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On molecular dynamics of the diamond D_5 seeds

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Abstract Diamond D_5 is a hyperdiamond with the rings being mostly pentagonal and built up on the frame of *mtn* structure, appearing in clathrate hydrates of type II. As the seed of D_5 , the centrohexaquinane C_{17} was proposed (Diudea, Stud Univ Babes-Bolyai Chem 55(4):11–17, [2010\)](#page-5-0). In this article, we study the molecular dynamics MD of four structures based on C_{17} skeleton, as all carbon or partly oxygenated derivatives. The results are discussed in terms of structural stability as given by DFT calculations as well as by the stable fluctuations of root mean square deviations and total, potential and kinetic energies provided by MD calculations.

Keywords Diamond $D_5 \cdot$ Centrohexaquinane \cdot HF \cdot DFT - Molecular dynamics

Introduction

In the Nano-era, a period starting with the discovery of C_{60} , in 1985, the carbon allotropes played a dominant role. Among the carbon structures, fullerenes (zero-dimensional), nanotubes (one dimensional), graphene (two dimensional), diamond, and spongy carbon (three dimensional) were the most studied $[1-3]$, both from theoretical reasons and applications perspective.

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Diamond D_6 , the beautiful classical diamond, with allhexagonal rings of sp^3 carbon atoms (Fig. [1](#page-1-0)), crystallized in a face-centered cubic fcc network (space group $Fd3m$), has kept its leading interest among the carbon allotropes, even as the ''nano'' varieties [\[4–9\]](#page-5-0). Its aesthetical appeal and mechanical characteristics are of great importance in jewelry and technique. Synthetic diamonds are currently produced by a variety of methods, including high pressure–high temperature, chemical vapor deposition, ultrasound cavitation [\[10\]](#page-5-0), etc.

However, the diamond D_6 is not unique: a hexagonal network called lonsdaleite (space group $P6_3/mmc$) [\[11](#page-5-0)], was discovered in a meteorite in the Canyon Diablo, Arizona, in 1967. Several diamond-like networks have also been proposed [[2,](#page-5-0) [12](#page-5-0), [13](#page-5-0)].

In a previous study, Diudea and Ilic^{[\[14](#page-5-0)]} described some multi-tori (i.e., structures showing multiple hollows—see Ref. [[15\]](#page-5-0)), one of them illustrated in Fig. [2,](#page-1-0) left.

The reduced graph of this multi-torus provided the structure for the seed of diamond D_5 : C_{17} , (Fig. [2,](#page-1-0) right) consisting of a tetravalent atom surrounded by six pentagons, the maximum possible number of pentagons around a $sp³$ carbon atom. According to the chemical nomenclature, C_{17} is a centrohexaquinane, a class of structures previously studied by Gund [\[16](#page-5-0)], Paquette [[17](#page-5-0)], and more recently by Kuck [\[18–20\]](#page-5-0).

Diamond D_5 is the name given by Diudea to diamondoids consisting mostly of pentagonal rings $[14, 21]$ $[14, 21]$ $[14, 21]$ $[14, 21]$ $[14, 21]$. D₅ is a hyperdiamond built up as a trinodal net in the frame of *mtn* structure; it belongs to the space group Fd-3m [\[22](#page-5-0)] while its seed is eventually the centrohexaquinane C_{17} . However, D_5 belongs to the family of Clathrates; it is precisely the type II clathrate C_{34} [[23\]](#page-5-0), of which Si_{34} -analogue was already synthesized.

 C_{17} can dimerize to $2 \times C_{17} = C_{34}$ $2 \times C_{17} = C_{34}$ $2 \times C_{17} = C_{34}$ (Fig. 3), the repeating unit, in crystallographic terms, of the diamond D_5 network. In crystallochemical terms, an adamantane-like

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Fig. 1 Diamond D_6 (left) and its repeating unit, adamantane (right)

Fig. 2 A multi-torus (left) and its reduced graph C_{17} (right), the seed of diamond D₅

Fig. 3 A joint of two C_{17} units (left) to give a dimer C_{34} (right), the repeating unit (in crystallographic terms) of the diamond D_5 network

Fig. 4 Adamantane-like structure (left) of the diamond D_5 network (right)

Table 1 The single point energies of the optimized structures at DFT (B3LYP/6-311+G**) level of theory

B3LYP	C_{17}	D,	\mathbf{D}_2	P_1 -766.479 6.274	
$B3LYP$ (a.u.)	-655.058	-766.491	-766.480		
B3LYP_Gap (eV)	5.868	6.461	6.264		

structure, as ada_20_158 (Fig. 4, left) is the monomer which will probably condense to form the diamond D_5 network (Fig. 4, right).

