Data Article

Dimethylsulfoxide (DMSO) clusters dataset: DFT relative energies, non-covalent interactions, and cartesian coordinates

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A R T I C L E  I N F O

Article history:
Received 19 February 2022
Revised 1 March 2022
Accepted 2 March 2022
Available online 7 March 2022

Keywords:
Dimethylsulfoxide clusters
Non-covalent interactions
QTAIM analysis
Relative energies

A B S T R A C T

Theoretical understanding of dimethylsulfoxide (DMSO) liquid depends on the understanding of the DMSO clusters. In this work, we provide the structures and the energetics of the DMSO clusters. The structures have been generated using ABCluster and further optimized at the MP2/aug-cc-pVDZ level of theory. The final structures have been optimized at two different levels of theory: PW6B95D3/aug-cc-pVDZ and \(\omega\)B97XD/aug-cc-pVDZ. The Cartesian coordinates of the structures optimized at the MP2/aug-cc-pVDZ level of theory are also reported. The relative energies of the structures can be used to locate the most favorable structures of the DMSO clusters. The Cartesian coordinates of the structures can be used for further investigations on DMSO clusters. In addition, we report the data related to the quantum theory of atoms in molecule (QTAIM) analysis of the investigated clusters. The QTAIM data reported in this work can be used to understand and determine the nature of non-covalent interactions in DMSO clusters. For further reading and discussion on the data reported here, please report to the original manuscript Malloum and Conradie (2022) [1].

DOI of original article: 10.1016/j.molliq.2022.118522
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https://doi.org/10.1016/j.dib.2022.108024
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Specifications Table

| Subject               | Chemistry |
|-----------------------|-----------|
| Specific subject area | Physical and Theoretical Chemistry |
| Type of data          | Figure Table |
| How data were acquired| Electronic energies and the Cartesian coordinates of the structures are generated using Gaussian 16, at the three different levels of theory. The data related to the quantum theory of atoms in molecule (QTAIM) analysis are generated using the AIMAll program. |
| Data format           | Analyzed Raw |
| Description of data collection | Raw data are retrieve from the output of the Gaussian calculations. QTAIM data (analyzed data) are from the AIMAll program. Relative energies (analyzed data) are calculated using the electronic energies from Gaussian output files. Geometries are optimized using the resources of the South African Center of High Performance Computing (CHPC). We used our Laboratory clusters to perform QTAIM analysis. |
| Data source location  | Institution: Physical Chemistry Laboratory of the Department of Chemistry, University of the Free State |
|                       | City/Town/Region: Bloemfontein |
|                       | Country: South Africa |
| Data accessibility     | Repository name: Mendeley Data |
|                       | Data identification number (DOI number): 10.17632/bwfjhwvhcz.1 |
|                       | Direct link to the dataset: https://doi.org/10.17632/bwfjhwvhcz.1 |
| Related research article| A. Malloum, J. Conradie, Non-Covalent Interactions in Dimethylsulfoxide (DMSO) Clusters and DFT Benchmarking, J. Mol. Liq. 350 (2022) 118522. |
|                       | https://doi.org/10.1016/j.molliq.2022.118522. |

Value of the Data

- The data reported in this work are important to understand the hydrogen bond network of the dimethylsulfoxide (DMSO) clusters. This understanding is important for proper theoretical description of the liquid DMSO.
- The relative energies at different levels of theory provide the quantitative data necessary to select relevant structures of the DMSO clusters.
- The QTAIM data can be used to determine covalent and non-covalent bonds of the DMSO clusters, as well as their strength.
- The Cartesian coordinates of the located structures will be of extreme help for further investigations involving DMSO clusters.

1. Data Description

The data reported in this work are constituted of analyzed data and raw data. The analyzed data are constituted of the structures of the DMSO clusters from $n = 2$ to $n = 4$ and their relative energies as calculated at two different levels of theory: PW6B95D3/aug-cc-pVDZ and $\omega$B97XD/aug-cc-pVDZ. The structures and their relative energies are reported in Figs. 1 and 2. Each sub-caption of the structures in Figs. 1 and 2 reports the name of the structure (in accordance with the naming in the main paper [1]); the relative energies at the PW6B95D3/aug-cc-pVDZ level of theory (and at the $\omega$B97XD/aug-cc-pVDZ level of the theory in brackets); and the symmetry point group of the structure. In addition to the structures and their relative energies, analyzed data are also constituted of data from quantum theory of atoms in molecule...
Fig. 1. Structures of the DMSO dimer and trimer as optimized at the PW6B95D3/aug-cc-pVDZ level of theory. Numbers represent the zero point corrected relative energies at the PW6B95D3/aug-cc-pVDZ level of theory (and ωB97XD/aug-cc-pVDZ level of theory in brackets).

(QTAIM) analysis. Data from QTAIM analysis of the DMSO dimers are provided in the supplementary material. These data are constituted of the bond critical points information of the investigated structures of the DMSO clusters. To avoid cumbersome results, the QTAIM analysis data of the DMSO dimers, trimers, and tetramers are reported in the supplementary material. The reader is referred to the supplementary file for clear assessment of the QTAIM data. Each of the tables in the supplementary material has seven columns which describes respectively, the bond critical point’s name (name), the bond critical point’s atoms (Atoms), the bond critical point’s electron density (ρ), the bond critical point’s Laplacian of the electron density (∇²ρ), the bond’s ellipticity (Ellipticity), the bond critical point’s kinetic energy density (K) and the difference between the geometric bond length and the bond path length (BPL-GBL_I). As far as raw data of this work are concerned, they are constituted of the structures’ Cartesian coordinates (as
Fig. 2. Structures of the DMSO tetramer as optimized at the PW6B95D3/aug-cc-pVDZ level of theory. Numbers represent the zero point corrected relative energies at the PW6B95D3/aug-cc-pVDZ level of theory (and ωB97XD/aug-cc-pVDZ level of theory in brackets).

provided in Figs. 1 and 2). The MP2/aug-cc-pVDZ level of theory has been used for the optimization of the Cartesian coordinates. These Cartesian coordinates are reported in the supplementary material.

