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Propeller-like Conformation of Diphenylacetic Acid

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Abstract Crystal structure of diphenylacetic acid has been solved by X-ray diffraction. The crystals are monoclinic, space group P2₁/c, with a = 12.254(4) Å, b = 7.2260(8) Å, c = 17.521(4) Å, $\beta = 133.38(1)^{\circ}$, Mr = 212.24, V = 1127.6(5) Å³, Z = 4 and R = 0.045. A strong hydrogen bond links the molecules in dimers. The dimers are connected by weaker C-H··· π and π ··· π interactions. A calculation was performed for the isolated molecule and for the dimer within the Hartree-Fock (HF) level with a 6-311G(d) basis set. In both calculations, the minimum of the energy is achieved with the phenyl rings assuming a more symmetric arrangement around the central carboxylic plane than is experimentally observed.

Keywords Diphenylacetic acid · Liquid crystal · Crystal structure · Hydrogen bonds · Hartree-Fock calculation

Introduction

Liquid crystals are substances that exhibit a phase of matter with properties between those of a conventional liquid, and those of a solid crystal. Many substances are liquid

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crystals, some of them as common as soap, and in most of them the liquid crystalline structure consists of a strip-like or disc-like core in the centre and long alkyl chains in the outer region. A few are of the flying-seed-like type. Ohta et al. have shown that sodium diphenylacetate is one of this rare liquid crystals showing a hexagonal columnar mesophase [1]. The arrangement of the molecules in the liquid crystal phase correlates with and resembles that observed in the solid state, according to our study [2]. In the crystal structure of sodium diphenylacetate, each Na⁺ ion is coordinated by four carboxylate O atoms at distances in the range 2.207 (2)-2.467 (3) Å to form cubes of Na and O atoms which are linked via the carboxylate C atoms into a columnar structure. The two phenyl groups are in the outside of the columns and are oriented like the blades of a propeller. Preliminary results on potassium diphenylacetate show the same formation of columns while the rubidium diphenylacetate crystal structure has less symmetry giving rise to the formation of ladders instead of columns [3].

Within our project of studying diphenylacetate complexes, we have undertaken the study of pure crystalline diphenylacetic acid. A search on the CSD database showed that the diphenylacetic acid atomic coordinates were unavailable. Minimum crystal data have been published by Koshima et al. [4].

The molecule of diphenylacetic acid has the ability of promoting absolute asymmetric synthesis, that is to assist the formation of enantiomerically enriched products from achiral precursors without the intervention of chiral chemical reagents or catalysts. Diphenylacetic acid has been used to cocrystallize with acridines originating chiral crystals [4, 5]. It is the propeller like arrangement of the phenyl rings in the diphenylacetic acid molecule that causes a chiral conformation in an otherwise achiral molecule.

To investigate the effect of the intermolecular interactions in the conformation of the molecule we have performed an optimization of the geometry of the isolated molecules and of the strongly H-bonded dimer by ab-initio Hartree-Fock calculations.

Experimental and Computational Methods

Preparation of Diphenylacetic Acid

Single crystals of the title compound were collected directly from the flask purchased from Sigma-Aldrich (99%).

Crystal Structure Determination

A crystal of the title compound having approximate dimensions of 0.49 mm \times 0.24 mm \times 0.22 mm was glued on a glass fiber and mounted on a Bruker Apex II difractometer. Diffraction data were collected at room temperature 293(2) K using graphite monochromated Mo K α ($\lambda = 0.71073$ Å). The crystallographic structure was solved by direct methods (SHELXS-97) [6]. Refinements were carried out with SHELXL-97 [6] package. All refinements were made by full-matrix least-squares on F [2], with anisotropic displacement parameters for all nonhydrogen atoms. Hydrogen atoms were included in the refinement in calculated positions. Exception made to the carboxylic H atom whose coordinates were allowed to refine and $U_{iso}(H) = 1.2U_{eq}(O)$. The final least-squares cycle was based on 2,296 observed reflections $[I > 2\sigma(I)]$ and 145 variable parameters, converged with R = 0.0451and wR = 0.1271. Additional information to the structure determinations is given in Table 1. Hydrogen bond geometric data is given on Table 2. Selected bond distances and angles can be seen in Table 3. Supplementary data have been deposited at the Cambridge Crystallographic Data Centre (CCDC No. 652158).

Calculations

The structure optimization calculations of the isolated molecule and of the dimer were performed using the code GAMESS [7] within the Hartree-Fock approximation with a 6-311G(d) basis sets [8–11]. The calculation was relaxed from the X-ray geometry. At the final equilibrium geometry the maximum gradient was 10^{-5} Hartree Bohr⁻¹. The molecules of the dimer were related by an imposed inversion centre.

