

Acta Crystallographica Section E

## Structure Reports

Online

ISSN 1600-5368

## 2,3-Dimethyl-N-[(E)-2,4,5-trimethoxybenzylidene]aniline

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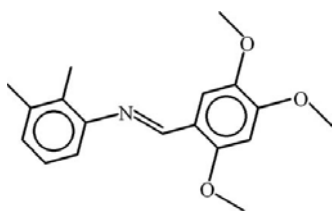
Received 29 June 2010; accepted 1 July 2010

Key indicators: single-crystal X-ray study;  $T = 296$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.045;  $wR$  factor = 0.139; data-to-parameter ratio = 19.4.

In the title compound,  $\text{C}_{18}\text{H}_{21}\text{NO}_3$ , the  $\text{C}=\text{N}$  bond has a *trans* conformation and the benzene rings are oriented at a dihedral angle of  $61.32(6)^\circ$ . The C atoms of the three methoxy groups are all roughly coplanar with their attached ring [deviations =  $0.219(2)$ ,  $-0.097(2)$  and  $-0.137(2)$  Å]. In the crystal, a weak  $\text{C}-\text{H}\cdots\pi$  interaction may help to establish the packing.

## Related literature

For background information on Schiff bases and related crystal structures, see: Tahir *et al.* (2010a,b); Tariq *et al.* (2010).



## Experimental

## Crystal data

 $\text{C}_{18}\text{H}_{21}\text{NO}_3$  $M_r = 299.36$ Triclinic,  $P\bar{1}$  $a = 7.0040(2)$  Å $b = 11.0396(4)$  Å $c = 11.1585(4)$  Å $\alpha = 73.941(1)^\circ$  $\beta = 76.022(2)^\circ$  $\gamma = 82.079(1)^\circ$  $V = 802.24(5)$  Å<sup>3</sup> $Z = 2$ Mo  $K\alpha$  radiation $\mu = 0.08$  mm<sup>-1</sup> $T = 296$  K $0.32 \times 0.14 \times 0.12$  mm

## Data collection

Bruker Kappa APEXII CCD

diffractometer

Absorption correction: multi-scan

(SADABS; Bruker, 2005)

 $T_{\min} = 0.980$ ,  $T_{\max} = 0.985$ 

13855 measured reflections

3957 independent reflections

2935 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.024$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.045$  $wR(F^2) = 0.139$  $S = 1.07$ 

3957 reflections

204 parameters

H-atom parameters constrained

 $\Delta\rho_{\max} = 0.23$  e Å<sup>-3</sup> $\Delta\rho_{\min} = -0.16$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

 $\text{Cg1}$  is the centroid of the  $\text{C1}-\text{C6}$  ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C16}-\text{H16B}\cdots\text{Cg1}^i$	0.96	2.99	3.5694 (19)	120

Symmetry code: (i)  $-x + 2, -y + 1, -z + 1$ .

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINTE* (Bruker, 2009); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

The authors acknowledge the provision of funds for the purchase of the diffractometer and encouragement by Dr Muhammad Akram Chaudhary, Vice Chancellor, University of Sargodha, Pakistan.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5535).

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