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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ R factor = 0.066 wR factor = 0.184 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.

2,6-Bis(4-aminophenyl)benzo[1,2-d:5,4-d']dioxazole 1-methylpyrrolidin-2-one trisolvate

The title compound, $C_{20}H_{14}N_4O_2 \cdot 3C_5H_9NO$, crystallizes with three molecules of 1-methyl-2-pyrrolidinone. The amino groups at each end of the fused-ring heterocycle each interact with two of the solvent molecules.

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Comment

2,6-Bis(4-aminophenyl)benzo(1,2-d:5,4-d')dioxazole, which can be used for the synthesis of thermally stable polyheterocycles, has been known for a long time (Evstaf'ev et al., 1970; Preston et al., 1968). A recent study described its synthesis from 1,2,3-trichlorobenzene and its purification (Chen et al., 2003). We have used another method to synthesize this monomer, but our choice of solvent for recrystallization led to the isolation of the compound as a trisolvate, (I) (Fig. 1). The compound has the two aminophenyl substituents of the benzodioxazole group coplanar with the fused-ring system. Bond dimensions are similar to those found in the only other report on this class of 2,6-bis(phenyl)benzo(1,2-d:5,4-d')dioxazole derivatives, viz. the parent unsubstituted compound itself (Fratini, 1978). The amino groups interact with two of the three solvent molecules through N-H···O hydrogen bonds (Table 1). The third solvent molecule makes no significant contacts. The bond dimensions of the three methylpyrrolidinone solvent molecules compare well with those found in the parent compound itself, whose structure was determined at low temperature (Müller et al., 1996).



Experimental

Equimolar quantities of 4,6-diamino-1,3-benzenediol (14.0 g, 0.1 mol) and 4-aminobenzoic acid (13.7 g, 0.1 mol) were dissolved in polyphosphoric acid (50 ml). The mixture was heated at 423 K for five h. The cool solution yielded a pale-yellow powder that was purified by recrystallization from 1-methylpyrrolidone. The recrystallization process took about a month, and yellow prisms were obtained. Analysis calculated for $C_{35}H_{41}N_7O_5$: C 65.71, H 6.46, N 15.33%; found: C 65.74, H 6.48, N 15.35%.

Crystal data

$C_{20}H_{14}N_4O_2 \cdot 3C_5H_9NO$	Z = 2
$M_r = 639.75$	$D_x = 1.293 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 7.586 (2) Å	Cell parameters from 14761
b = 12.525 (2) Å	reflections
c = 17.627 (2) Å	$\theta = 3.0–27.4^{\circ}$
$\alpha = 94.92 \ (1)^{\circ}$	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 92.88 \ (2)^{\circ}$	T = 295 (2) K
$\gamma = 99.19 \ (1)^{\circ}$	Prism, yellow
V = 1643.8 (6) Å ³	$0.39 \times 0.27 \times 0.20$ mm

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Data collection

Rigaku R-AXIS RAPID IP diffractometer ω scans Absorption correction: multi-scan (ABSCOR; Higashi, 1995) T_{min} = 0.654, T_{max} = 0.983 15592 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.066$ $wR(F^2) = 0.184$ S = 1.027333 reflections 427 parameters H-atom parameters constrained 7333 independent reflections 4014 reflections with $I > 2\sigma(I)$ $R_{int} = 0.038$ $\theta_{max} = 27.4^{\circ}$ $h = -9 \rightarrow 9$ $k = -16 \rightarrow 16$ $l = -22 \rightarrow 22$

$$\begin{split} &w = 1/[\sigma^2(F_{\rm o}^{-2}) + (0.0898P)^2 \\ &+ 0.1894P] \\ &where \ P = (F_{\rm o}^{-2} + 2F_{\rm c}^{-2})/3 \\ (\Delta/\sigma)_{\rm max} = 0.001 \\ \Delta\rho_{\rm max} = 0.50 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.19 \ {\rm e} \ {\rm \AA}^{-3} \end{split}$$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
N1-H1n1···O3	0.86	2.25	3.092 (3)	166
$N1-H1n2\cdots O4^{i}$	0.86	2.21	3.060 (3)	172
N4-H4 $n1$ ···O3 ⁱⁱ	0.86	2.34	3.122 (3)	152
$N4-H4n2\cdots O4$	0.86	2.37	3.130 (3)	148
	0.00	2107	01100 (0)	110

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

The C-bound H atoms were placed at calculated positions (C– $H_{aromatic} = 0.93$ Å, C– $H_{methylene} = 0.97$ Å and C– $H_{methyl} = 0.96$ Å), and were included in the refinement in the riding-model approximation, with $U_{iso}(H)$ set to $1.2U_{eq}(C)$ for the aromatic and methylene C atoms, and to $1.5U_{eq}(C)$ for the methyl groups. The methyl groups were rotated to fit the electron density. The N-bound H atoms were treated similarly [N–H = 0.86 Å and $U_{iso}(H) = 1.2U_{eq}(C)$].

Data collection: *RAPID-AUTO* (Rigaku Corporation, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSC, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.



Figure 1

ORTEPII (Johnson, 1976) plot of (I). Displacement ellipsoids are drawn at the 70% probability level, and H atoms are drawn as spheres of arbitrary radii.

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