Table 1 Crystal data and structure refinement the title compound
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Table I Crystal data and structure	rennement the title compound
Empirical formula	$C_{14}H_{12}O_2$
Formula weight	212.24
Temperature (K)	293(2)
Wavelength (Å)	0.71073
Crystal system	Monoclinic
Space group	P2 ₁ /c
<i>a</i> (Å)	12.254(4)
b (Å)	7.2260(8)
<i>c</i> (Å)	17.521(4)
α (°)	90
β (°)	133.38(1)
γ (°)	90
Volume (Å ³)	1127.6(5)
Ζ	4
Calculated density (g/cm ³)	1.250
Absorption coefficient (cm ⁻¹)	0.083
<i>F</i> (000)	448
Crystal size (mm)	$0.49\times0.24\times0.22$
θ range for data collection (°)	2.29-30.89
Index ranges	$\begin{array}{l} -16 < h < 17, \ -10 < k < 10, \\ -25 < l < 25 \end{array}$
Reflections collected/unique	$32,797/3,483 \ [R(int) = 0.0282]$
Completeness to $\theta = 30.89$	97.6%
Refinement method	Full-matrix least-squares on F^2
Data/restraints/parameters	2296/0/145
Goodness–of–fit on F^2	1.071
Final <i>R</i> indices $[I > 2\sigma(I)]$	$R_1 = 0.0451 \ wR_2 = 0.1102$
R indices (all data)	$R_1 = 0.0772 \ wR_2 = 0.1271$
Largest diff. peak and hole (e \AA^{-3})	0.250 and -0.176

Results and Discussion

Crystal Structure of the Title Compound

The structure of the title compound together with the atomnumbering scheme, is illustrated in Fig. 1. The packing diagram is shown in Fig. 2. Geometric details of the intermolecular interactions are listed on Tables 2–3. Selected bond lengths and angles are listed in Table 4.

The title compound, diphenylacetic acid (Fig. 1), crystallizes in a centrosymmetric space group P2₁/c with the acid molecules assembled in dimers via strong hydrogen bonds between the carboxylic groups. The donor-acceptor distance is 2.660(2) Å with the bond angle $175(3)^{\circ}$. The conformation of the phenyl rings is like the blades of a propeller making C2 a chiral centre. An inversion centre relates the two dimers thus making the whole structure achiral. Besides the strong hydrogen bond, other intermolecular interactions play a role in governing the crystal packing: C–H… π and π … π interactions.

Table 2 Intermolecular H-bond geometry (Å, °)

	D–H	Н…А	D····A	D–H…A
O2-H21O1 ⁱ	0.94(2)	1.73(2)	2.660(2)	175(3)
(Symmetry code	i: -1 - x, -	-y, -1 - z		

Table 3 C–H··· π interactions geometry (Å, °)

	$d_{H\cdots\pi}$	θ	α	$d=d_{H\cdots\pi}cos\theta$	
C2–H2···Cg1 ⁱⁱ	3.053	65.3	171.7	1.278	Type II
C7–H7…Cg2 ⁱⁱⁱ	3.325	74.6	126.5	0.881	Type III
C8–H8····Cg2 ⁱⁱⁱ	3.265	60.8	129.4	1.592	Type V
C12–H12···Cg2 ^{iv}	3.099	85.7	146.5	0.233	Type III

(Symmetry codes ii: -x, -1/2 + y, -1/2 - z, iii: x, 1 + y, z iv: -1 - x, -1/2 + y, -1/2 - z; Cg1-centroid of ring C3–C8, Cg2-centroid of ring C9–C14, definition of geometric parameters according to [12])

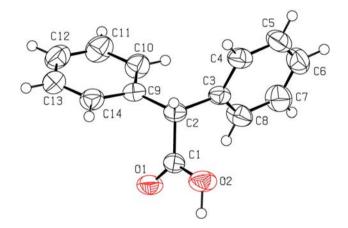


Fig. 1 ORTEP diagram of the title compound with the ellipsoids drawn at the 50% probability level, with the atomic labelling scheme

There are four C–H··· π intermolecular interactions, involving four hydrogen atoms and the π electron system of the two rings (Table 3). In two of the bonds, the hydrogen atom is directly above the centre of the ring but the C–H bond points towards a ring carbon. This geometry corresponds to a type III interaction as classified by Malone et al. [12]. In another bond (C2–H2··· π), the classical T-shape geometry is almost accomplished, but a slight deviation of the H atom from the centre of the ring makes it more of the type II. In the remaining bond (C8–H8··· π) the hydrogen atom interacts with a carbon at the edge of the acceptor ring (type V). Two of the above bonds (C7–H7··· π and C8–H8··· π) share the H atoms with the same π system. The C–H··· π interactions join the molecules in layers parallel to the *ab* plane (Fig. 2).

