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Key indicators

Single-crystal X-ray study T = 294 K Mean $\sigma(C-C) = 0.004$ Å R factor = 0.062 wR factor = 0.160 Data-to-parameter ratio = 15.6

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

The solid-state structure of the title compound, $C_{32}H_{40}N_2O_2$, is characterized by benzoxazine units with a *cisoid* geometry, and with the N atoms displaced from the planes of the heterocycles. The two benzoxazine groups are nearly coplanar.

1,4-Bis(8-tert-butyl-6-methyl-4H-1,3-

benzoxazin-3-yl)benzene

Comment

Solid-state structures of benzoxazine derivatives are of interest partly due to their potential to inhibit the enzyme γ aminobutyric acid transaminase (Andrews et al., 1986). The benzoxazine system is also a critical component of a series of resorcinarene derivatives (Arnecke et al., 1995, El Gihani et al., 1995), which have been employed as macrocyclic receptors to encapsulate small molecules or ions selectively (Atwood & Szumna, 2002; Kröck et al., 2004).



The reaction of 1,4-diaminobenzene with 2-tert-butyl-4methylphenol and four equivalents of formaldehyde results in the formation of the title bis-benzoxazine derivative, (I). The bond lengths in the heterocyclic groups of (I) are comparable with those of related compounds (Andrews et al., 1986), with average distances of 1.455 (3) Å for O-C(methylene) and 1.419 (3) Å for N–C(methylene), and an average N–C–O angle of 114.4 (2)°. Atoms N9 and N24 are displaced from the benzoxazine planes, defined by C10-C18/O7 and C25-C33/ O22, by 0.522 (2) and 0.541 (2) Å, respectively. The methylene group located between O and N lies at distances of 0.203 (2) (C8) and 0.212 (3) Å (C23) for C8 and C23, respectively, from the benzoxazine plane. A cisoid arrangement of the tert-butyl groups on the two heterocyclic units is observed.

The solid-state structure of (I) is further characterized by the nearly perpendicular orientation of the benzoxazine planes with respect to the phenylenediamine unit. The angles between the the phenylenediamine plane, defined by atoms C1-C6/N9/N24, and the heterocyclic planes containing O7 and O22 are 81.14(8) and $83.39(8)^\circ$, respectively. This results in a small angle of $2.61 (6)^{\circ}$ between the two benzoxazine systems.

Molecules of (I) pack into columns parallel to the *a* axis, with $C-H\cdots\pi$ contacts between H8A and H8B on the methylene atom C8, and the benzene rings of an adjacent molecule (see Table 2). A second set of $C-H \cdots \pi$ interactions is observed between H23A and H23B on the methylene atom C23, and the benzene rings on another adjacent molecule.

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21975 measured reflections

 $R_{\rm int} = 0.062$

 $\theta_{\rm max} = 25.4^{\circ}$

5180 independent reflections

3379 reflections with $I > 2\sigma(I)$



Figure 1

The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.





View of the crystal packing, showing the columns of (I) aligned parallel to the *a* axis. $C-H\cdots\pi$ interactions are shown as blue lines.

Experimental

A mixture of 1,4-diaminobenzene (0.50 g, 4.62 mmol) and aqueous formaldehyde solution (37% wt, 1.52 g, 18.50 mmol) in ethanol (25 ml) was stirred for 16 h at room temperature. 2-*tert*-Butyl-4-methylphenol (1.52 g, 9.25 mmol) was added and the mixture was then refluxed for 48 h. The reaction mixture was allowed to cool to room temperature and the volatile materials were removed under vacuum. The aqueous mixture obtained was chilled before acidifying with 5 ml of 3N HCl, followed by extraction with diethyl ether (60 ml) and chloroform (40 ml). The combined organic phases were washed with an aqueous potassium carbonate solution (20 ml) and distilled water (20 ml), followed by drying with anhydrous MgSO₄. After evaporation of the solvents under reduced pressure, (I) was purified

by column chromatography on silica gel eluting with a 3:1 hexanesdichloromethane mixture (yield: 0.90 g, 40%; m.p. 422–424 K).

Crystal data

Data collection

Bruker SMART APEX CCD diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 2004) $T_{\min} = 0.965, T_{\max} = 0.996$

Refinement

Table 1

Selected geometric parameters (Å, °).

C1-N9	1.424 (3)	N9-C10	1.464 (3)	
C4-N24	1.427 (3)	O22-C30	1.374 (3)	
O7-C15	1.373 (3)	O22-C23	1.457 (3)	
O7-C8	1.452 (3)	C23-N24	1.417 (4)	
C8-N9	1.419 (3)	N24-C25	1.460 (4)	
C15-07-C8	115.29 (18)	C30-O22-C23	115.3 (2)	
N9-C8-O7	114.25 (19)	N24-C23-O22	114.5 (2)	
C8-N9-C1	117.7 (2)	C23-N24-C4	117.3 (2)	
C8-N9-C10	108.66 (19)	C23-N24-C25	108.8 (2)	
C1-N9-C10	116.4 (2)	C4-N24-C25	116.0 (2)	
N9-C10-C16	111.5 (2)	N24-C25-C31	111.4 (2)	

Table 2

Hydrogen-bond geometry (Å, °).

Cg1 represents the centroid of phenyl ring C1–C6, Cg2 that of ring C11–C16 and Cg3 that of ring C26–C31.

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C8-H8A\cdots Cg3^{i}$	0.97	2.66	3.608 (3)	166
$C8 - H8B \cdot \cdot \cdot Cg1^{i}$	0.97	2.75	3.592 (3)	145
$C23-H23A\cdots Cg2^{ii}$	0.97	2.68	3.650 (3)	174
$C23-H23B\cdots Cg1^{ii}$	0.97	2.95	3.807 (3)	148

Symmetry codes: (i) -x + 1, -y + 1, -z; (ii) -x + 2, -y + 1, -z.

H atoms were positioned geometrically and refined as riding, with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}$ of the parent methylene and aromatic C atoms, and $U_{\rm iso}({\rm H}) = 1.5U_{\rm eq}$ of the parent methyl C atoms, with C–H distances in the range 0.93–0.97 Å.

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT-Plus* (Bruker, 1999); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2000); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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