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Key indicators

Single-crystal X-ray study T = 100 KMean σ (C–C) = 0.002 Å R factor = 0.034 wR factor = 0.098 Data-to-parameter ratio = 12.4

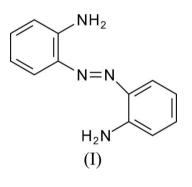
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The title compound, $C_{12}H_{12}N_4$, is a *trans*-azobenzene which crystallizes with an inversion center at the midpoint of the N=N bond.

trans-Bis(2-aminophenyl)diazene

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Comment

The family of azobenzenes has been investigated intensively due to the their remarkable photochemically induced *trans* \rightarrow *cis* isomerism. Although the structures of many azobenzenes have been determined, some of the fundamental and synthetically important members of this family, such as the title compound, (I), (Fig. 1) have not been characterized. This report describes the 100 K structure of this highly functional member of the azobenzene family.



In (I) there is an inversion center at the mid-point of the N=N bond. The donating oamino substituents of (I) allow for numerous potential derivatives to be prepared (Skrabel & Hohl-Blumer, 1976; de Mendoza *et al.*, 1981) and they are also expected to influence the electronic structure of the azobenzene system. The N1=N1ⁱ [symmetry code: (i) -x, -y, 1 - z] bond length of 1.2641 (18) Å in (I) is slightly greater than the average value of 1.257 Å for azobenzene N=N bonds found in the Cambridge Structural Database (Version 5.27; Allen, 2002). In addition, the relatively short C1-N1 bond length of 1.4131 (14) Å suggests some delocalization of electron density into the N=N π^* orbital, although this needs to be confirmed by spectroscopic measurements. The molecular conformation of (I) appears to be stabilized by a weak intramolecular N-H···N hydrogen bond (Table 1).

Experimental

The title compound was prepared by surfactant-mediated aerial oxidation of *o*-phenylenediamine in toluene by literature methods (de Mendoza *et al.*, 1981). Crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of a toluene solution.

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organic papers

Crystal data

 $C_{12}H_{12}N_4$ $M_r = 212.26$ Monoclinic, C2/c a = 22.278 (6) Å b = 5.8647 (16) Å c = 8.254 (2) Å $\beta = 106.357$ (3)° V = 1034.8 (5) Å³ Z = 4

Data collection

Siemens SMART CCD diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Siemens, 1996) $T_{\min} = 0.983, T_{\max} = 0.990$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.034$ $wR(F^2) = 0.098$ S = 1.031003 reflections 81 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$N2-H2A\cdots N1$	0.884 (18)	2.330 (17)	2.7064 (15)	105.7 (13)

The N-bound H atoms were located in difference maps and their positions and $U_{\rm iso}$ values were freely refined. The C-bound H atoms were located in difference maps and then repositioned in idealized locations (C-H = 0.93 Å) and refined as riding, with $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm C)$.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve

 $D_x = 1.362 \text{ Mg m}^{-3}$ $D_m = 1.32 \text{ Mg m}^{-3}$ $D_m \text{ measured by flotation in ????}$ Mo K\alpha radiation $\mu = 0.09 \text{ mm}^{-1}$ T = 100 (2) KBlock, orange-red $0.20 \times 0.10 \times 0.07 \text{ mm}$

3927 measured reflections 1003 independent reflections 929 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.021$ $\theta_{\text{max}} = 25.9^{\circ}$

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0584P)^2 \\ &+ 0.6927P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} < 0.001 \\ \Delta\rho_{\text{max}} &= 0.27 \text{ e } \text{ Å}^{-3} \\ \Delta\rho_{\text{min}} &= -0.18 \text{ e } \text{ Å}^{-3} \end{split}$$

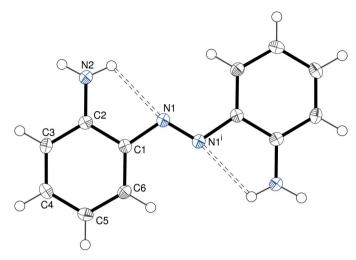


Figure 1

View of the molecular structure of (I), showing 50% displacement ellipsoids (arbitrary spheres for the H atoms). [Symmetry code: (i) -x, -y, 1 - z.]

structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Siemens, 1996); software used to prepare material for publication: *SHELXTL*.

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