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Key indicators

Single-crystal X-ray study T = 294 K Mean σ (C–C) = 0.003 Å R factor = 0.045 wR factor = 0.139 Data-to-parameter ratio = 15.4

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

N,N'-Bis(3,4,5-trimethoxybenzylidene)hydrazine

In the title compound, $C_{20}H_{24}N_2O_6$, the molecule possesses a crystallographically imposed centre of symmetry at the midpoint of the N–N bond. The central C—N–N—C linkage is therefore planar.

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Comment

To date, a large number of azine compounds containing both a diimine linkage and N–N bonding have been synthesized because they are used in coordination chemistry (Xu *et al.*, 1997; Armstrong *et al.*, 1998; Kesslen *et al.*, 1999; Kundu *et al.*, 2005; Zheng *et al.*, 2005). In this context, an X-ray crystal structure determination of the title compound, (I), was carried out.



In the molecule of (I), there is an inversion centre at the mid-point of the N1-N1ⁱ bond [symmetry code: (i) -x + 1, -y + 2, -z + 1] and the C=N-N=C linkage is planar (Fig. 1). The N-N bond length, 1.419 (3) Å, is slightly greater than that observed in related azine compounds (Liu *et al.*, 2004; Sengül *et al.*, 2004; Xu *et al.*, 2005). The C=N-N angle, 112.2 (2)°, is similar to that observed in *N*,*N*'-bis(3-hydroxy-4-methoxybenzylidene)hydrazine (Duan *et al.*, 2005), but



Figure 1

© 2006 International Union of Crystallography All rights reserved View of the molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level for non-H atoms. [Symmetry code: (A) 1 - x, 2 - y, 1 - z.]

organic papers

significantly smaller than the ideal value of 120° expected for an sp^2 N atom, as a consequence of the repulsion between the N lone pairs and the adjacent C=N bond. The O atoms of the methoxy groups attached to the benzene ring do not deviate substantially from the plane of the ring; a maximum deviation of 0.0699 (2) Å is observed for atom O2.

Experimental

The title compound was synthesized by the reaction of 3,4,5trimethoxybenzaldehyde with hydrazine hydrate in refluxing ethanol (Liu *et al.*, 2004). Single crystals suitable for X-ray analysis were obtained by slow evaporation at 298 K of a tetrahydrofuran solution.

Z = 4

 $D_x = 1.287 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

Block, light vellow

 $0.22 \times 0.18 \times 0.14 \text{ mm}$

 $\mu = 0.10 \text{ mm}^{-1}$

T = 294 (2) K

Crystal data

 $\begin{array}{l} C_{20}H_{24}N_2O_6\\ M_r = 388.41\\ \text{Monoclinic, } C2/c\\ a = 29.879\ (10)\ \text{\AA}\\ b = 4.9185\ (18)\ \text{\AA}\\ c = 14.040\ (5)\ \text{\AA}\\ \beta = 103.754\ (6)^\circ\\ V = 2004.2\ (12)\ \text{\AA}^3 \end{array}$

Data collection

Bruker SMART 1000 CCD areadetector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.979, T_{\max} = 0.987$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.046$ $wR(F^2) = 0.139$ S = 1.012004 reflections 130 parameters 2004 independent reflections 1280 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.033$ $\theta_{\text{max}} = 26.2^{\circ}$

5201 measured reflections

H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0802P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.16 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.22 \text{ e} \text{ Å}^{-3}$

Table 1

			0		
Selected	geometric	parameters	(Å, '	°)	

O1-C4	1.365 (2)	O3-C6	1.370 (2)
O2-C5	1.375 (2)	$N1-N1^i$	1.419 (3)
C1 N1 N1 ⁱ	112.2 (2)		
CI-NI-NI	112.2 (2)		
$N1^{i} - N1 - C1 - C2$	-179.4 (2)		
Symmetry code: (i) $-x$ -	+1, -y + 2, -z + 1		

All H atoms were positioned geometrically and refined using a riding model. Constrained distances: 0.93 Å for Csp^2 -H and 0.96 Å for Csp^3 -H. U_{iso} (H) values were fixed at $1.2U_{eq}$ (aromatic C) and $1.5U_{eq}$ (methyl C).

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

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