerenes should interact with amino acids and act as potential drug delivery vessels.

The present study aims to investigate the stable binding sites between the C_{20} fullerene and $C_{12}B_8$ heterofullerenes (here after named as $C_{12}X_8$ ($X = C, B$)) with glycine based on three active sites of glycine and one X atom in $C_{12}X_8$ ($X = C, B$) (shown in Fig. 1 for C_{20} , $C_{12}B_8$ and glycine). The C_{20} cage, which consists solely by pentagons, is the smallest and unconventional fullerene [32-34]; it might generate the larger vibronic coupling than C_{60} cage. In the continuation of our interest in fullerene chemistry, it was very interesting for us to explore more detailed insights into the structural, and electronic properties of the interacting molecules

with the aid of quantum chemical calculations.

COMPUTATIONAL METHODS

It is well known that a glycine molecule has three active sites, the amino nitrogen (N), the hydroxyl oxygen (OH) and the carbonyl oxygen (O) sites. The $C_{12}X_8$ ($X=C, B$) has two active sites, being C atom and B atom. As shown in Figure 2, six $C_{12}X_8$ – glycine conformers, were generated from the $C_{12}X_8$ cage and glycine at different binding sites. Full geometry optimizations were accomplished by means of hybrid functional B3LYP and the 6-31+G* basis set, as implemented in Gaussian 98 [35]. DFT methods were generally flawed when discussing dispersion forces, the methods employed appear to partially account for this. The strength of the dispersion forces for simple van der Waals complexes were adequately computed by the B3LYP method. The applied basis set was composed of Pople's well-known 6-31G* basis set and an extra plus due to the importance of diffuse functions.

This value was a measure of the stability of the complex however our calculations show that as this value increases entropy values decrease as a result of increased affinity for the fullerene surfaces.

Vibrational frequencies were calculated for $C_{12}B_8$ to establish the nature of stationary points as true minima. If the vibrational frequencies were positive then the structures calculated were minimum structures.

RESULTS AND DISCUSSION

The smallest fullerene that satisfies Euler's theorem is C_{20} . It is highly strained due to the extreme pyramidalization of the C=C double bonds. Cao et al. studied the stable binding sites between the C_{20} fullerene and glycine based on three active sites of

glycine and an active top site and plane for C_{20} . Furthermore they considered and explored endohedral metallo fullerenes $Gd@C_{20}$ – glycine [36].

Since The $Gd@C_{20}$ -glycine is also carried out as a hypothetical system, not available, our interesting was exploration the structures, stabilities, and electronic properties of $C_{12}B_8$ -glycine with the aid of quantum chemical calculations. Although proteins are much more complicated than glycine, however, all proteins contain amino nitrogen (–N), hydroxyl oxygen (–O), and carbonyl oxygen (=O) active sites. The equilibrium geometries for the C_{20} , $C_{12}B_8$ and isolated glycine molecule were as shown in Figure1. Therefore, from the calculation results involving in this paper, one can predict that proteins might form bindings with C_{20} . The calculated harmonic vibrational frequencies at B3LYP/6-31+G* level confirm that the optimized $C_{12}B_8$ and all of the complexes were true minima (NIMAG = 0).

All C–B bonds of $C_{12}B_8$ showed the similar length of 1.588 Å. The C–B bond lengths of $C_{12}B_8$ systems are quite close to the sum of covalent radii of C and B atoms. The covalent radius of B is about 0.84 Å. C–B–C angles were narrower than normal 120.0 (in a range of 104.8–109.2). C=C double bonds were expectedly shorter than those of C_{20} (1.394 vs. 1.445 Å, Table 1).

The binding energy of $C_{12}B_8$ was 5.82 eV/atom which was not so far from that of C_{20} (6.34 eV/atom). Accordingly, the computed ν_{\min} of $C_{12}B_8$ was 317 cm^{-1} which was noticeably high (compare to 32 cm^{-1} of C_{20}). The fullerene used initially had C_i symmetry but upon complexation it became slightly distorted. For stable complexes, the N – C bond length of glycine was lengthened from 1.447 Å to 1.460 Å. As well at the same time, the C=O and C–O bond lengths slightly increase, 0.02 Å. It means the binding

between glycine and $C_{12}X_8$ also changed the structure of glycine.

