

***N*-(4-Methylbenzoyl)-*N'*-(4-nitrophenyl)thiourea**

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Key indicators

Single-crystal X-ray study

T = 293 K

Mean $\sigma(\text{C}-\text{C}) = 0.004 \text{ \AA}$

R factor = 0.046

wR factor = 0.128

Data-to-parameter ratio = 12.9

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

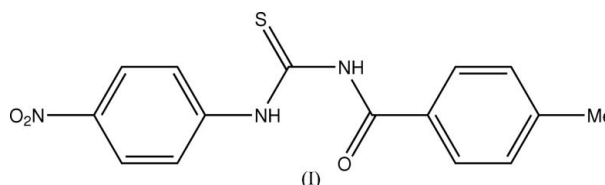
The title compound, $\text{C}_{15}\text{H}_{13}\text{N}_3\text{O}_3\text{S}$, adopts a *trans*–*cis* configuration of the tolyl and nitrophenyl groups, respectively, with respect to the thiono S atom across the thiourea C–N bonds. The molecule is stabilized by intermolecular hydrogen bonds, forming one-dimensional chains parallel to the *a* axis.

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Comment

Some thiourea derivatives have shown promising biological activity against bacteria, fungi and cells (Baruah *et al.*, 2002). The title compound, (I), is isostructural with *N*-benzoyl-*N'*-phenylthiourea, (II) (Yamin & Yusof, 2003), and *N*-(4-chloro-3-nitrophenyl)-*N'*-(4-nitrobenzoyl)thiourea, (III) (Yusof *et al.*, 2006), but having nitro and methyl groups attached to the benzene rings (Fig. 1).



The bond lengths and angles (Table 1) are normal (Allen *et al.*, 1987) and in agreement with those in (II) and (III). The central carbonylthiourea (S1/N1/N2/C7/O1/C8), 4-methylphenyl (C1–C6/C15) and 4-nitrophenyl (C9–C14/N3/O2/O3) groups are all planar, with a maximum deviation of 0.151 (2) Å for atom O2 from the least-squares plane of the 4-nitrophenyl fragment. The central carbonylthiourea fragment makes dihedral angles of 24.67 (10) and 6.65 (10)° with the 4-methylphenyl and 4-nitrophenyl fragments, respectively. The two aryl rings are inclined to each other at an angle of 28.12 (12)°.

There is one intramolecular hydrogen bond, N2–H2A···O1 (Table 2), forming a pseudo-six-membered ring (O1···H2A–N2–C8–N1–C7–O1). In the crystal structure, the molecules are linked by intermolecular interactions, N1–H1A···S1 and C5–H5···O2 (symmetry codes as in Table 2), forming one-dimensional chains along the *a* axis (Fig. 2).

Experimental

To a stirred acetone solution (75 ml) of 4-methylbenzoyl chloride (2.0 g, 13 mmol) and ammonium thiocyanate (0.98 g, 13 mmol), a solution of 4-nitrophenylamine (1.79 g, 13 mmol) in acetone (30 ml) was added dropwise. The resulting mixture was refluxed for 1 h. The resulting solution was poured into a beaker containing some ice

blocks. The light-yellow precipitate was filtered off, washed with distilled water and cold ethanol, and then dried under vacuum. Good quality crystals were obtained by recrystallization from chloroform (yield 76%, 2.53 g; m.p. 460.29–462.93 K).

Crystal data

$C_{15}H_{13}N_3O_3S$ $Z = 4$
 $M_r = 315.34$ $D_x = 1.430 \text{ Mg m}^{-3}$
 Monoclinic, $P2_1/c$ Mo $K\alpha$ radiation
 $a = 12.237 (3) \text{ \AA}$ $\mu = 0.24 \text{ mm}^{-1}$
 $b = 6.3181 (15) \text{ \AA}$ $T = 293 (2) \text{ K}$
 $c = 19.362 (5) \text{ \AA}$ Slab, light yellow
 $\beta = 101.976 (4)^\circ$ $0.40 \times 0.31 \times 0.11 \text{ mm}$
 $V = 1464.4 (6) \text{ \AA}^3$

