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Key indicators

Single-crystal X-ray study T = 296 KMean σ (C–C) = 0.003 Å R factor = 0.030 wR factor = 0.061 Data-to-parameter ratio = 15.3

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

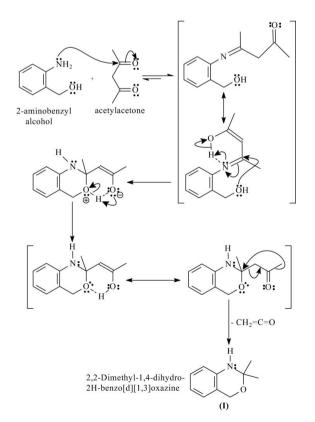
2,2-Dimethyl-1,4-dihydro-2H-3,1-benzoxazine

The title compound, $C_{10}H_{13}NO$, crystallizes with two independent but essentially identical molecules in the asymmetric unit. The molecules are linked by two N-H···O and two C-H···O hydrogen bonds into chains of edge-fused rings in which $R_3^3(10)$ and $R_8^8(33)$ rings alternate.

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Comment

1,3-Oxazines have generated interest as antipsychotic agents and as possible effectors for serotonin and dopamine receptors (Gentles *et al.*, 1991; Petterson *et al.*, 1990; Peglion *et al.*, 1997). In addition, benzoxazines have been evaluated as antimalarial agents (Ren *et al.*, 2001). The title compound, (I), was prepared by reaction of acetylacetone and 2-aminobenzyl alcohol. This is a new method for synthesizing oxazabenzocyclohexanes (see scheme). The steps proposed in the scheme are based on the more stable character of the reaction products with respect to the intermediate products. The structure of (I) has been determined in order to establish both the geometry of the oxazahetero ring and the nature of the supramolecular interactions.



© 2006 International Union of Crystallography All rights reserved Compound (I) (Fig. 1) crystallizes in space group $P2_12_12_1$, with two independent molecules in the asymmetric unit

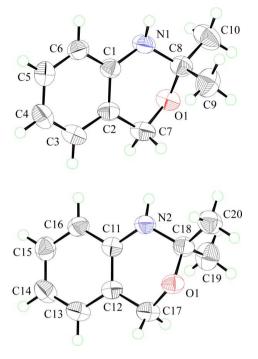


Figure 1

Views of the independent molecules in (I), showing the atom-numbering schemes and 50% probability displacement ellipsoids.

(Fig. 1). The intramolecular dimensions are very similar in the two molecules and the bond distances show no unusual features. The angles at the ring atoms N1 and N2 show considerable expansion from 109° (Table 1).

Molecules of (I) are linked into a chain of edge-fused rings by the combination of two nearly linear $N-H\cdots O$ hydrogen bonds and two rather weak $C-H\cdots O$ hydrogen bonds; geometric parameters are listed in Table 2. Within the asymmetric unit, atom N1 in the first independent molecule acts as hydrogen-bond donor to O2 in the second molecule and, in a similar way, atom N2 in the second independent molecule acts as hydrogen-bond donor to O4 in the symmetry-related 'second' molecule. These hydrogen bonds generate, by translation, a C(5) chain (Bernstein *et al.*, 1995) that runs parallel to the [100] direction, as illustrated in Fig. 2.

Two such chains pass through each unit cell and these two chains are weakly linked by $C-H\cdots O$ hydrogen bonds $(H\cdots O < 3.00 \text{ Å} \text{ and } C-H\cdots O > 110^\circ)$ (Jeffrey, 1997). The aryl C6 atom in the first independent molecule acts as a hydrogen-bond donor to atom O1, thereby generating two $R_3^3(10)$ rings and one $R_8^8(33)$ ring. Propagation by translation of the $R_3^3(10)$ motif linking antiparallel C(4) chains then generates an [010] chain of edge-fused rings, with two $R_3^3(10)$ rings alternating with $R_8^8(33)$ rings. There are no directionspecific aromatic $\pi-\pi$ interactions between adjacent chains of rings in (I).