As shown earlier, glycine and $C_{12}X_8$ can form complexes by forming a new bond by breaking one of the original bonds of glycine. The new bond formation can increase the stability of the complexes, whereas the breaking of the original bond of glycine can decrease their stability. To evaluate the stability of $C_{12}X_8$ -glycine complexes, we calculated the energy of formation of a complex between the glycine and $C_{12}X_8$ molecules (glycine + $C_{12}X_8 = C_{12}X_8$ - glycine), by using the equation:

$$E_b = E_{gly-C_{12}X_8} - E_{gly} - E_{C_{12}X_8} \quad (1)$$

where E_{gly} , $E_{C_{12}X_8}$ and $E_{gly-C_{12}X_8}$ were the energy values of glycine, $C_{12}X_8$ and glycine- $C_{12}X_8$ complex, respectively. The results were listed as shown in Table 1. A negative E_b shows a thermodynamic

stability; and the more negative the value of E_b , the more stable the complex. One can see that the formations of all six complexes were endothermic by 4.06 to 22.00 eV. This indicates that the binding of glycine to $C_{12}B_8$ was slightly unstable via its carbonyl oxygen (=O) sites and strongly unstable via its hydroxyl oxygen (-O) and amino nitrogen (-N) sites. Glycine preferred to interact with the $C_{12}B_8$ cage via its carbonyl oxygen (=O) active site, which was similar to the previous study of $C_{59}B$ -glycine [26] but disagrees with the C_{60} -glycine, [27] and C_{20} -glycine with the amino nitrogen (-N) active site [36]. Among two types of active atoms on the $C_{12}B_8$, it was easy to see that the B atoms were relatively favored in energy over the C atoms in the $C_{12}B_8$ -glycine while the stable ordering of three active sites on glycine molecule was =O site > -O site > -N site.

Table 1. Binding energies (E_b in eV) and ranges of carbon-carbon, C-B and C-X bond lengths (Å), for the $C_{12}X_8$ -glycine at the B3LYP/6-31+G* level

Species	E_b	C=C	C-B	C-X	B-X
C12B8-Gly(C=O)	4.28	1.395	1/654	1/572	-
C12B8-Gly(C-O)	22.00	1/398	1/582	1/439	-
C12B8-Gly(C-N)	21.36	1/397	1/559	1/396	-
C12B8-Gly(B=O)	4.06	1/394	1/640	-	1/546
C12B8-Gly(B-O)	20.28	1/388	1/567	-	1/373
C12B8-Gly(B-N)	19.99	1/390	1/598	-	1/389

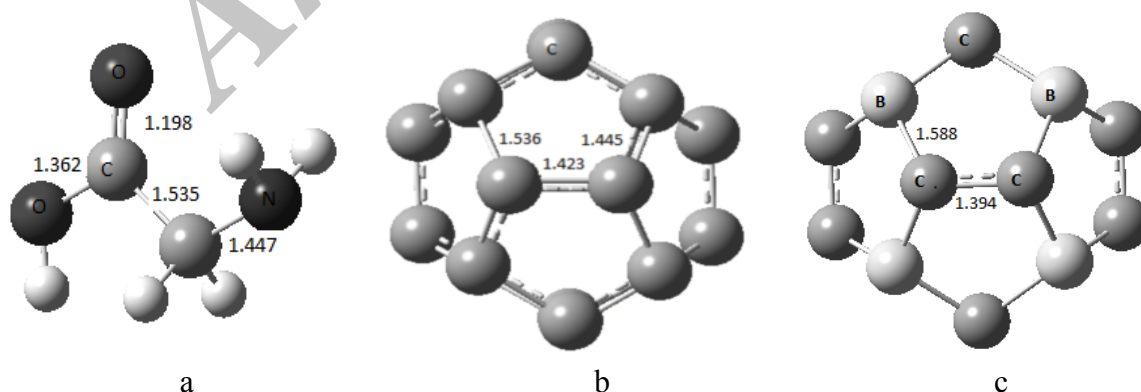
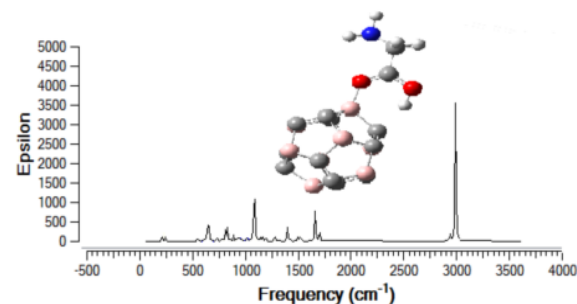


