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Crystal structure and DFT study of 8-hydroxy-1,2,3,5,6,7-hexahydropyrido[3,2,1-*ij*]quinoline-9carbaldehyde

Md. Serajul Haque Faizi,^a Necmi Dege^b and Maria L. Malysheva^{c*}

^aDepartment of Chemistry, College of Science, Sultan Qaboos University, PO Box 36 Al-Khod 123, Muscat, Oman, ^bOndokuz Mayıs University, Arts and Sciences Faculty, Department of Physics, 55139 Samsun, Turkey, and ^cDepartment of Chemistry, Taras Shevchenko National University of Kyiv, 64, Vladimirska Str., Kiev 01601, Ukraine. *Correspondence e-mail: maria_malysheva@mail.univ.kiev.ua

In the title compound, $C_{13}H_{15}NO_2$, the fused non-aromatic rings of the juloidine moiety adopt envelope conformations. The hydroxy group forms an intramolecular hydrogen bond to the aldehyde O atom, generating an S(6) ring motif. Weak intermolecular $C-H\cdots O$ hydrogen bonds help to stabilize the crystal structure. Density functional theory (DFT) optimized structures at the B3LYP/ 6–311 G(d,p) level are compared with the experimentally determined molecular structure in the solid state.

1. Chemical context

Julolidine is chemically an aniline derivative with two N-alkyl substituents forming rings back to the aromatic ring; the fused rings lock the nitrogen lone-pair of electrons into conjugation with the aromatic ring leading to unusual reactivity. The presence of the julolidine ring system in some molecules makes them useful for chromogenic naked-eye detection of copper, zinc, iron and aluminium ions as well as fluoride ions (Wang et al., 2013; Choi et al., 2015; Kim et al., 2015; Jo et al., 2015). Julolidine dyes exhibit excited-state intramolecular proton transfer (Nano et al., 2015). Compounds containing lulolidine rings are also used as fluorescent probes for the measurement of cell-membrane viscosity. Julolidine-based materials are also used as red emitters in OLEDs when linked to dicyanomethylpyran modules (Lee, et al., 2012). The julolidine unit plays an important role as it has strong electronicdonating properties for chelating (Nano, et al., 2013). Julolidine malononitrile acts as a 'push-pull' molecule with large hyperpolarizability and is used as a model system for understanding the non-linear optical properties of molecules (Mennucci et al., 2009).

There are many reports in the literature on julolidine-based Schiff bases and their applications as sensors for metal ions (Park *et al.*, 2014; Lee *et al.*, 2014; Kim *et al.*, 2016). The present work is a part of an ongoing structural study of Schiff bases based on the julolidine ring system (Faizi *et al.*, 2016, 2017). We report here the crystal structure and DFT computational calculation of the title julolidine compound (I). The results of calculations by density functional theory (DFT) on (I) carried out at the B3LYP/6–311 G(d,p) level are compared with the experimentally determined molecular structure in the solid state.

research communications



Table 1	
Hydrogen-bond geome	try (Å, °).

, , ,	2	,		
$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdot \cdot \cdot A$
O2−H2···O1	0.82	1.89	2.621 (2)	148
$C9-H9A\cdots O2^{i}$	0.97	2.50	3.324 (3)	143
$C9-H9B\cdots O1^{ii}$	0.97	2.55	3.438 (3)	152
$O2-H2 \cdot \cdot \cdot O1$	0.82	1.89	2.621 (2)	148
$C9-H9A\cdots O2^{i}$	0.97	2.50	3.324 (3)	143
$C9-H9B\cdotsO1^{ii}$	0.97	2.55	3.438 (3)	152

Symmetry codes: (i) x, y + 1, z; (ii) -x + 1, -y + 1, -z + 1.

3. Supramolecular features

In the crystal, molecules are linked by $C-H\cdots O$ hydrogen bonds, forming an A-B-A-B-A-B arrangement through the inversion centre and propagating along the *c*-axis direction (see Fig. 2 and Table 1). There are no other significant intermolecular contacts present in the molecule.

4. DFT study

The DFT quantum-chemical calculations were performed at the B3LYP/6–311 G(d,p) level (Becke, 1993; Lee *et al.*, 1988) as implemented in *GAUSSIAN09* (Frisch *et al.*, 2009). DFT structure optimization of (I) was performed starting from the X-ray geometry and the values compared with experimental values (see Table 2). From these results we can conclude that basis set 6–311 G(d,p) is well suited in its approach to the experimental data.

