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3-(3-Methoxybenzoyl)-1,1-diphenylthiourea

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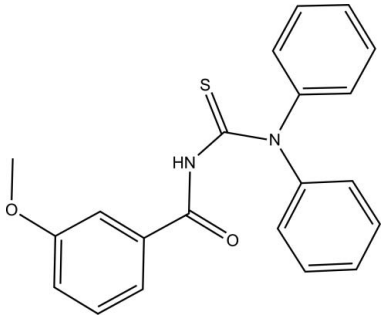
Key indicators: single-crystal X-ray study; T = 298 K; mean σ(C–C) = 0.003 Å; R factor = 0.038; wR factor = 0.110; data-to-parameter ratio = 15.1.

The thiono and carbonyl groups in the title compound, $C_{21}H_{18}N_2O_2S$, adopt an *anti* disposition with respect to the central C—N bond. The diphenylamine rings are twisted relative to each other by a dihedral angle of 82.55 (10)°. The 3-methoxybenzoyl fragment is twisted relative to one of the diphenylamine rings, forming a dihedral angle of 74.04 (9)°. In the crystal, pairs of intermolecular N—H···S hydrogen bonds link the molecules into centrosymmetric dimers, forming columns parallel to the *a* axis.

Related literature

For related structures and background references, see: Al-abbasi *et al.* (2010); Al-abbasi & Kassim (2011); Md Nasir *et al.* (2011). For metal complexes of benzoylthioureas, see: Circu *et al.* (2009); Weiqun *et al.* (2005). For the synthetic procedure, see: Hassan *et al.* (2008). For a standard bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

$C_{21}H_{18}N_2O_2S$
M_r = 362.43
Triclinic, $P\overline{1}$
a = 6.056 (3) Å
b = 12.895 (7) Å
c = 13.344 (7) Å
α = 112.796 (9)°
β = 100.336 (10)°
γ = 97.951 (10)°
V = 920.0 (8) Å^3
Z = 2
Mo $K\alpha$ radiation
μ = 0.19 mm^{-1}
T = 298 K
0.52 × 0.23 × 0.03 mm

Data collection

Bruker SMART APEX CCD area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2000)
T_{min} = 0.906, T_{max} = 0.994
10189 measured reflections
3612 independent reflections
2663 reflections with $I > 2\sigma(I)$
R_{int} = 0.029

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.038
$wR(F^2)$ = 0.110
S = 1.04
3612 reflections
240 parameters
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.17 e Å^{-3}
$\Delta\rho_{min}$ = −0.27 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1A···S1^{i}	0.87 (2)	2.54 (2)	3.380 (2)	163.6 (17)

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

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: JH2306).

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supplementary materials

Acta Cryst. (2011). E**67**, o1947-o1948 [**doi:10.1107/S1600536811025785**]

3-(3-Methoxybenzoyl)-1,1-diphenylthiourea

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Comment

Benzoylthiourea compounds act as a chelating ligands by deprotonation of the amide group and coordinate *via* a thiolate form through the S and O atoms to form neutral complexes with cobalt (Weiqun *et al.*, 2005) and platinum (Circu *et al.*, 2009).

The title compound, I, is a thiourea derivative analogous to our previously reported compounds (Md Nasir *et al.*, 2011; Al-abbasi & Kassim, 2011). The thiono and the carbonyl groups are *trans* positioned with respect to N1—C8 bond with C7N1C8S1 torsion angle of -129.10 (16)°. The methoxy group is coplanar with the parent benzene ring with the largest deviation from the mean planes of (O1/C1/C2/C3/C4/C5/C6/C7) of -0.051 (2)° for C7 and the dihedral angle between the same plane and the thioamide mean planes (S1/N1/N2/C8) is 87.45 (8)°.

In the crystal structure, Intermolecular N1—H1A···S1 hydrogen bond link the molecules into a centrosymmetric dimers forming channel parallel to the *a*-axis.

Experimental

The title compound was prepared according to a previously reported procedure (Al-abbasi *et al.*, 2010) using 3-methoxybenzoyl chloride and diphenylamine as the appropriate starting materials. A yellowish crystal, suitable for X-ray crystallography, was obtained by a slow evaporation from ethanol solution at room temperature (yield 75%).

Refinement

Hydrogen atom of the amide group was determined from the difference Fourier map and N—H was initially fixed at 0.86(0.01) Å and allowed to be refined on the parent N atom with U_{iso}(H) = 1.2U_{eq}(N). All other H atoms were positioned geometrically with C—H bond lengths in the range 0.93 - 0.97 Å and refined in the riding model approximation with U_{iso}(H)=1.2U_{eq}(C,*N*), except for methyl group where U_{iso}(H)= 1.5U_{eq}(C).

Figures

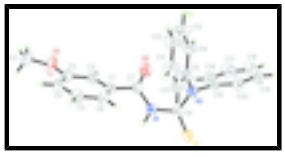

Fig. 1. The molecular structure of the title compound with displacement ellipsoids drawn at the 50% probability level.

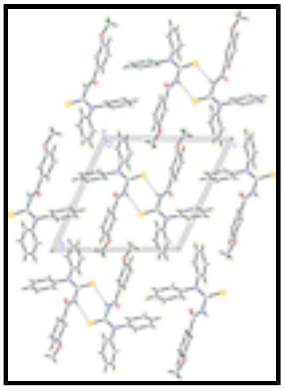

Fig. 2. A packing diagram of the title compound viewed down the *a*-axis showing the intermolecular hydrogen bonds N1—H1···S1 (*-x + 1, -y + 1, -z + 1*) centrosymmetric dimers along *a*-axis.