Data collection

Bruker SMART APEX CCD area-detector diffractometer 7216 measured reflections
 2572 independent reflections
 ω scans 2004 reflections with $I > 2\sigma(I)$
 Absorption correction: multi-scan $R_{int} = 0.027$
 (SADABS; Bruker, 2000) $\theta_{max} = 25.0^\circ$
 $T_{min} = 0.911, T_{max} = 0.974$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0714P)^2 + 0.407P]$
 $R[F^2 > 2\sigma(F^2)] = 0.046$ where $P = (F_o^2 + 2F_c^2)/3$
 $wR(F^2) = 0.128$ $(\Delta/\sigma)_{max} < 0.001$
 $S = 1.00$ $\Delta\rho_{max} = 0.19 \text{ e \AA}^{-3}$
 2572 reflections $\Delta\rho_{min} = -0.19 \text{ e \AA}^{-3}$
 200 parameters
 H-atom parameters constrained

Table 1

Selected geometric parameters ($\text{\AA}, ^\circ$).

S1—C8	1.649 (2)	N1—C8	1.392 (3)
N1—C7	1.383 (3)	N2—C8	1.339 (3)
C7—N1—C8	129.90 (19)	N2—C8—S1	128.49 (18)
N2—C8—N1	114.04 (19)	N1—C8—S1	117.48 (16)

Table 2

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2—H2A \cdots O1	0.86	1.91	2.640 (3)	142
N1—H1A \cdots S1 ⁱ	0.86	2.75	3.490 (2)	145
C5—H5 \cdots O2 ⁱⁱ	0.93	2.57	3.474 (4)	165

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

After their location in a difference map, all H atoms were positioned geometrically and allowed to ride on their parent C or N atoms, with C—H = 0.93–0.96 \AA , N—H = 0.86 \AA and $U_{iso}(H) = 1.2$ (CH_2 and NH) or 1.5 (CH_3) times $U_{eq}(C,N)$.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); 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, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

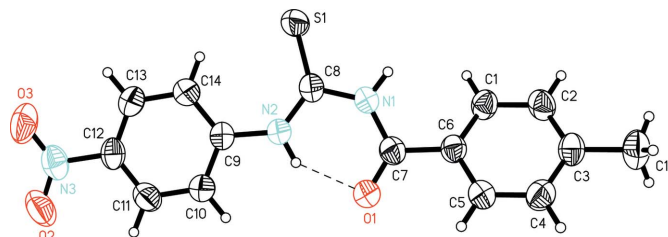


Figure 1

The molecular structure of the title compound, (I), shown with 50% probability displacement ellipsoids. The dashed line indicates a hydrogen bond.

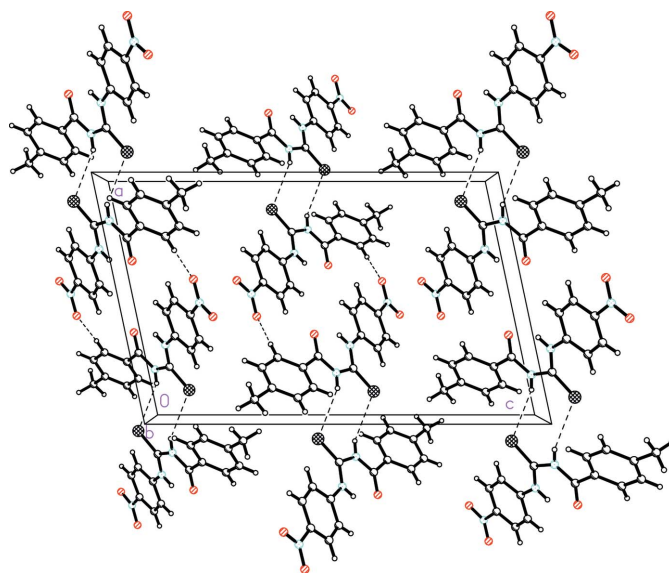


Figure 2

Packing diagram of (I), viewed down the b axis. Dashed lines denote the N—H \cdots S and C—H \cdots O hydrogen bonds.

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