The DFT study of (I) shows that the HOMO and LUMO are localized in the plane extending from the whole julolidine ring to the salicylaldehyde ring. The electron distribution of the HOMO-1, HOMO, LUMO and the LUMO+1 energy levels are shown in Fig. 3. The molecular orbital of HOMO





A view of the A-B-A-B-A-B arrangement in the crystal structure of the title compound. The hydrogen bonds are shown as dashed lines (see Table 1). For clarity, only the H atoms involved in hydrogen bonding have been included. The packing structure exhibits $R_2^2(16)$ and $R_4^4(10)$ graphset motifs.

2. Structural commentary

The molecular structure of the title compound (I) is shown in Fig. 1. The π -conjugated system is nearly planar, with a 2.5 (1)° twist between the aromatic and aldehyde groups. The julolidine ring system comprises three fused rings and one locked nitrogen atom. The C1-O1 and C3-O2 bond lengths are of 1.231 (3) and 1.345 (3) Å, respectively, indicate double- and single-bond character for these bonds. The two fused nonaromatic rings of the julolidine moiety adopt slightly distorted envelope conformations with atoms C9 and C12 displaced from the plane through the remaining ring atoms by 0.654(2)and 0.648 (2) Å, respectively. The intramolecular O2-H2···O1 hydrogen bond forms an S(6) ring motif (Fig. 1 and Table 1) between the phenol and aldehyde groups. Such an intramolecular hydrogen bond is common in salicylaldehyde derivatives, and the metrical parameters are comparable to those for related structures such as hydroxybenzaldehyde (Kirchner et al., 2011).



Figure 1

The molecular structure of the title compound, with the atom labelling. Displacement ellipsoids are drawn at the 30% probability level. The intramolecular $O-H\cdots O$ hydrogen bond is shown as a dashed line (see Table 1).

Table 2 Comparison of selected geometric data for (I) (Å, °) from calculated (DFT) and X-ray data.

Bonds	X-ray	B3LYP/6-311G(d,p)
C1-O1	1.231 (3)	1.231
C3-O2	1.345 (3)	1.345
C1-C2	1.431 (3)	1.431
N1-C5	1.381 (2)	1.381
O1-C1-C2	126.2 (2)	126.22
C1-C2-C3	121.34 (18)	120.25
C11-N1-C10	116.83 (15)	116.81

contain both σ and π character whereas HOMO-1 is dominated by π -orbital density. The LUMO is mainly composed of σ density while LUMO+1 has both σ and π electronic density. The HOMO-LUMO gap was found to be 0.154 a.u. and the frontier molecular orbital energies, E_{HOMO} and E_{LUMO} were f -0.19624 and -0.04201 a.u., respectively.



Electron distribution of the HOMO-1, HOMO, LUMO and the LUMO+1 energy levels for (I).

Table 3 Experimental details.

$\begin{array}{llllllllllllllllllllllllllllllllllll$
Chemical formula $C_{13}H_{15}NO_2$ M_r 217.26 Crystal system, space group Monoclinic, $P2_1/c$ Temperature (K) 100 a, b, c (Å) 8.546 (3), 9.137 (3), 13.662 (4) β (°) 95.984 (6) V (Å ³) 1061.0 (6) Z 4 Radiation type Mo K α μ (mm ⁻¹) 0.09 Crystal size (mm) 0.18 × 0.15 × 0.11 Data collection Bruker SMART CCD area detector
$\begin{array}{ll} M_{\rm r} & 217.26 \\ {\rm Crystal system, space group} & {\rm Monoclinic, $P2_1/c$} \\ {\rm Temperature (K)} & 100 \\ a, b, c~({\rm \AA}) & 8.546~(3), 9.137~(3), 13.662~(4) \\ \beta~(^{\circ}) & 95.984~(6) \\ V~({\rm \AA}^3) & 1061.0~(6) \\ Z & 4 \\ {\rm Radiation type} & {\rm Mo} \ K\alpha \\ \mu~({\rm mm}^{-1}) & 0.09 \\ {\rm Crystal size (mm)} & 0.18 \times 0.15 \times 0.11 \\ \end{array}$
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β (°)95.984 (6) V (Å3)1061.0 (6) Z 4Radiation typeMo K α μ (mm ⁻¹)0.09Crystal size (mm)0.18 × 0.15 × 0.11Data collectionDiffractometerDiffractometerBruker SMART CCD area detector
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Data collection Diffractometer Bruker SMART CCD area detector
Diffractometer Bruker SMART CCD area detector
Absorption correction Multi-scan (SADABS; Sheldrick, 2004)
T_{\min}, T_{\max} 0.985, 0.991
No. of measured, independent and 5787, 2083, 1530
observed $[I > 2\sigma(I)]$ reflections
R _{int} 0.026
$(\sin \theta / \lambda)_{\rm max} ({\rm \AA}^{-1}) \qquad 0.617$
Refinement
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$ 0.066, 0.231, 1.11
No. of reflections 2083
No. of parameters 145
H-atom treatment H-atom parameters constrained
$\Delta \rho_{\text{max}}, \Delta \rho_{\text{min}} \left(e \text{\AA}^{-3} \right) \qquad \qquad 0.64, -0.27$

