# organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.004 Å R factor = 0.056 wR factor = 0.154 Data-to-parameter ratio = 17.4

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

# 1-{2-[(4-Nitrobenzylidene)amino]phenyl}-3-phenyl-thiourea

The molecular structure of the title compound,  $C_{20}H_{16}N_4O_2S$ , is stabilized by  $N-H\cdots N$  and weak  $C-H\cdots S$  intramolecular interactions. The crystal packing is stabilized by  $N-H\cdots S$  and  $C-H\cdots O$  intermolecular interactions.

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## Comment

Considerable attention has been focused on the design of host molecules that recognize and sense anion species selectively through visible, electrochemical and optical responses (Martinez-Manez & Sancenon, 2003). The development of colorimetric anion sensing is very important and increasingly appreciated since naked eye detection can offer qualitative and quantitative information (Gale, 2001). Color changes, as signaling an event detected by the naked eye, are widely used owing to the low cost or lack of equipment required.

The bond lengths and bond angles of the title compound, (I), are comparable to the literature values (Allen *et al.*, 1987). The C2–C1–N1–O1  $[-4.7 (4)^{\circ}]$  and C2–C1–N1–O2  $[173.2 (3)^{\circ}]$  torsion angles indicate a slight deviation of the nitro group from the plane of the benzene ring. The dihedral angles made by the benzene rings C1–C6 and C8–C13 with C15–C20 are 41.7 (1) and 69.9 (1)°, respectively, and that between C1–C6 and C8–C13 is 28.4 (1)°.



The molecular geometry is stabilized by an  $N-H\cdots N$  hydrogen bond and a weak  $C-H\cdots S$  interaction. The crystal packing is stabilized by  $N-H\cdots S$  and  $C-H\cdots O$  intermolecular interactions.

### **Experimental**

1-{2-[4-Nitrobenzylidene)amino}phenyl}-3-phenylthiourea was synthesized by Schiff base condensation between 1-(2-aminophenyl)thiourea and 4-nitrobenzaldehyde. To a solution of 1-(2-aminophenyl)thiourea (0.3 g, 1.2 mmol) in methanol (25 ml), 4-

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#### Figure 1



nitrobenzaldehyde (0.206 g, 1.2 mmol) in methanol (25 ml) was added with stirring. The resulting mixture was heated at reflux for 3 h and cooled to room temperature. The solid product was collected by filtration and washed with cold methanol. The microcrystalline compound was recrystallized from hot chloroform; yellow coloured crystals suitable for X-ray diffraction were obtained on slow evaporation. Yield: 71%; m.p. 476 K.

#### Crystal data

 $\begin{array}{l} C_{20}H_{16}N_4O_2S\\ M_r = 376.43\\ Triclinic, P\overline{1}\\ a = 7.2705 \ (6) \\ b = 11.0178 \ (9) \\ c = 12.0137 \ (9) \\ a = 84.776 \ (1)^\circ\\ \beta = 85.675 \ (1)^\circ \end{array}$ 

#### Data collection

Bruker SMART APEX CCD diffractometer Absorption correction: none 10737 measured reflections  $\gamma = 75.405 (1)^{\circ}$   $V = 926.06 (13) \text{ Å}^3$  Z = 2Mo K\alpha radiation  $\mu = 0.20 \text{ mm}^{-1}$  T = 293 (2) K $0.28 \times 0.23 \times 0.22 \text{ mm}$ 

4234 independent reflections 3312 reflections with  $I > 2\sigma(I)$  $R_{int} = 0.021$  Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.056$  $wR(F^2) = 0.155$ S = 1.034234 reflections 244 parameters H-atom parameters constrained 
$$\begin{split} &\Delta\rho_{max}=0.64\ e\ {\mbox{\AA}}^{-3}\\ &\Delta\rho_{min}=-0.42\ e\ {\mbox{\AA}}^{-3} \end{split}$$

# Table 1 Hydrogen-bond geometry (Å, $^\circ).$

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N3-H3A\cdots N2$ $C12-H12\cdots S1$ $N4-H4\cdots S1^{i}$ $C18-H18\cdots O2^{ii}$	0.86 0.93 0.86 0.93	2.18 2.51 2.46 2.56	2.640 (2) 3.188 (2) 3.301 (2) 3.472 (5)	113 130 165 166
			()	

Symmetry codes: (i) -x + 2, -y, -z + 1; (ii) -x, -y, -z.

All H atoms were refined using a riding model with C-H = 0.93 Å, N-H = 0.86 Å and  $U_{iso}(H) = 1.2U_{eq}(C,N)$ .

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

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