Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

D. Gayathri,^a D. Velmurugan,^a* K. Ravikumar,^b E. Poonguzhali^c and H. Surya Prakash Rao^c

^aDepartment of Crystallography and Biophysics, University of Madras, Guindy Campus, Chennai 600 025, India, ^bLaboratory of X-ray Crystallography, Indian Institute of Chemical Technology, Hyderabad 500 007, India, and ^cDepartment of Chemistry, Pondicherry University, Pondicherry 605 014, India

Correspondence e-mail: d_velu@yahoo.com

Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.056 wR factor = 0.173 Data-to-parameter ratio = 19.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. 2-(2,6-Diphenylpiperidin-1-yl)ethanol

The title compound, $C_{19}H_{23}NO$, crystallizes with two molecules in the asymmetric unit. The piperidine ring adopts a chair conformation in both molecules. Intramolecular O- $H \cdots N$ and $C-H \cdots \pi$ interactions are observed in only one molecule in the asymmetric unit. The crystal packing is stabilized by O- $H \cdots O$ intermolecular interactions, generating $R_4^4(8)$ rings.

Comment

The strong base, piperidine, is widely used as a building-block molecule in several industries. It is used as an intermediate in agrochemicals, pharmaceuticals and rubber vulcanization accelerators. Several 2,6-substituted piperidine derivatives possess fungicidal, herbicidal and bactericidal properties (Mobio *et al.*, 1989). In view of the above importance, the X-ray crystal structure determination of the title compound, (I), has been undertaken.



Compound (I) crystallizes in the triclinic system with two molecules, *A* and *B*, in the asymmetric unit. The bond lengths and angles are comparable with the corresponding values in a related structure which we published recently (Gayathri *et al.*, 2006). The sums of the bond angles around atom N1 [335.9 and 332.1° for molecules *A* and *B*] indicate sp^3 hybridization. The C1-N1-C6-C7, C5-N1-C6-C7 and N1-C6-C7-O1 torsion angles are 58.5 (2), -68.1 (2) and 174.1 (1)°, respectively, in molecule *A*, and -139.7 (1), 95.9 (2) and 50.7 (2)°, respectively, in molecule *B*.

The piperidine ring adopts a chair conformation in both molecules. The puckering parameters (Cremer & Pople, 1975) are $q_2 = 0.009$ (2) Å, $q_3 = 0.571$ (2) Å, $Q_T = 0.571$ (2) Å and $\varphi = 2.1$ (2)°, respectively, for molecule *A*, and $q_2 = 0.024$ (2) Å, $q_3 = 0.565$ (2) Å, $Q_T = 0.566$ (2) Å and $\varphi = 2.4$ (2)°, respectively, for molecule *B*.

Received 6 October 2006 Accepted 19 October 2006

© 2006 International Union of Crystallography

All rights reserved





The asymmetric unit of (I), showing 30% probability displacement ellipsoids.



Figure 2

The crystal packing of (I), viewed down the a axis. H atoms which are not involved in hydrogen bonding have been omitted for clarity.

Intramolecular O-H···N and C-H··· π interactions are observed only in molecule B and not in molecule A, because of the difference in the torsion angles N1A - C6A - C7A - O1A $[174.1 (1)^{\circ}]$ and N1B-C6B-C7B-O1B [50.7 (2)^{\circ}] in molecules A and B, respectively. In the $O-H \cdots N$ interaction, atom O1B acts as donor to N1B, generating an S(5) ring motif. In the C-H··· π interaction, atom C7*B* acts as a donor to the centroid, Cg, of the C14B-C19B ring, with an H7B2 \cdots Cg separation of 2.659 Å.

The crystal packing of (I) is stabilized by $O-H \cdots O$ intermolecular interactions (Table 2), generating $R_4^4(8)$ rings.

Experimental

To a homogenous solution of 1,3-diphenylpropane (0.992 mmol, 250 mg, 1 equivalent), ethanolamine (9.92 mmol, 605 mg, 10 equivalents) and polyethyleneglycol-200 (10 ml) in a 25 ml Erlenmeyer flask, 85% formic acid (9.92 mmol, 465 mg, 1 ml, 10 equivalents) was added at 273–278 K. The reaction mixture was then irradiated in a domestic microwave oven for 3 min at 370 W, after which time no 1.3diphenylpropane was detected by thin-layer chromatography. After completion of the reaction, the reaction mixture was added to icecold water (25 ml) and the pH of the aqueous solution was adjusted to 11 with 1 N NaOH. The organic compounds were extracted with dichloromethane (DCM: 3×15 ml). The DCM solution was dried over anhydrous Na₂SO₄ and evaporated under reduced pressure. The crude product was subjected to column chromatography on basic alumina by eluting with increasing amounts of ethyl acetate in hexanes (2-10%). After purification, compound (I) was obtained as a white crystalline solid (yield 76%, 212 mg). Single crystals were obtained by recrystallization from about 5% DCM in hexanes (b.p. 333-353 K).