In the above symbols, "20" refers to C_{20} , which is the main unit of the hyper-diamond D_5 , while the last number counts the carbon atoms in structures.

The ratio C-sp³/C-total trends to 1 in a large enough network. As the content of pentagons $R \, [5]$ $R \, [5]$ per total rings trends to 90% in this yet hypothetical carbon allotrope, the net was called the diamond D_5 [\[21](#page-5-0)].

Method

The optimized structures, at Hartree–Fock HF (HF/6- $31G^{**}$) and DFT (B3LYP/6-311+ G^{**}) levels of theory, were submitted to molecular dynamics MD procedure. All calculations were performed in gas phase by Gaussian 09 [\[24](#page-5-0)] while MD calculations were done in vacuum, using Amber 10.0 software [[25\]](#page-5-0). The single point energy minima

Fig. 5 C_{17} hexaquinane trioxo-derivatives: Paquette P₁ (left) and Diudea, D_1 (*middle*) and D_2 (*right*)

obtained for the investigated structures are shown in Table 1. Before the MD was run, the atomic charges were calculated according to Merz-Kollmann scheme via the RESP [[26\]](#page-5-0) procedure at HF/6-31G** level. The AMBER force field [\[27](#page-5-0)] was used for dynamic trajectory generation. There were several steps of molecular dynamics. After stabilization of energies and root mean square deviation (RMSD) values during run, the actual molecular dynamics were performed, in a cascade way. Each tested system was heated by 20 ps while MD simulations were 100 ns long. The visualizations were prepared in the GaussView program. After MD run, the values of RMSD and energies of analyzed structures were

Table 2 The averages total energy (E_{tot}) values estimated, by MD, on geometries in the gas phase

300	600	900	1200	1400	1600	1800
127.405	180.379	233.358	286.460	322.227	357.665	393.123
5.789	11.585	17.373	23.121	27.070	30.932	34.938
300	600	900		1200	1400	1600
144.996	197.907	250.689		303.114	337.947	373.136
5.782	11.593	17.273		23.106	26.739	30.745
	300			600		900
	116.45			169.266		222.049
	5.783			11.542		17.363
300	600	900		1200	1400	1600
142.574	195.448			300.329	335.524	370.602
5.798	11.535			23.076	26.931	30.678
				247.889 17.300		

The averaged values were calculated on all the generated points of energies in every 1 ps of molecular dynamics. Symbol δ means the standard deviation

Fig. 6 The plot of total energy (E_{tot}) versus temperature (Temp)

recorded: total energy (E_{tot}) , kinetic energy (E_{kin}) , and potential energy (E_{pot}) . In the analysis, averaged values of all generated points of energies and values of RMSD in every 1 ps of MD were used.

Results and discussion

Stability evaluation was performed on four hypothetical seeds of D_5 , the all-carbon structure C_{17} (Fig. [2](#page-1-0), right) and

Fig. 7 The plot of RMSD versus temperature (Temp)

three trioxa-derivatives of C_{17} . The isomer in Fig. [5](#page-1-0), left was synthesized by Paquette and Vazeux [[17\]](#page-5-0), and is hereafter denoted P_1 . The other two structures, denoted D_1 and D_2 (Fig. [5,](#page-1-0) middle and right) were proposed, by Diudea, as possibly appearing in rearrangements of the Paquette's P_1 structure. The last two structures would be the appropriate ones in the next step of dimerization to C_{34} , in fact the repeating unit $[23]$ $[23]$ of the structure we call here D_5 .

