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## Key indicators

Single-crystal X-ray study  
 $T = 292$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å  
 $R$  factor = 0.056  
 $wR$  factor = 0.152  
Data-to-parameter ratio = 17.6For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.*N,N'*-Bis(3,4-dimethoxybenzylidene)ethylenediamineThe molecule of the title compound,  $\text{C}_{20}\text{H}_{24}\text{N}_2\text{O}_4$ , has crystallographic inversion symmetry. In the crystal structure, the molecules are linked *via* weak intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds into a three-dimensional structure.Received 12 December 2005  
Accepted 16 January 2006

## Comment

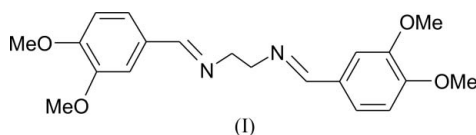
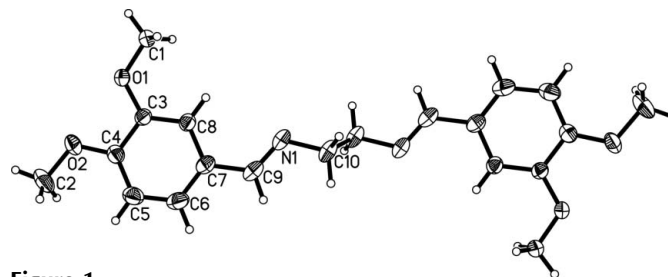
Ethylenediamine and its derivatives play an important role in developing molecular architectures in coordination chemistry (Brown *et al.*, 2001; Dapporto *et al.*, 2000). Similarly, the condensation products of ethylenediamine with aldehydes, *i.e.* Schiff bases, form an interesting class of ligands for transition metal complexes (Goswami & Eichhorn, 1999; Henson *et al.*, 1999). Recently, we have reported a Schiff base compound derived from the condensation of ethylenediamine with 2-nitrobenzaldehyde, which has been structurally characterized (Li *et al.*, 2005). As an extension of this work, the crystal structure of the title compound, (I), has been determined.In (I) (Fig. 1), two 3,4-dimethoxybenzylidene groups are bridged by the ethylenediamine fragment *via* two  $\text{C}=\text{N}$  double bonds in a roughly linear geometry. The torsion angle  $\text{C}6-\text{C}7-\text{C}9-\text{N}1$  of  $172.34$  ( $15$ ) $^\circ$  indicates an extended configuration. The molecule is disposed about a centre of inversion at the mid-point of the central  $\text{C}-\text{C}$  bond. The  $\text{C}9=\text{N}1$  bond length of  $1.260$  ( $2$ ) Å is comparable to the corresponding values of  $1.250$  ( $2$ ) Å observed in bis(2-nitrobenzaldehyde)ethylenediamine (Li *et al.*, 2005). All other bond lengths are within normal ranges (Allen *et al.*, 1987).In the crystal packing, the molecules are linked *via* weak intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds, forming a three-dimensional structure (Table 1 and Fig. 2).

Figure 1

The structure of centrosymmetric (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme. Unlabelled atoms are related to labelled atoms by the symmetry code  $(1-x, 2-y, 1-z)$ .

## Experimental

Ethylenediamine (1 mmol, 60 mg) and 3,4-dimethoxybenzaldehyde (2 mmol, 332 mg) were dissolved in methanol (10 ml) at 323 K. The mixture was stirred for 10 min to give a clear and colourless solution. After the solution had been allowed to stand in air for 8 d, colourless crystals formed, in about 87% yield, on slow evaporation of the solvent. Analysis found: C 67.35, H 6.91, N 7.85%;  $C_{20}H_{24}N_2O_4$  requires C 67.39, H 6.79, N 7.86%.

### Crystal data

$C_{20}H_{24}N_2O_4$	$D_x = 1.269 \text{ Mg m}^{-3}$
$M_r = 356.41$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 2443 reflections
$a = 13.126 (3) \text{ \AA}$	$\theta = 2.7\text{--}23.0^\circ$
$b = 8.531 (2) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$c = 8.393 (2) \text{ \AA}$	$T = 292 (2) \text{ K}$
$\beta = 97.125 (4)^\circ$	Block, colourless
$V = 932.6 (4) \text{ \AA}^3$	$0.20 \times 0.10 \times 0.10 \text{ mm}$
$Z = 2$	

### Data collection

Bruker APEX area-detector diffractometer	2111 independent reflections
$\varphi$ and $\omega$ scans	1415 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$R_{\text{int}} = 0.142$
$T_{\text{min}} = 0.897$ , $T_{\text{max}} = 0.995$	$\theta_{\text{max}} = 27.5^\circ$
10357 measured reflections	$h = -16 \rightarrow 16$
	$k = -10 \rightarrow 10$
	$l = -10 \rightarrow 10$

### Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.056$	$w = 1/[\sigma^2(F_o^2) + (0.0823P)^2]$
$wR(F^2) = 0.152$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.94$	$(\Delta/\sigma)_{\text{max}} < 0.001$
2111 reflections	$\Delta\rho_{\text{max}} = 0.23 \text{ e \AA}^{-3}$
120 parameters	$\Delta\rho_{\text{min}} = -0.21 \text{ e \AA}^{-3}$

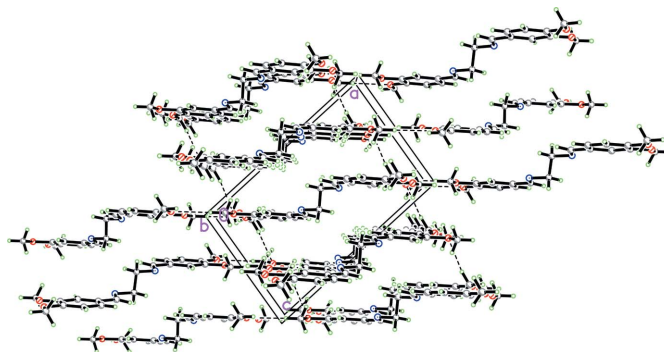
**Table 1**

Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D\text{--}H\cdots A$	$D\text{--}H$	$H\cdots A$	$D\cdots A$	$D\text{--}H\cdots A$
$C2\text{--}H2C\cdots O1^i$	0.96	2.56	3.490 (2)	164
$C1\text{--}H1B\cdots O2^{ii}$	0.96	2.57	3.521 (2)	171

Symmetry codes: (i)  $x, -y + \frac{1}{2}, z + \frac{1}{2}$ ; (ii)  $-x, -y + 1, -z$ .

The H atoms were included in calculated positions and refined using a riding-model approximation, with aromatic C—H = 0.93  $\text{\AA}$ , methylene C—H = 0.97  $\text{\AA}$  and methyl C—H = 0.96  $\text{\AA}$ , and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  for aromatic and methylene, and  $1.5U_{\text{eq}}(\text{C})$  for



**Figure 2**

The crystal packing of (I), viewed along the  $b$  axis. Dashed lines show intermolecular C—H...O hydrogen bonds.

methyl H atoms. The high value of  $R_{\text{int}}$  is due to the weak precision of single-crystal reflection data.

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINTE* (Bruker, 2003); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997a); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997a); molecular graphics: *SHELXTL* (Sheldrick, 1997b); software used to prepare material for publication: *SHELXL97*.

We thank the Scientific Research Foundation for Returned Overseas Chinese Scholars, the State Education Ministry, and the Natural Science Foundation of Hubei Province, People's Republic of China, for research grant No. 2003ABB010.

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