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Hong-Qing Hao^a and Hui Zhang^b*

^aDepartment of Chemistry, Xiamen University, Xiamen 361005, People's Republic of China, and ^bState Key Laboratory for Physical Chemistry of Solid Surfaces, Department of Chemistry, Xiamen University, Xiamen 361005, People's Republic of China

Correspondence e-mail: huizhang@jingxian.xmu.edu.cn

Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.004 Å R factor = 0.079 wR factor = 0.230 Data-to-parameter ratio = 15.8

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

N,*N*'-Bis[4-(dimethylamino)benzylidene]ethane-1,2-diamine

The title compound, $C_{20}H_{26}N_4$, shows a distinctly V-shaped structure. In the crystal structure, parallel rings and short intermolecular contacts indicate the presence of aromatic stacking interactions. A twofold rotation axis passes through the mid-point of the central C–C bond.

Comment

The crystal structure of N,N'-bis(4-(dimethylamino)benzylidene)-1,2-diaminocyclohexane has been reported previously (Jones *et al.*, 1998); we report here the crystal structure of N,N'-bis[4-(dimethylamino)benzylidene]-1,2-diaminoethane, (I). The molecular structure of (I) is illustrated in Fig.1, with selected bond lengths and bond angles listed in Table 1.



The molecular structure of (I) exhibits a distinct V shape. A twofold rotation axis passes through the mid-point of the $C10-C10^{i}$ bond (symmetry code as in Table 1). The C9-N2 bond length of 1.251 (4) Å is consistent with double-bond character. Other bond lengths and angles in (I) lie within the expected ranges. In the molecular packing, the distance between parallel rings is 3.597 Å, which indicates the presence of aromatic stacking interactions.

Experimental

A mixture of 1,2-diaminoethane (1 mmol) and 4-dimethylaminobenzaldehyde (2 mol) in ethanol (40 ml) was refluxed for 3 h and then cooled to room temperature with stirring. The solution was filtered and concentrated to dryness. The resulting residue was washed with ethanol and dried in a vacuum. The yellow product was dissolved in dichloromethane and the solution was evaporated slowly in a dark place at room temperature. After two weeks, red block crystals were obtained. The overall yield was 75%. CHN elemental analysis on the red platelets: found (calculated) for $C_{20}H_{26}N_4$ (%): C 74.35 (74.50), H 8.10 (8.13), N 17.55 (17.38). IR (KBr): 3423, 3030, 1645, 1630, 1450, 1380, 1280, 1115, 850, 803, 771, 680, 637 cm⁻¹.

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Crystal data

 $\begin{array}{l} C_{20}H_{26}N_4 \\ M_r = 322.45 \\ \text{Monoclinic, } C2/c \\ a = 24.151 \ (10) \text{ \AA} \\ b = 6.503 \ (3) \text{ \AA} \\ c = 12.390 \ (5) \text{ \AA} \\ \beta = 114.869 \ (8)^{\circ} \\ V = 1765.5 \ (12) \text{ \AA}^3 \\ Z = 4 \end{array}$

Data collection

Bruker APEX diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{min} = 0.978, T_{max} = 0.992$ 4624 measured reflections 1721 independent reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.1093P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.079$	+ 2.2949P]
$wR(F^2) = 0.230$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.08	$(\Delta/\sigma)_{\rm max} < 0.001$
1721 reflections	$\Delta \rho_{\rm max} = 0.32 \text{ e } \text{\AA}^{-3}$
109 parameters	$\Delta \rho_{\rm min} = -0.27 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

 $D_x = 1.213 \text{ Mg m}^{-3}$

Cell parameters from 1200

 $0.31 \times 0.30 \times 0.11 \text{ mm}$

1347 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

reflections

 $\mu = 0.07 \text{ mm}^{-1}$

T = 298 (2) K

 $\theta = 3.2 - 25^{\circ}$

Plate, red

 $R_{\rm int} = 0.028$

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

 $h = -20 \rightarrow 29$

 $k = -8 \rightarrow 7$

 $l = -15 \rightarrow 8$

Table 1

Selected geometric parameters (Å, °).

C9-N2 C10-N2	1.251 (4) 1.440 (3)	C1-N1 C2-N1	1.432 (4) 1.428 (4)
$C10 - C10^{1}$	1.506 (6)		
N2-C10-C10 ⁱ C9-N2-C10 C3-N1-C2	111.5 (2) 116.6 (2) 121.2 (2)	C3-N1-C1 C2-N1-C1	120.9 (3) 117.3 (3)
C7-C6-C9-N2 C5-C6-C9-N2 C4-C3-N1-C2	177.8 (3) -2.7 (4) -7.6 (4)	C4-C3-N1-C1 C10-C10 ⁱ -N2-C9	-178.3 (3) -130.5 (3)

Symmetry code: (i) $-x + 1, y, -z + \frac{3}{2}$.

The aromatic and aliphatic H atoms were placed at calculated positions and refined using the riding-model approximation, with C-H = 0.93 - 0.97 Å and $U_{iso} = 1.2U_{eq}$ (C).

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve



Figure 1

ORTEPII (Johnson, 1976) plot of (I). Displacement ellipsoids are drawn at the 30% probability level. [Symmetry code: (i) -x + 1, y, $-z + \frac{3}{2}$.]

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*.

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