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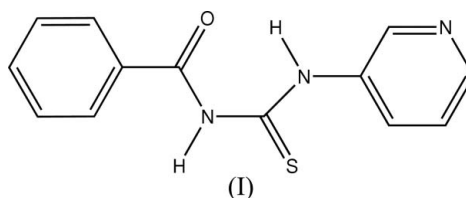
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Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.043
 wR factor = 0.106
Data-to-parameter ratio = 13.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.*N*-Benzoyl-*N'*-(3-pyridyl)thioureaThe molecule of the title compound, $\text{C}_{13}\text{H}_{11}\text{N}_3\text{OS}$, is non-planar; the dihedral angles between the thiourea and pyridine planes and between the thiourea and benzene planes are $41.28(8)$ and $27.47(8)^\circ$, respectively.Received 12 June 2006
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Comment

Thiourea and its derivatives have attracted much attention because of their applications in fields such as herbicides, insecticides and plant-growth regulators (Anthiline & Taketa, 1982).



The molecular structure of the title thiourea derivative, (I), is shown in Fig. 1. The molecule is non-planar; the dihedral angles between the thiourea and pyridine planes and between the thiourea and benzene planes are $41.28(8)$ and $27.47(8)^\circ$, respectively. The carbonyl and thiocarbonyl groups are located on the opposite sides of the $\text{N}3-\text{C}6$ bond. The carbonyl group forms an intramolecular hydrogen bond with the $\text{N}2$ -imino group (Table 1). This is similar to that found in *N*-benzoyl-*N'*-(2-hydroxyethyl)thiourea (Zhang *et al.*, 2006).

$\pi-\pi$ stacking is observed in the crystal structure of (I) (Fig. 2). The centroid-to-centroid separation between parallel benzene rings related by an inversion center at $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$ is $3.7275(17)$ Å; the centroid-to-centroid separation between parallel pyridine rings related by an inversion center at $(0, -\frac{1}{2}, 0)$ is $3.7767(18)$ Å.

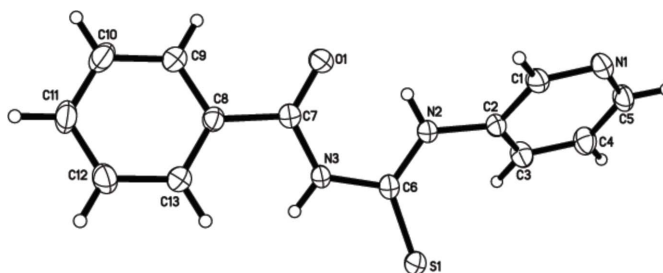


Figure 1
The molecular structure of (I), with 30% probability displacement ellipsoids (arbitrary spheres for H atoms).

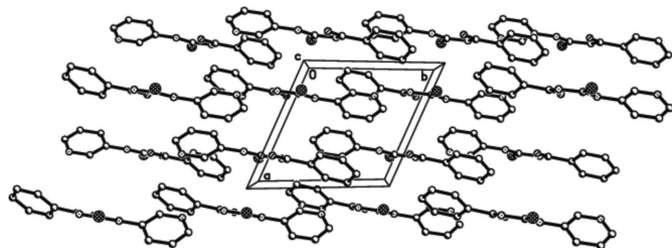


Figure 2
The packing of (I); H atoms have been omitted for clarity.

Experimental

The reaction of 2.82 g (0.02 mol) of benzoyl chloride, 2.29 g (0.03 mol) of ammonium thiocyanate and 1.70 g (0.02 mol) of 3-aminopyridine in 25 ml of CH₂Cl₂ under solid-liquid phase-transfer catalysis conditions (0.37 g of 3% polyethylene glycol-400) gave the title compound. The solid product was filtered off, washed with CH₂Cl₂ and water in turn, and dried under reduced pressure. Recrystallization from chloroform solution gave single crystals of (I).

Crystal data

C ₁₃ H ₁₁ N ₃ OS	$V = 627.0 (3) \text{ \AA}^3$
$M_r = 257.31$	$Z = 2$
Triclinic, $P\bar{1}$	$D_x = 1.363 \text{ Mg m}^{-3}$
$a = 7.828 (2) \text{ \AA}$	Mo $K\alpha$ radiation
$b = 8.688 (2) \text{ \AA}$	$\mu = 0.25 \text{ mm}^{-1}$
$c = 10.892 (3) \text{ \AA}$	$T = 293 (2) \text{ K}$
$\alpha = 111.785 (3)^\circ$	Prism, colorless
$\beta = 92.152 (4)^\circ$	$0.25 \times 0.15 \times 0.06 \text{ mm}$
$\gamma = 111.499 (4)^\circ$	

Data collection

Bruker SMART CCD area-detector diffractometer	3239 measured reflections
φ and ω scans	2263 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 2002)	1879 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.930$, $T_{\max} = 0.985$	$R_{\text{int}} = 0.013$
	$\theta_{\text{max}} = 25.5^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.106$
 $S = 1.01$
 2263 reflections
 163 parameters
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0512P)^2 + 0.1913P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.19 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.21 \text{ e \AA}^{-3}$

Table 1

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2—H2 \cdots O1	0.86	1.87	2.597 (2)	141
N3—H3A \cdots N1 ⁱ	0.86	2.34	3.164 (3)	162

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

H atoms were placed in calculated positions with C—H = 0.93 \AA and N—H = 0.86 \AA , and refined in riding mode with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C,N})$.

Data collection: SMART (Bruker, 2003); cell refinement: SAINT (Bruker, 2003); data reduction: SAINT; 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: SHELXTL.

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