organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(\text{C}-\text{C}) = 0.002 \text{ Å}$ R factor = 0.039 wR factor = 0.086 Data-to-parameter ratio = 17.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

7-(Benzyloxy)spiro[2H-1,3-benzoxazine-2,1'-cyclohexan]-4(3H)-one

In the title compound, $C_{20}H_{21}NO_3$, the dihedral angle between the two benzene rings is 8.79 (7)° and the cyclohexane ring adopts a chair conformation. The molecules are linked by paired N-H···O hydrogen bonds into centrosymmetric $R_2^2(8)$ dimers.

Comment

The molecular structure of (I) is shown in Fig. 1. The dihedral angle between the benzene ring of the 2*H*-benzoxazine system and the phenyl ring is $8.79 (7)^{\circ}$. The bulky, six-membered cyclohexane chair ring in the compound could be used in asymmetric induction, as we have reported previously (Jian *et al.*, 2005).



In the crystal structure, atom N1 acts as a hydrogen donor to atom O1 of a symmetry-related molecule (Table 1), leading to the formation of centrosymmetric $R_2^2(8)$ dimers (Fig. 2).

Experimental

To a mixture of 7-hydroxy-[2H-1,3-benzoxazine-2,1'-cyclohexan]-4(3H)-one (2.0 g, 8.6 mmol) and NaH (0.40 g, 11.7 mmol, 70%) in dimethylformamide (30 ml), benzyl chloride (0.9 ml, 8 mmol) was added dropwise. This mixture was stirred at room temperature for 30 min and then at 333 K overnight. CH₂Cl₂ (30 ml) and water



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Figure 1 The molecule of compound (I) in the crystal. Displacement ellipsoids are drawn at the 50% probability level.

Received 5 September 2005 Accepted 9 September 2005 Online 14 September 2005 (10 ml) were then added to the reaction mixture. The organic layer was washed successively with water (3 × 10 ml), dried over anhydrous Na₂SO₄ and evaporated *in vacuo*. Recrystallization of the resulting white solid from MeOH gave colourless crystals of (I) (m.p. 470–471 K). Spectroscopic analysis: ¹H NMR (500 MHz, CDCl₃, δ , p.p.m.): 1.39–2.11 (*m*, 10H), 5.08 (*s*, 2H), 6.52 (*s*, 1H), 6.66 (*brs*, 1H), 6.67 (*dd*, 1H), 7.35–7.44 (*m*, 5H), 7.83 (*d*, 1H).

 $D_x = 1.283 \text{ Mg m}^{-3}$

Cell parameters from 6953

Mo $K\alpha$ radiation

reflections $\theta = 2.6-27.5^{\circ}$

 $\mu = 0.09 \text{ mm}^{-1}$ T = 295 (2) K

Prism, colourless

 $R_{\rm int}=0.026$

 $\theta_{\rm max} = 27.5^{\circ}$

 $h = -7 \rightarrow 7$

 $k = -11 \rightarrow 11$

 $l = -39 \rightarrow 40$

 $0.26 \times 0.25 \times 0.2 \text{ mm}$

3819 independent reflections

2271 reflections with $I > 2\sigma(I)$

Crystal data

$C_{20}H_{21}NO_3$
$M_r = 323.38$
Monoclinic, $P2_1/c$
a = 6.0904 (2) Å
b = 8.9171 (3) Å
c = 30.8699 (8) Å
$\beta = 93.1470 \ (10)^{\circ}$
$V = 1673.98 (9) \text{ Å}^3$
Z = 4

Data collection

Rigaku R-AXIS RAPID diffractometer ω scans Absorption correction: multi-scan (ABSCOR; Higashi, 1995) T_{min} = 0.976, T_{max} = 0.983 7297 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.037P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.039$	where $P = (F_0^2 + 2F_c^2)/3$
$wR(F^2) = 0.086$	$(\Delta/\sigma)_{\rm max} < 0.001$
S = 1	$\Delta \rho_{\rm max} = 0.14 \ {\rm e} \ {\rm \AA}^{-3}$
3819 reflections	$\Delta \rho_{\rm min} = -0.13 \ {\rm e} \ {\rm \AA}^{-3}$
222 parameters	Extinction correction: SHELXL97
H atoms treated by a mixture of	(Sheldrick, 1997)
independent and constrained	Extinction coefficient: 0.0164 (11)
refinement	

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1 - H1 \cdots O1^i$	0.87 (1)	1.98 (1)	2.8541 (15)	177 (1)
Symmetry code: (i)	-x + 2, -v + 1	-z + 1.		

Atom H1 was found in a difference Fourier map and refined freely. The H atoms of the methylene groups and of the aromatic ring were





A view of the dimer formed by paired N-H···O hydrogen bonds (dashed lines) in the crystal structure of (I). H atoms not involved in hydrogen bonds have been omitted for clarity. [Symmetry code: (i) 2 - x, 1 - y, 1 - z.]

placed in calculated positions, with C–H distances of 0.97 and 0.93 Å, respectively, and were included in the final cycles of the least-squares refinement as riding on their carrier atoms, with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}$ of the corresponding carrier atom.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/ MSC, 2004); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON* (Spek, 2003).

We thank the National Natural Science Foundation of China (grant No. 20272051) and the Teaching and Research Award Programme for Outstanding Young Teachers in Higher Education Institutions of the MoE, China.

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