Computer programs: SMART and SAINT (Bruker, 2003), SHELXTL and SHELXL97 (Sheldrick, 2008).

5. Database survey

A search of the Cambridge Structural Database (CSD, Version 5.37, update May 2016; Groom et al., 2016) gave 121 hits for the julolidine moiety. Of these, six have an OH group in position 8, and four also have a C—N group in position 1. The very similar compound 2-[(2,3,6,7-tetrahydro-1H,5Hbenzo[*ij*]-quinolizin-9-yl)methylene]propanedinitrile (II) reported by Liang et al. (2009) has the aldehydic group in (I) replaced by dicyanovinyl groups and the hydroxy group replaced by hydrogen. The N1–C5 bond length [1.381 (2) Å] in the title compound is longer than in (II) [1.365 (3) Å] due to conjugation with dicyanovinyl group. In the julolidine-1,6dione compound reported by Wu et al. (2007), the N atom of the julolidine moiety lies approximately in the plane of the benzene ring with a deviation of 0.023 (2) Å, similar to that in title compound [0.043 (2) Å], as might be expected for the maximum conjugation normally found for N-atom substituents on benzene rings.

6. Crystallization

2,3,6,7-Tetrahydro-8-hydroxy-1H,5H-benzo[ij]quinolizine-9carboxaldehyde was purchased from Sigma Aldrich and crystallized by slow evaporation of methanol solution over a period of 2-3 days to yield quality crystal suitable for X-ray data collection.

7. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. All H atoms were located from difference-Fourier maps but in the final cycles of refinement they were included in calculated positions and treated as riding atoms: O-H = 0.84 Å, C-H = 0.93-0.98 Å with $U_{iso}(H) = 1.5U_{eq}(O)$ and $1.2U_{eq}(C)$ for other H atoms.

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Crystal structure and DFT study of 8-hydroxy-1,2,3,5,6,7-hexahydropyrido[3,2,1-*ij*]quinoline-9-carbaldehyde

Md. Serajul Haque Faizi, Necmi Dege and Maria L. Malysheva

Computing details

Data collection: *SMART* (Bruker, 2003); cell refinement: *SMART* (Bruker, 2003); data reduction: *SAINT* (Bruker, 2003); program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

8-Hydroxy-1,2,3,5,6,7-hexahydropyrido[3,2,1-ij]quinoline-9-carbaldehyde

Crystal data

C₁₃H₁₅NO₂ $M_r = 217.26$ Monoclinic, $P2_1/c$ a = 8.546 (3) Å b = 9.137 (3) Å c = 13.662 (4) Å $\beta = 95.984$ (6)° V = 1061.0 (6) Å³ Z = 4

Data collection

Bruker SMART CCD area detector diffractometer Radiation source: sealed tube Graphite monochromator phi and ω scans Absorption correction: multi-scan (SADABS; Sheldrick, 2004) $T_{\min} = 0.985, T_{\max} = 0.991$

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.066$ $wR(F^2) = 0.231$ S = 1.112083 reflections 145 parameters 0 restraints F(000) = 464 $D_x = 1.360 \text{ Mg m}^{-3}$ Mo K\alpha radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 1494 reflections $\theta = 2.4-28.1^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ T = 100 KNeedle, colorless $0.18 \times 0.15 \times 0.11 \text{ mm}$

5787 measured reflections 2083 independent reflections 1530 reflections with $I > 2\sigma(I)$ $R_{int} = 0.026$ $\theta_{max} = 26.0^\circ, \ \theta_{min} = 2.4^\circ$ $h = -10 \rightarrow 10$ $k = -8 \rightarrow 11$ $l = -16 \rightarrow 16$

Hydrogen site location: inferred from neighbouring sites H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.1465P)^2 + 0.0857P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.64$ e Å⁻³ $\Delta\rho_{min} = -0.27$ e Å⁻³