Fig. 1. Optimized structures of (a) glycine molecule (b) C20 fullerene and (c) $C_{12}B_8$ heterofullerene obtained at the B3LYP/6-31+G* level of theory.

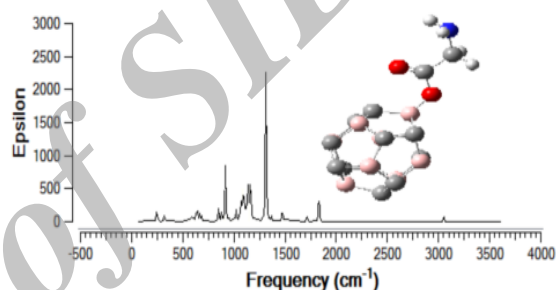
The lowest vibrational frequencies of C₁₂B₈ –glycine (B=O), being 9.32 cm⁻¹, involve mainly the wagging vibration of glycine backbone. The strongest band is found at 3299 cm⁻¹, which arise from the hydroxyl O–H stretching vibration. In order to further provide the distinctive spectroscopic fingerprints of our complexes, their calculated IR spectra were depicted in Fig. 2.

It worth to mentioned the HOMO – LUMO energy gap, (highest occupied molecular orbital (HOMO) - lowest

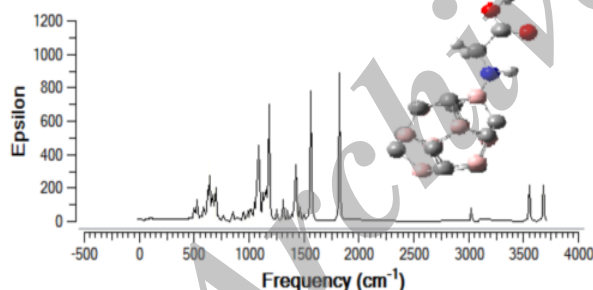
unoccupied molecular orbital (LUMO), because it can be associated with the optical and electrochemical properties of complexes. The electrons donated by a molecule in a reaction should be from its highest occupied molecular orbital (HOMO), while the electrons captured by the molecule should be located on its lowest unoccupied molecular orbital (LUMO). Furthermore, the atom, on which the HOMO mainly distributes, should have the ability for detaching electrons, whereas the atom with the occupation of the LUMO