Experimental

Compound (I) was prepared by combining an *n*-butanol solution (50 ml) containing acetylacetone (1.00 g, 10 mmol) with an *n*-butanol

(50 ml) solution containing 2-aminobenzyl alcohol (1.23 g, 10 mmol). The reaction mixture was stirred for 2 h under reflux. Solid (I) was obtained by allowing the clear reaction mixture to stand overnight. Crystals of (I) suitable for single-crystal X-ray diffraction were grown by slow evaporation of a tetrahydrofuran solution of (I) (yield 75%, m.p. 390–392 K).

Mo $K\alpha$ radiation Cell parameters from 9459

reflections

Block, colourless $0.44 \times 0.34 \times 0.19 \text{ mm}$

2027 reflections with $I > 2\sigma(I)$

 $\begin{array}{l} \theta = 1.6 {-} 26.0^{\circ} \\ \mu = 0.08 \ \mathrm{mm}^{-1} \end{array}$

T = 296 K

 $R_{\rm int}=0.054$

 $\theta_{\rm max} = 26.0^{\circ}$

 $h = -10 \rightarrow 10$

 $k = -10 \rightarrow 10$

 $l=-31\rightarrow 27$

Crystal data

C₁₀H₁₃NO $M_r = 163.21$ Orthorhombic, $P2_12_12_1$ a = 8.2844 (9) Å b = 8.4743 (10) Å c = 25.388 (3) Å V = 1782.4 (4) Å³ Z = 8 $D_x = 1.216$ Mg m⁻³

Data collection

Stoe IPDS-2 diffractometer ω scans Absorption correction: integration (X-RED32; Stoe & Cie, 2002) $T_{min} = 0.969, T_{max} = 0.986$ 9973 measured reflections 3493 independent reflections

Refinement

Refinement on F^2	H atoms treated by a mixture of
$R[F^2 > 2\sigma(F^2)] = 0.030$	independent and constrained
$wR(F^2) = 0.061$	refinement
S = 0.81	$w = 1/[\sigma^2(F_0^2) + (0.0275P)^2]$
3493 reflections	where $P = (F_0^2 + 2F_c^2)/3$
229 parameters	$(\Delta/\sigma)_{\rm max} = 0.001$
	$\Delta \rho_{\rm max} = 0.11 \ {\rm e} \ {\rm \AA}^{-3}$
	$\Delta \rho_{\rm min} = -0.13 \text{ e} \text{ Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

C1-N1	1.398 (2)	C11-N2	1.394 (2)
C7-O1	1.439 (2)	C17-O2	1.430 (2)
C8-O1	1.436 (2)	C18-O2	1.438 (2)
C8-N1	1.447 (2)	C18-N2	1.460 (3)
C7-O1-C8	113.33 (14)	O1-C8-C9	110.92 (15)
C17-O2-C18	113.63 (14)	O1-C8-C10	104.84 (17)
C1-N1-C8	117.64 (16)	N1-C8-C9	112.72 (17)
C11-N2-C18	117.91 (17)	N1-C8-C10	109.30 (18)
O1-C7-C2	111.53 (16)	C9-C8-C10	111.20 (19)
O1-C8-N1	107.51 (16)		

Table 2		
Hydrogen-bond geometry	(Å,	°).

$N1 - H1N \cdots O1^{i}$ 0.882	2 (19) 2.25 (a) a 121 (2)	
		2) 3.131 (2)	176.5 (17)
$N2-H2N\cdots O2^{ii}$ 0.88	(2) 2.20 (2) 3.082 (2)	177 (2)
C20-H20C···O1 ⁱⁱ 0.96	2.86	3.707 (3)	148
$C6{-}H6{\cdot}{\cdot}{\cdot}O2^{iii} \qquad 0.93$	2.68	3.308 (2)	125

In the absence of significant anomalous dispersion effects, 1460 Friedel pairs were averaged. All C-bound H atoms were refined using the riding-model approximation, with C-H = 0.93 Å for aromatic, 0.97 Å for methylene and 0.96 Å for methyl H atoms, and with

 $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$ for aromatic and methylene H atoms, and $U_{\rm iso}({\rm H}) = 1.5 U_{\rm eq}({\rm C})$ for methyl H atoms. N-bound H atoms were refined freely; see Table 2 for geometric parameters.

Data collection: X-AREA (Stoe & Cie, 2002); cell refinement: X-AREA; data reduction: X-RED32 (Stoe & Cie, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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