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#### **Key indicators**

Single-crystal X-ray study T = 298 K Mean  $\sigma$ (C–C) = 0.002 Å R factor = 0.052 wR factor = 0.167 Data-to-parameter ratio = 16.0

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

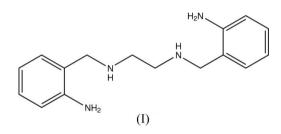
# The asymmetric unit of the title compound, $C_{16}H_{22}N_4$ ,

N,N'-Bis(2-aminobenzyl)ethane-1,2-diamine

### Comment

lographic inversion center.

The title compound, (I), a Schiff base ligand, has been synthesized as a chemical precursor for a variety of simple and macrocyclic multidentate ligands and metal complexes. Several crystal structures of complexes using (I) have been reported, with  $Cd^{II}$  (Ansell *et al.*, 1983), and with Ni<sup>II</sup> (Anacona *et al.*, 2005; Taylor *et al.*, 2004).



Compound (I) crystallizes with the molecule lying on a crystallographic inversion center, located at the mid-point of the C8–C8<sup>i</sup> bond [symmetry code: (i) 1 - x, 1 - y, -z] (Fig. 1). The conformation of the central chain is described by torsion angles C6–C7–N2–C8 = 175.26 (13)°, C7–N2–C8–C8<sup>i</sup> = 81.3 (2)° and N2–C8–C8<sup>i</sup>–N2<sup>i</sup> constrained by symmetry to 180°. This *trans-gauche-trans* conformation stabilized in the solid state for (I) is less common than the all-*trans* conformation generally found in aliphatic systems. On the other hand, all the N atoms in (I) may coordinate to a metal center, giving another conformation for this ligand. These observations suggest that (I) is a highly flexible molecule, with almost free rotation about all  $\sigma$  bonds.

The molecule shows a weak intramolecular hydrogen bond involving N1 and N2 and an intermolecular contact formed between amine group N1 and a symmetry-related N2 group (Table 1), forming a chain along the *a* axis (Fig. 2). Finally, the H atom of the N2 amine group does not participate in any hydrogen bonds with a symmetry-related molecule. This preferential and more extended hydrogen-bonding formation, using the NH<sub>2</sub> substituents rather than the NH functionalities of the central diaminoethane group, probably gives a more stable crystal structure.

#### **Experimental**

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Published procedures were used (Elizondo-Martínez *et al.*, 2006; Obregón-Solís *et al.*, 2001) to prepare *N*,*N*'-bis(2-nitrobenzyl)-1,2-

# organic papers

diaminoethane. A Pd/C-catalysed selective reduction reaction of this compound using  $NH_2NH_2\cdot H_2O$  afforded (I) together with a 2*H*-indazole compound, [2,2'-(ethane-1,2-diyl)bis(2*H*-indazole)] (Rodríguez de Barbarín *et al.*, 2006). Suitable crystals were obtained as yellow prisms from ethanol by slow evaporation at 298 K. The solid was characterized by IR and <sup>1</sup>H NMR spectroscopy and elemental analysis, which are in agreement with the X-ray crystal structure.

Z = 4

 $D_x = 1.241 \text{ Mg m}^{-3}$ 

 $0.60 \times 0.60 \times 0.45 \ \mathrm{mm}$ 

3 standard reflections

every 97 reflections

intensity decay: 3%

Mo  $K\alpha$  radiation

 $\mu = 0.08 \text{ mm}^{-1}$ 

T = 298 (2) K

Prism, yellow

 $R_{\rm int} = 0.062$ 

 $\theta_{\rm max} = 27.5^{\circ}$ 

#### Crystal data

 $\begin{array}{l} C_{16}H_{22}N_4 \\ M_r = 270.38 \\ \text{Orthorhombic, } Pccn \\ a = 9.974 \ (3) \ \text{\AA} \\ b = 22.729 \ (7) \ \text{\AA} \\ c = 6.384 \ (2) \ \text{\AA} \\ V = 1447.2 \ (8) \ \text{\AA}^3 \end{array}$ 

#### Data collection

Bruker P4 diffractometer  $\omega$  scans Absorption correction: none 3702 measured reflections 1661 independent reflections 1191 reflections with  $I > 2\sigma(I)$ 

#### Refinement

-	
Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0954P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.052$	+ 0.1803P]
$wR(F^2) = 0.167$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} < 0.001$
1661 reflections	$\Delta \rho_{\rm max} = 0.25 \ {\rm e} \ {\rm \AA}^{-3}$
104 parameters	$\Delta \rho_{\rm min} = -0.21 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	Extinction correction: SHELXTL-
independent and constrained	Plus
refinement	Extinction coefficient: 0.060 (14)

#### Table 1

Hydrogen-bond geometry (Å, °).

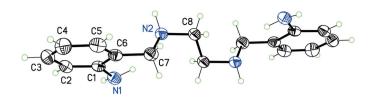
$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} N1 - H1B \cdots N2 \\ N1 - H1A \cdots N2^{i} \end{array}$	0.91 (2)	2.32 (2)	2.994 (2)	131.3 (17)
	0.88 (2)	2.39 (2)	3.264 (2)	172.3 (16)

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

The atoms H1*A*, H1*B* and H2*A* bonded to N atoms were found in difference maps and refined isotropically with free coordinates. H atoms bonded to C atoms were included in calculated positions, and refined using the riding method, with C–H distances constrained to 0.93 (aromatic CH) and 0.97 Å (methylene CH<sub>2</sub>) and  $U_{iso}(H) = 1.2U_{eq}(carrier C)$ .

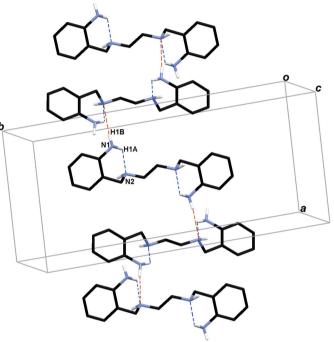
Data collection: XSCANS (Siemens, 1996); cell refinement: XSCANS; data reduction: XSCANS; program(s) used to solve structure: SHELXTL-Plus (Sheldrick, 1997); program(s) used to refine structure: SHELXTL-Plus; molecular graphics: SHELXTL-Plus and Mercury (Macrae et al., 2006); software used to prepare material for publication: SHELXTL-Plus.

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## Figure 1

Molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radius. Unlabeled atoms are related to labeled atoms by (1 - x, 1 - y, -z).



#### Figure 2

Molecular packing for (I), showing the hydrogen-bonding scheme (dashed bonds). H atoms not involved in this network have been omitted.

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