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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 = 0.034
 wR factor = 0.098
 Data-to-parameter ratio = 15.0

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

5-(3-Methylphenoxyethyl)-4-phenyl-1,2,4-triazole-3-thione

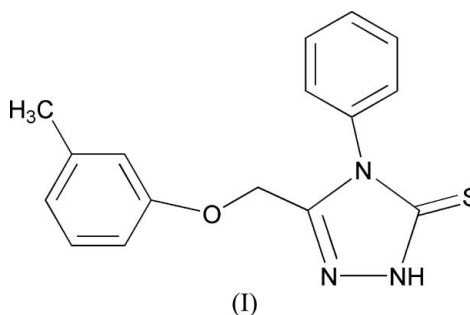
In the title compound, $\text{C}_{16}\text{H}_{15}\text{N}_3\text{OS}$, molecules form inversion-related dimers *via* $\text{N}-\text{H}\cdots\text{S}$ hydrogen bonds. The structure is further stabilized by intermolecular $\pi-\pi$ stacking interactions down the a axis.

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Comment

Substituted triazole derivatives display significant biological activity including antimicrobial (Holla *et al.*, 1998), analgesic (Turan-Zitouni *et al.*, 1999), antitumor (Demirbas *et al.*, 2002), antihypertensive (Paulvannan *et al.*, 2000) and antiviral activities (Kritsanida *et al.*, 2002). The biological activity is closely related to the structure, possibly being due to the presence of the $-\text{N}-\text{C}=\text{S}$ unit (Omar *et al.*, 1986). We are interested in the synthesis and biological activity of aryl-oxyacetyl hydrazide derivatives and report here the synthesis and crystal structure of the title compound, (I) (Fig. 1).



Compound (I) contains a planar triazolethione ring (mean deviation from the ring plane = 0.0008 Å). The dihedral angle between the C11–C16 phenyl ring and the triazole ring is 72.6 (2)°. In the crystal structure, molecules are linked into inversion-related dimers in the ac plane by $\text{N}-\text{H}\cdots\text{S}$ hydrogen bonds (Table 1 and Fig. 2).

The C11–C16 and C11ⁱ–C16ⁱ phenyl rings [symmetry code: (i) $2 - x, -y, 1 - z$] are parallel by symmetry, with a centroid-centroid distance of 4.070 Å. They thus form slipped-parallel dimers (Tsuzuki *et al.*, 2002; Hobza *et al.*, 1996); these intermolecular $\pi-\pi$ interactions further stabilize the structure.

Experimental

The synthesis of the title compound was carried out by refluxing a solution of 1-(3-methylphenoxyacetyl)-4-phenylthiosemicarbazide (10 mmol) in 2 M NaOH for 2 h. Colorless single crystals of (I) were obtained by slow evaporation of an ethanol solution over a period of about one week.

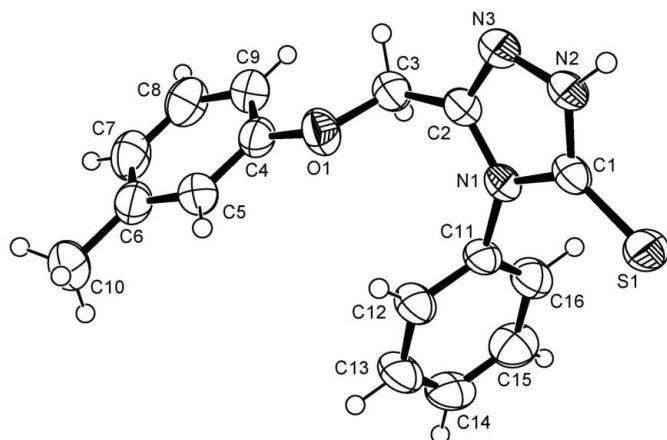


Figure 1
The molecular structure of (I), with displacement ellipsoids drawn at the 50% probability level.

Crystal data

$C_{16}H_{15}N_3OS$	$V = 750.2 (2) \text{ \AA}^3$
$M_r = 297.37$	$Z = 2$
Triclinic, $P\bar{1}$	$D_x = 1.316 \text{ Mg m}^{-3}$
$a = 6.9579 (12) \text{ \AA}$	Mo $K\alpha$ radiation
$b = 9.8402 (18) \text{ \AA}$	$\mu = 0.22 \text{ mm}^{-1}$
$c = 11.686 (2) \text{ \AA}$	$T = 293 (2) \text{ K}$
$\alpha = 74.127 (3)^\circ$	Block, colorless
$\beta = 78.039 (3)^\circ$	$0.55 \times 0.47 \times 0.32 \text{ mm}$
$\gamma = 82.196 (3)^\circ$	

Data collection

Bruker SMART CCD area-detector diffractometer	6291 measured reflections
φ and ω scans	2920 independent reflections
Absorption correction: multi-scan (SADABS; Bruker, 1998)	2337 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.890$, $T_{\max} = 0.934$	$R_{\text{int}} = 0.016$
	$\theta_{\max} = 26.0^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0466P)^2 + 0.192P]$
$R[F^2 > 2\sigma(F^2)] = 0.034$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.098$	$(\Delta/\sigma)_{\max} = 0.001$
$S = 1.03$	$\Delta\rho_{\max} = 0.17 \text{ e \AA}^{-3}$
2920 reflections	$\Delta\rho_{\min} = -0.16 \text{ e \AA}^{-3}$
195 parameters	
H atoms treated by a mixture of independent and constrained refinement	

Table 1

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N2-H2\cdots S1^i$	0.86 (2)	2.40 (2)	3.2588 (14)	176.6 (17)

Symmetry code: (i) $-x+3, -y, -z$.

The N-bound H atom was located in a difference Fourier map and refined freely, with an isotropic displacement parameter. All other H

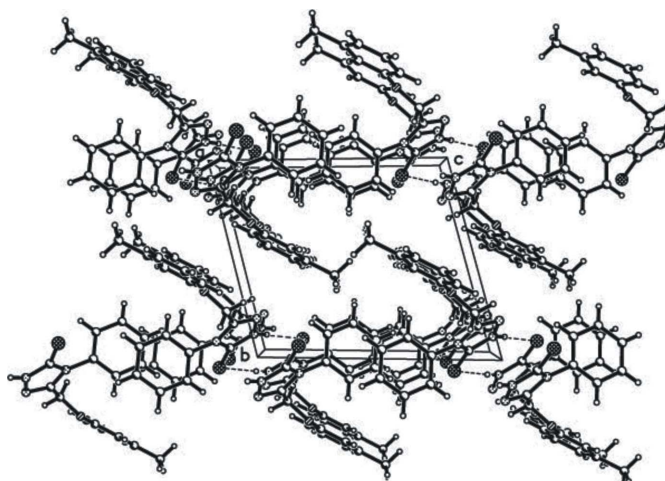


Figure 2
Packing diagram for (I), with hydrogen bonds shown as dashed lines.

atoms were positioned geometrically, with $C-H = 0.93-0.97 \text{ \AA}$, and refined using a riding model, with $U_{\text{iso}}(H) = 1.5U_{\text{eq}}(\text{methyl C})$ and $1.2U_{\text{eq}}(C)$.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

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