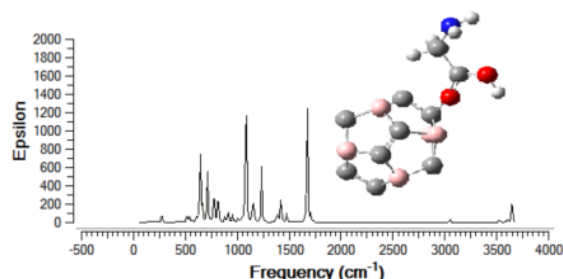
C12B8–B=O



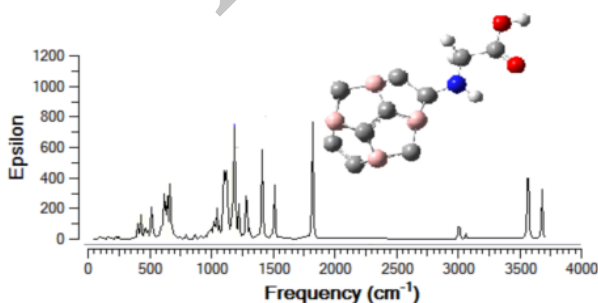
C12B8–B–O



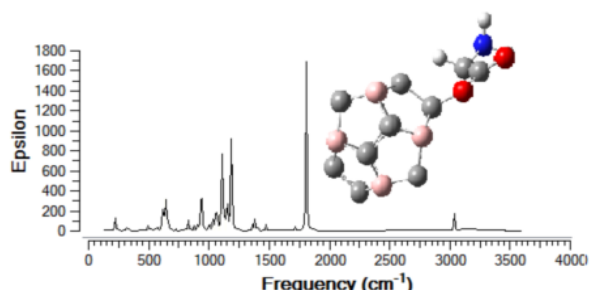
C12B8–C=O



C12B8–B–N



C12B8–C–O



C12B8–C–N

Fig. 2. The calculated IR spectra and optimized structures of C12X8–gly at B3LYP/6-31+G*.

Table 2. the smallest vibrational frequencies (ν_{\min}), the number of imaginary frequencies (NIMAG), $\Delta E_{\text{HOMO-LUMO}}$, (eV) and $\Delta E_{\text{reaction}}$ for $\text{C}_{12}\text{B}_8\text{-gly}$ at B3LYP/6-31+G*

System	ν_{\min}	NIMAG	$\Delta E_{\text{HOMO-LUMO}}$ (eV)	$\Delta E_{\text{reaction}}$ (kcal/mol)
$\text{C}_{12}\text{B}_8\text{-Gly-C=O}$	23.69	0	2.76	-64.48
$\text{C}_{12}\text{B}_8\text{-Gly-C-O}$	27.97	0	1.86	-62.60
$\text{C}_{12}\text{B}_8\text{-Gly-C-N}$	24.89	0	1.86	-59.62
$\text{C}_{12}\text{B}_8\text{-Gly-B=O}$	9.32	0	2.71	-65.35
$\text{C}_{12}\text{B}_8\text{-Gly-B-O}$	18.44	0	2.91	-63.09
$\text{C}_{12}\text{B}_8\text{-Gly-B-N}$	13.84	0	2.86	-63.94

should gain electrons. The HOMO–LUMO gap is traditionally associated with chemical stability against electronic excitation, with larger gap corresponding to greater stability. The gap of C_{12}B_8 is calculated for comparison with the results of gap of complexes.

The energy gap of hollow C_{12}B_8 cage was 2.86 eV. As a result, the C_{12}B_8 – glycine (C–O) and $\text{C}_{12}\text{B}_8\text{-glycine}$ (C–N) were 1.86eV, being smaller than that of C_{12}B_8 cage. The HOMO– LUMO gaps of C_{12}B_8 – glycine (C=O), C_{12}B_8 – glycine (B=O), C_{12}B_8 – glycine (B–O) and C_{12}B_8 – glycine (B–N) were 2.76, 2.71, 2.91 and 2.86 eV respectively. The results are shown in table 2. Molecular high-kinetic stability can be represented by a large HOMO – LUMO gap, because the molecule with large gap was unfavorable in energy to extract electrons from a low-lying HOMO orbital or to add electrons to a high-lying LUMO orbital; so it was relatively hard to form an activated compound.

CONCLUSION

We studied the interaction of the C_{12}B_8 fullerene with the smallest amino acid, glycine, by means of density-functional theory calculation. Six $\text{C}_{12}\text{B}_8\text{-glycine}$ conformations, generated from the C_{12}B_8 cage and glycine at different active sites, were considered to be further explored. Glycine preferred to interact with the C_{12}B_8

cage via its carbonyl oxygen (=O) active site. The $\text{C}_{12}\text{B}_8\text{-glycine}$ (C–O) and $\text{C}_{12}\text{B}_8\text{-glycine}$ (C–N) were 1.86 eV, being smaller than that of C_{12}B_8 cage. The HOMO – LUMO gaps of C_{12}B_8 – glycine (C=O), C_{12}B_8 – glycine (B=O), C_{12}B_8 – glycine (B–O) and C_{12}B_8 – glycine (B–N) are 2.76, 2.71, 2.91 and 2.86 eV respectively so C_{12}B_8 – glycine (B–O) had the more molecular kinetic stability.

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