The stability of molecules was evaluated both in static and dynamic temperature conditions. The isomer D_1 seems the most stable among all studied structures, as given by

Table 3 The values of standard deviations of E_{tot} at a given temperature (see the center of each slide), for the four investigated structures

Table 4 The averaged RMSD values estimated by molecular dynamics MD on the geometries in the gas phase

The averaged values were calculated on all the generated points of RMSD in every 1 ps of MD. Symbol δ represents the standard deviation

Fig. 8 The structure of the tested hypothetical seeds of the diamond $D₅$ during molecular dynamics

optimization in gas phase at DFT level (Table [1\)](#page-1-0). In decreasing the stability, it follows P_1 and D_2 . However, the all-carbon C_{17} appears the most stable at MD treatment, even at DFT level is the last one. This is probably because the C–C bond is more stable at temperature variations (see Fig. 8).

In MD, C_{17} keeps its structure up to about 1800 K, while its destruction starts at 2,000 K (Tables [2](#page-2-0), [4;](#page-3-0) Figs. [6,](#page-2-0) [7](#page-2-0)). Kuck has reported a centrohexaindane, as the most symmetric structure in this series but also a benzo-centro-hexaquinane [[19,](#page-5-0) [20](#page-5-0)] as the last step structure in the synthesis of a non-planar 3D structure, designed according to mathematical rules. However, in the synthesis of centrohexaquinane-derivatives, C_{17} remained yet elusive.

Very close to C_{17} behaves the oxygen-containing isomer D_1 , as expected from its highest stability at DFT level (Table [1](#page-1-0)-the values in bold).

Despite a very long time of molecular dynamics (100 ns) was leaded, it is believed that prolonged annealing at 1,800 K for both P_1 and D_1 isomers finally resulted in the destruction of these molecules. Thus, P_1 and D_1 isomers

behave similarly in MD conditions. The isomer D_2 was the least stable one, as the largest RMSD values were recorded for this isomer.

According to molecular dynamics, it is clear that by increasing temperature resulted in higher values of energy and RMSD (and their standard deviations) of all the analyzed structures, with high values of correlation. The plots of E_{tot} versus temperature for all tested systems are given in (Fig. [6\)](#page-2-0), while for RMSD the plots are given in Fig. [7.](#page-2-0) As expected, the correlations in the RMSD plot are a little lower than those for E_{tot} . The MD calculations, listed in Tables [2](#page-2-0), [3,](#page-3-0) and [4](#page-3-0) show the followings.

As can be seen from Tables [2](#page-2-0) and [3,](#page-3-0) the values of standard deviations of the averaged values of E_{tot} are closely correlated with the values of temperature, in the range the molecular dynamics simulations were done. The values of these standard deviations at a given temperature are similar for all four studied structures (Tables [2,](#page-2-0) [3](#page-3-0)), due to their structural relatedness.

The smallest values of the RMS deviation are observed for C₁₇, with the lowest values of standard deviation (δ) at all the studied values of temperature (Table [4\)](#page-3-0).

In the case of P_1 , one can see a similar behavior but somewhat with larger values of RMSD (as well as values of the standard deviation), compared to the all-carbon structure C_{17} (Table [4\)](#page-3-0). It confirms the structural stability of the above structures. The largest values of the RMS deviation were recorded for D_2 isomer (Table [4](#page-3-0); Fig. [7](#page-2-0)).

Visualization of the structural changes (first step destruction, the right column) is presented in Fig. 8.

Conclusions

Structural stability of several hypothetical seeds of the diamond D_5 was investigated. It was evaluated both in static and dynamic temperature conditions by molecular dynamics MD. During MD, the all-carbon C_{17} appeared the most resistant to changes of temperature. Structural and energetic stability of the other three seeds of D_5 vary both with the values of temperature and evolution time in molecular dynamics and the arrangement of oxygen atoms in the molecules. Among all the studied structures, the D_2 isomer is the most sensitive to changes in temperature. After optimization by B3LYP, D_1 isomer seemed to be the most stable one. The structure stability of D_1 and P_1 isomers in MD are similar. These two isomers are only slightly more sensitive to temperature as compared with the all-carbon C_{17} . These results could be useful in guiding further reactions, e.g., the dimerization to C_{34} and condensation to adamantane-like structures, finally leading to the diamond D_5 .

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