The $\pi \cdots \pi$ interaction displays the usual slipped stacking geometry, with the interacting π systems parallel displaced.

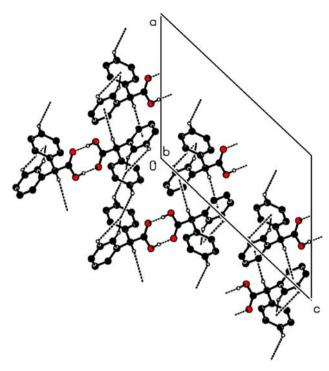


Fig. 2 Partial packing diagram of the title compound with the Hbonds and C-H $\cdots \pi$ interactions drawn as dashed lines

 Table 4
 Selected structural parameters by X-ray and theoretical calculations

Bond distances (Å)	Experimental	Calculated		
and angles (°)		Monomer	Dimmer	
C101	1.215(1)	1.182	1.196	
C1O2	1.309(1)	1.327	1.304	
O1C1O2	122.7(1)	122.19	123.27	
O2-C1-C2-C3	67.1(1)	105.27	102.90	
O2-C1-C2-C9	-165.3(1)	-126.85	-128.85	
C1C2C3C8	40.5(2)	33.74	35.20	
C1C2C3C14	63.9(1)	105.65	106.38	

The centroid–centroid distance is 4.867(2) Å and the angle between the ring normal and the vectors between the ring centroids is 39.9(1)°. The shorter atom-atom contact between the two parallel planes is 3.771(2) Å (C7 to C7^{*i*}. *i*: -x, -y, -z), characteristic of a weak interaction [13]. The $\pi \cdots \pi$ interactions join the dimers in chains that run along the *c* axis (Fig. 2).

The geometry optimization for the isolated molecule converges to a molecular conformation more symmetric than the one observed in the solid state (Fig. 3). The calculated parameters show an angle between the least-squares planes of both phenyl rings of 74.1°. The angle between the mean plane of phenyl ring A (C3–C8) and the carboxylic mean plane is 64.4°. The angle between the

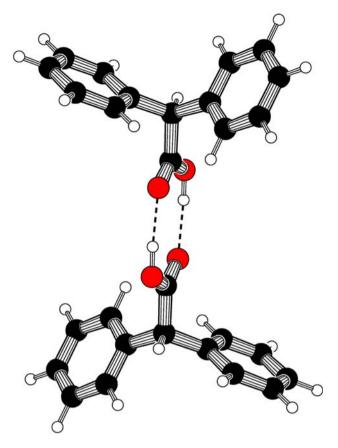


Fig. 3 The molecular conformation of the dimer as resulted from the Hartree-Fock calculations

mean plane of phenyl ring B (C9–C14) and the carboxylic mean plane is 64.2° . There is a strong asymmetry between the C–O bond lengths (see Table 3).

The geometry optimization for the dimer converges to a less symmetric molecular conformation. The strong hydrogen bond observed in the solid state is maintained. The asymmetry between the C–O bond lengths diminishes since the H atom is shared between O2 and O1 of inversion related molecules. The calculated parameters show an angle between the least-squares plane of both phenyl rings of 72.8°. The angle between the mean plane of phenyl ring A and the carboxylic mean plane is 66.3° . The angle between the mean plane of phenyl ring B and the carboxylic mean plane is 63.5° .

In the solid state the intermolecular interactions distort even more the symmetry of the isolated monomer. There is a great involvement of the rings in intermolecular interactions both as donors and acceptors and different conformations may be favoured energetically. The energy of a mole of dimers is -861984 kcal/mol for the observed conformation and -862255 kcal/mol for the optimized geometry, which indicates the role of the packing forces in the cohesion of the crystal. The angle between the least-squares planes of the phenyl rings is $83.99(9)^\circ$. The angle between the mean plane of phenyl ring A and the carboxylic mean plane is $86.81(8)^\circ$. The angle between the mean plane of phenyl ring B and the carboxylic mean plane is $70.03(8)^\circ$.

The HOMA parameter [14] describing aromaticity, which has a value of 1.000 for benzene, is equal to 0.961 (for ring A) and 0.982 (for ring B). In both calculations the converged geometry results in equal HOMA parameters for the phenyl rings [0.996 for the dimer and 0.997 for the monomer].

Supplementary Material

Crystallographic data for structural analysis have been deposited with the Cambridge Crystallographic Data Center, CCDC 652158. Copies of this information may be obtained free of charge on application to CCDC, 12 Union Road, Cambridge CB2 1EZ, UK (fax: +44 1223 336 033; e-mail: deposit@ccdc.cam.ac.uk or http://www.ccdc.cam.ac.uk).

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