2. Experimental Design, Materials and Methods

The methodology (or methods) used to generate the data presented in this paper has been extensively presented in the main article (related research article) [1]. Interested reader are invited to read the related research article for details on the methodology [1]. Nevertheless, we will provide in the next few sentences the methodological procedure used to acquire the data.

Initially, we use ABCluster code [2,3] to generate structures. We have already used ABCluster to generate the structures of clusters in our previous works [4–7]. In addition, we have pointed out clearly the efficiency of the ABCluster as compared to density functional theory (DFT) and
MP2 [4]. These works have shown that ABCluster is efficient, easy to use and reliable for the generation of initial structures of molecular clusters. The structures of the DMSO clusters generated by ABCluster are optimized at the MP2/aug-cc-pVDZ level of theory. The output files of the optimization, using MP2/aug-cc-pVDZ, have been used to retrieve the optimized Cartesian coordinates of the DMSO clusters which are reported in this work as supplementary material. The structures obtained at the MP2 are re-optimized at two more levels of theory: PW6B95D3/aug-cc-pVDZ and ωB97XD/aug-cc-pVDZ. The output files of these optimizations have been used to retrieve the relative energies of the DMSO clusters reported in Figs. 1 and 2. Furthermore, the images of the structures provided in Figs. 1 and 2 are also retrieved from the output files of the calculations. It is worth noting that all optimization have been performed using the Gaussian 16 suite of programs.

Regarding the data related to the QTAIM analysis, we used the AIMAll program [8] to generate the data. To generate the data with QTAIM through AIMAll, we use the Gaussian checkpoint files containing the orbitals and the electron density information of the structures. AIMAll generates the data reported in the supplementary file as an excel spreadsheet file. The data contains mainly the critical points of the electron density topology and their related information (properties). These properties can be used to understand and quantify the strength of the bondings of the structures [9,10]. The properties can also be used to differentiate and classify the bondings (covalent and non-covalent) based on the electron density and the Laplacian of the electron density [11,12].

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships which have, or could be perceived to have, influenced the work reported in this article.

CRediT Author Statement

Alhadji Malloum: Conceptualization, Methodology, Validation, Formal analysis, Investigation, Data curation, Writing – original draft, Visualization; Jeanet Conradie: Resources, Visualization, Writing – review & editing, Supervision, Funding acquisition, Project administration.

Acknowledgments

The South African Center for High Performance Computing (CHPC) (Grant no. CHEM0947) is acknowledged for computational resources and clusters access. The Central Research Fund of the University of the Free State is also acknowledged for funding. The Norwegian Supercomputing Program (UNINETT Sigma2, Grant no. NN9684K) is acknowledged for computer time.

Supplementary Material

Cartesian coordinates of the structures, that have been fully optimized using the MP2 method associated with aug-cc-pVDZ basis set, are given. In addition, bond critical points QTAIM analysis properties of the isomers of the DMSO clusters are provided herein.

Supplementary material associated with this article can be found in the online version at doi:10.1016/j.dib.2022.108024

References

[1] A. Malloum, J. Conradie, Non-covalent interactions in dimethylsulfoxide (DMSO) clusters and DFT benchmarking, J. Mol. Liq. 350 (2022) 118522.
[2] J. Zhang, M. Dolg, Abcluster: the artificial bee colony algorithm for cluster global optimization, Phys. Chem. Chem. Phys. 17 (2015) 24173–24181.

[3] J. Zhang, M. Dolg, Global optimization of clusters of rigid molecules using the artificial bee colony algorithm, Phys. Chem. Chem. Phys. 18 (2016) 3003–3010.

[4] A. Malloum, J.J. Fifen, Z. Dhaouadi, S.G.N. Engo, J. Conradie, Structures, relative stabilities and binding energies of neutral water clusters, (H$_2$O)$_2$.30, New J. Chem. 43 (2019) 13020–13037.

[5] A. Malloum, J. Conradie, Solvent effects on the structures of the neutral ammonia clusters, Comput. Theor. Chem. 1191 (2020) 113042.

[6] A. Malloum, J. Conradie, Global and local minima of protonated acetonitrile clusters, New J. Chem. 44 (2020) 17558–17569.

[7] A. Malloum, J. Conradie, Structures of water clusters in the solvent phase and relative stability compared to gas phase, Polyhedron 193 (2021) 114856.

[8] Keith, T. A. (2019). TK gristmill software. Overland Park KS, USA, 11, 16. (aim.tkgristmill.com).

[9] S.J. Grabowski, What is the covalency of hydrogen bonding? Chem. Rev. 111 (2011) 2597–2625.

[10] R. Parthasarathi, V. Subramanian, N. Sathyamurthy, Hydrogen bonding without borders: an atoms-in-molecules perspective, J. Phys. Chem. A 110 (2006) 3349–3351.

[11] R.G. Bone, R.F. Bader, Identifying and analyzing intermolecular bonding interactions in van der Waals molecules, J. Phys. Chem. 100 (1996) 10892–10911.

[12] I. Alkorta, F. Blanco, J. Elguero, J.A. Dobado, S.M. Ferrer, I. Vidal, Carbon···carbon weak interactions, J. Phys. Chem. A 113 (2009) 8387–8393.