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

	x	У	Ζ	$U_{\rm iso}$ */ $U_{\rm eq}$
02	0.73084 (18)	0.25645 (16)	0.53899 (11)	0.0398 (5)
H2	0.691533	0.226583	0.587471	0.060*
O1	0.58740 (18)	0.27075 (18)	0.69921 (11)	0.0460 (5)
N1	0.83012 (18)	0.7058 (2)	0.37861 (11)	0.0288 (5)
C5	0.7743 (2)	0.6300 (2)	0.45521 (12)	0.0242 (5)
C4	0.7801 (2)	0.4768 (2)	0.45803 (13)	0.0269 (5)
C6	0.7080 (2)	0.7115 (2)	0.53120 (14)	0.0277 (5)
C2	0.6517 (2)	0.4816 (2)	0.60979 (14)	0.0282 (5)
C7	0.6468 (2)	0.6333 (2)	0.60434 (14)	0.0282 (5)
H7	0.599815	0.684412	0.652391	0.034*
C3	0.7204 (2)	0.4032 (2)	0.53505 (14)	0.0282 (5)
C10	0.8409 (2)	0.8646 (2)	0.37928 (14)	0.0333 (6)
H10A	0.840281	0.899757	0.312243	0.040*
H10B	0.939673	0.893952	0.415420	0.040*
C11	0.9103 (2)	0.6288 (2)	0.30450 (15)	0.0330 (6)
H11A	1.020944	0.618318	0.327631	0.040*
H11B	0.902214	0.685601	0.244223	0.040*
C13	0.8464 (2)	0.3888 (2)	0.37783 (14)	0.0336 (6)
H13A	0.786287	0.299427	0.365792	0.040*
H13B	0.954748	0.362366	0.398627	0.040*
C8	0.7061 (3)	0.8758 (2)	0.52940 (15)	0.0359 (6)
H8A	0.797720	0.912690	0.569753	0.043*
H8B	0.613101	0.910814	0.557226	0.043*
C1	0.5867 (2)	0.4043 (3)	0.68718 (15)	0.0358 (6)
H1	0.539410	0.460267	0.732734	0.043*
C12	0.8388 (2)	0.4791 (2)	0.28373 (14)	0.0330 (6)
H12A	0.895351	0.428934	0.235680	0.040*
H12B	0.730055	0.489689	0.256334	0.040*
С9	0.7061 (3)	0.9339 (2)	0.42599 (15)	0.0389 (6)
H9A	0.718103	1.039438	0.427433	0.047*
H9B	0.607130	0.910495	0.387813	0.047*

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\mathring{A}^2)

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
02	0.0514 (11)	0.0255 (10)	0.0406 (9)	-0.0001 (6)	-0.0050 (7)	0.0021 (6)
01	0.0469 (11)	0.0421 (11)	0.0470 (10)	-0.0075 (7)	-0.0054 (7)	0.0141 (7)
N1	0.0304 (9)	0.0301 (11)	0.0264 (9)	-0.0015 (7)	0.0044 (7)	-0.0016 (6)
C5	0.0233 (10)	0.0269 (13)	0.0217 (10)	-0.0009 (7)	-0.0011 (8)	-0.0033 (7)

supporting information

C4	0.0289 (12)	0.0273 (12)	0.0233 (11)	0.0013 (7)	-0.0034 (8)	-0.0030 (7)
C6	0.0278 (11)	0.0275 (12)	0.0270 (10)	0.0021 (8)	-0.0007 (8)	-0.0016 (8)
C2	0.0263 (11)	0.0314 (13)	0.0256 (11)	-0.0007 (8)	-0.0041 (8)	0.0020 (8)
C7	0.0289 (11)	0.0307 (13)	0.0244 (10)	0.0010 (8)	-0.0005 (8)	-0.0043 (8)
C3	0.0286 (12)	0.0220 (11)	0.0319 (12)	0.0005 (7)	-0.0069 (9)	-0.0005 (8)
C10	0.0408 (13)	0.0294 (13)	0.0298 (11)	-0.0084 (8)	0.0032 (9)	0.0002 (8)
C11	0.0290 (11)	0.0437 (14)	0.0264 (11)	0.0013 (9)	0.0036 (8)	-0.0040 (9)
C13	0.0372 (13)	0.0295 (13)	0.0332 (12)	0.0069 (8)	-0.0002 (9)	-0.0071 (8)
C8	0.0484 (14)	0.0272 (13)	0.0323 (12)	0.0008 (9)	0.0050 (9)	-0.0049 (8)
C1	0.0291 (12)	0.0411 (14)	0.0353 (12)	-0.0037 (9)	-0.0051 (9)	0.0087 (9)
C12	0.0329 (12)	0.0389 (13)	0.0265 (11)	0.0085 (9)	0.0001 (8)	-0.0099 (8)
C9	0.0543 (14)	0.0252 (12)	0.0368 (12)	-0.0019 (9)	0.0028 (10)	-0.0026 (9)