Crystal data

γ

C ₁₉ H ₂₃ NO	V = 1621.7 (2) Å ³
$M_r = 281.38$	Z = 4
Triclinic, P1	$D_x = 1.152 \text{ Mg m}^{-3}$
a = 11.7863 (9) Å	Mo $K\alpha$ radiation
b = 12.2482 (9) Å	$\mu = 0.07 \text{ mm}^{-1}$
c = 12.8201 (10) Å	T = 293 (2) K
$\alpha = 95.077 \ (1)^{\circ}$	Block, colourless
$\beta = 107.637 \ (1)^{\circ}$	$0.26 \times 0.24 \times 0.21 \text{ mm}$
$\gamma = 109.768 \ (1)^{\circ}$	

Data collection

Bruker SMART APEX CCD areadetector diffractometer ω scans Absorption correction: none 18837 measured reflections

Refinement

$w = 1/[\sigma^2(F_o^2) + (0.0901P)^2]$
+ 0.2212P]
where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} < 0.001$
$\Delta \rho_{\rm max} = 0.50 \ {\rm e} \ {\rm \AA}^{-3}$
$\Delta \rho_{\rm min} = -0.19 \text{ e} \text{ Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

C1A-N1A	1.472 (2)	C1B-N1B	1.487 (2)	
C5A - N1A	1.480 (2)	C5B-N1B	1.482 (2)	
C6A - N1A	1.471 (2)	C6B-N1B	1.476 (2)	
C7A-O1A	1.390 (2)	C7B - O1B	1.395 (2)	
C6A - N1A - C1A	111.9 (1)	C6B-N1B-C5B	110.8 (1)	
C6A-N1A-C5A	112.0 (1)	C6B-N1B-C1B	110.5 (1)	
C1A - N1A - C5A	112.0 (1)	C5B-N1B-C1B	111.8 (1)	
N1A-C6A-C7A-O1A	174.1 (1)	N1B-C6B-C7B-O1B	50.7 (2)	
C7A - C6A - N1A - C1A	58.5 (2)	C7B-C6B-N1B-C1B	-139.7 (1)	

7414 independent reflections 5335 reflections with $I > 2\sigma(I)$

 $R_{\rm int} = 0.018$

 $\theta_{\rm max} = 28.0^{\circ}$

Table 2

Hydrogen-bond geometry (Å, $^{\circ}$).

Cg is the centroid of the C14B-C19B ring.

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
$\begin{array}{c} \hline O1B-H1B1\cdots N1B\\ C7B-H7B2\cdots Cg\\ O1A-H1A1\cdots O1B^{i}\\ O1B-H1B1\cdots O1A^{ii} \end{array}$	0.82 0.97 0.82 0.82	2.54 2.66 1.93 2.09	2.884 (2) 3.528 (2) 2.744 (2) 2.816 (2)	107 149 170

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

All H atoms were positioned geometrically and allowed to ride on their parent C and O atoms, with the O-H distance fixed at 0.82 Å, with $U_{\rm iso}({\rm H}) = 1.5 U_{\rm eq}({\rm O})$, and with C-H distances fixed in the range 0.93–0.98 Å, with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm parent C})$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics:

PLATON (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

Financial support from the University Grants Commission (UGC-SAP) and the Department of Science and Technology (DST-FIST), Government of India, is acknowledged by DG and DV for providing facilities to the department. DV thanks CSIR, India, for a Major Research Project.

References

Bruker (2001). SMART (Version 5.625/NT/2000) and SAINT (Version 6.28a). Bruker AXS Inc., Madison, Wisconsin, USA.

Cremer, D. & Pople, J. A. (1975). J. Am. Chem. Soc. 97, 1354-1358.

Gayathri, D., Velmurugan, D., Ravikumar, K., Poonguzhali, E. & Rao, H. S. P. (2006). Acta Cryst. E62, 04169–04171.

Mobio, I. G., Soldatenkov, A. T., Federov, V. O., Ageev, E. A., Sargeeva, N. D., Lin, S., Stashenko, E. E., Prostakov, N. S. & Andreeva, E. I. (1989). *Khim. Farm. Zh.* 23, 421–427. (In Russian).

Nardelli, M. (1995). J. Appl. Cryst. 28, 659.

Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.

Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.