Geometric parameters (Å, °)

02—C3	1.345 (3)	C10—H10A	0.9700	
O2—H2	0.8200	C10—H10B	0.9700	
01—C1	1.231 (3)	C11—C12	1.512 (3)	
N1—C5	1.381 (2)	C11—H11A	0.9700	
N1-C10	1.454 (3)	C11—H11B	0.9700	
N1-C11	1.461 (2)	C13—C12	1.523 (3)	
C5—C4	1.401 (3)	C13—H13A	0.9700	
C5—C6	1.441 (3)	C13—H13B	0.9700	
C4—C3	1.390 (3)	C8—C9	1.509 (3)	
C4—C13	1.516 (3)	C8—H8A	0.9700	
С6—С7	1.376 (3)	C8—H8B	0.9700	
С6—С8	1.502 (3)	C1—H1	0.9300	
С2—С7	1.389 (3)	C12—H12A	0.9700	
С2—С3	1.423 (3)	C12—H12B	0.9700	
C2—C1	1.431 (3)	С9—Н9А	0.9700	
С7—Н7	0.9300	С9—Н9В	0.9700	
С10—С9	1.513 (3)			
С3—О2—Н2	109.5	N1-C11-H11B	109.5	
C5—N1—C10	121.51 (16)	C12—C11—H11B	109.5	
C5—N1—C11	120.49 (19)	H11A—C11—H11B	108.1	
C10-N1-C11	116.83 (15)	C4—C13—C12	109.65 (17)	
N1-C5-C4	120.54 (16)	C4—C13—H13A	109.7	
N1-C5-C6	118.7 (2)	C12—C13—H13A	109.7	
C4—C5—C6	120.79 (17)	C4—C13—H13B	109.7	
C3—C4—C5	119.32 (17)	C12—C13—H13B	109.7	
C3—C4—C13	119.00 (19)	H13A—C13—H13B	108.2	
C5—C4—C13	121.66 (17)	C6—C8—C9	111.45 (16)	
C7—C6—C5	117.6 (2)	C6—C8—H8A	109.3	
С7—С6—С8	121.81 (17)	C9—C8—H8A	109.3	
C5—C6—C8	120.62 (17)	C6—C8—H8B	109.3	
С7—С2—С3	118.46 (18)	C9—C8—H8B	109.3	
C7—C2—C1	121.34 (18)	H8A—C8—H8B	108.0	

C3—C2—C1	120.2 (2)	01—C1—C2	126.2 (2)
C6—C7—C2	123.07 (18)	01—C1—H1	116.9
C6—C7—H7	118.5	C2—C1—H1	116.9
C2—C7—H7	118.5	C11—C12—C13	110.52 (16)
O2—C3—C4	119.02 (17)	C11—C12—H12A	109.5
O2—C3—C2	120.24 (18)	C13—C12—H12A	109.5
C4—C3—C2	120.7 (2)	C11—C12—H12B	109.5
N1—C10—C9	111.68 (16)	C13—C12—H12B	109.5
N1—C10—H10A	109.3	H12A—C12—H12B	108.1
C9—C10—H10A	109.3	C8—C9—C10	108.80 (19)
N1—C10—H10B	109.3	C8—C9—H9A	109.9
C9—C10—H10B	109.3	C10—C9—H9A	109.9
H10A—C10—H10B	107.9	C8—C9—H9B	109.9
N1—C11—C12 N1—C11—H11A C12—C11—H11A	110.87 (16) 109.5 109.5	C10—C9—H9B H9A—C9—H9B	109.9 108.3

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	$D \cdots A$	<i>D</i> —H… <i>A</i>
02—H2…O1	0.82	1.89	2.621 (2)	148
С9—Н9А…О2 ^і	0.97	2.50	3.324 (3)	143
С9—Н9В…О1 ^{іі}	0.97	2.55	3.438 (3)	152
O2—H2…O1	0.82	1.89	2.621 (2)	148
С9—Н9А…О2 ^і	0.97	2.50	3.324 (3)	143
C9—H9 <i>B</i> ···O1 ⁱⁱ	0.97	2.55	3.438 (3)	152

Symmetry codes: (i) *x*, *y*+1, *z*; (ii) -*x*+1, -*y*+1, -*z*+1.