3-(3-Methoxybenzoyl)-1,1-diphenylthiourea

Crystal data

$C_{21}H_{18}N_2O_2S$	$Z = 2$
$M_r = 362.43$	$F(000) = 380$
Triclinic, $P\overline{1}$	$D_x = 1.308$ Mg m^{-3}
Hall symbol: -P 1	Melting point = 412.15–410.15 K
$a = 6.056$ (3) Å	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
$b = 12.895$ (7) Å	Cell parameters from 3612 reflections
$c = 13.344$ (7) Å	$\theta = 6.4$–$55.2°$
$\alpha = 112.796$ (9)°	$\mu = 0.19$ mm^{-1}
$\beta = 100.336$ (10)°	$T = 298$ K
$\gamma = 97.951$ (10)°	Plate, yellow
$V = 920.0$ (8) Å^3	0.52 × 0.23 × 0.03 mm

Data collection

Bruker SMART APEX CCD area-detector diffractometer	3612 independent reflections
Radiation source: fine-focus sealed tube	2663 reflections with $I > 2\sigma(I)$
graphite	$R_{int} = 0.029$
ω scan	$\theta_{max} = 26.0°$, $\theta_{min} = 1.7°$
Absorption correction: multi-scan (*SADABS*; Bruker, 2000)	$h = -7 \rightarrow 7$
$T_{min} = 0.906$, $T_{max} = 0.994$	$k = -15 \rightarrow 15$
10189 measured reflections	$l = -16 \rightarrow 16$

Refinement

Refinement on F^2	Primary atom site location: structure-invariant direct methods
Least-squares matrix: full	Secondary atom site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.038$	Hydrogen site location: inferred from neighbouring sites

$wR(F^2) = 0.110$	H atoms treated by a mixture of independent and constrained refinement
$S = 1.04$	$w = 1/[\sigma^2(F_o^2) + (0.0578P)^2 + 0.1546P]$ where $P = (F_o^2 + 2F_c^2)/3$
3612 reflections	$(\Delta/\sigma)_{max} < 0.001$
240 parameters	$\Delta\rho_{max} = 0.17$ e Å^{-3}
0 restraints	$\Delta\rho_{min} = -0.27$ e Å^{-3}

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on F^2, conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2. The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating *R*-factors(gt) *etc*. and is not relevant to the choice of reflections for refinement. *R*-factors based on F^2 are statistically about twice as large as those based on *F*, and *R*- factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
S1	0.28578 (10)	0.32442 (4)	0.43411 (4)	0.05684 (19)
O1	0.0958 (3)	0.92764 (11)	0.40494 (14)	0.0667 (4)
O2	−0.0666 (2)	0.49768 (10)	0.27758 (12)	0.0554 (4)
N1	0.2897 (3)	0.47668 (11)	0.34630 (12)	0.0403 (4)
N2	0.0990 (2)	0.28974 (11)	0.22465 (10)	0.0372 (3)
C6	0.4600 (3)	0.67429 (15)	0.30353 (15)	0.0458 (4)
H6	0.5408	0.6164	0.2821	0.055*
C5	0.5442 (3)	0.78217 (16)	0.30918 (17)	0.0504 (5)
H5	0.6817	0.7962	0.2898	0.060*
C4	0.4287 (3)	0.86899 (15)	0.34293 (15)	0.0458 (4)
H4	0.4889	0.9413	0.3471	0.055*
C3	0.2235 (3)	0.84874 (14)	0.37065 (15)	0.0438 (4)
C2	0.1354 (3)	0.74050 (14)	0.36416 (14)	0.0430 (4)
H2	−0.0031	0.7264	0.3827	0.052*
C1	0.2519 (3)	0.65424 (13)	0.33049 (13)	0.0380 (4)
C21	0.1482 (4)	1.03154 (16)	0.39139 (19)	0.0595 (5)
H21A	0.2992	1.0754	0.4378	0.089*
H21B	0.0366	1.0762	0.4132	0.089*
H21C	0.1443	1.0132	0.3141	0.089*
C7	0.1389 (3)	0.53717 (14)	0.31542 (14)	0.0405 (4)
C8	0.2192 (3)	0.36345 (13)	0.33048 (13)	0.0376 (4)
C9	0.1204 (3)	0.31150 (13)	0.12858 (13)	0.0369 (4)
C14	−0.0742 (3)	0.29310 (16)	0.04667 (14)	0.0476 (4)
H14	−0.2192	0.2659	0.0529	0.057*
C13	−0.0526 (4)	0.31537 (19)	−0.04475 (16)	0.0610 (6)

H13	−0.1836	0.3029	−0.1002	0.073*
C12	0.1610 (4)	0.35575 (19)	−0.05447 (17)	0.0618 (6)
H12	0.1745	0.3719	−0.1155	0.074*
C11	0.3535 (4)	0.37199 (18)	0.02626 (18)	0.0579 (5)
H11	0.4983	0.3987	0.0195	0.070*
C10	0.3355 (3)	0.34923 (15)	0.11738 (15)	0.0464 (4)
H10	0.4675	0.3592	0.1712	0.056*
C15	−0.0664 (3)	0.18826 (13)	0.20578 (13)	0.0388 (4)
C16	−0.2480 (3)	0.20071 (17)	0.25501 (16)	0.0523 (5)
H16	−0.2582	0.2736	0.3035	0.063*
C17	−0.4148 (4)	0.1046 (2)	0.2320 (2)	0.0697 (6)
H17	−0.5374	0.1125	0.2654	0.084*
C18	−0.4001 (4)	−0.0022 (2)	0.16034 (19)	0.0757 (8)
H18	−0.5141	−0.0668	0.1442	0.091*
C19	−0.2184 (4)	−0.01469 (17)	0.11210 (17)	0.0691 (7)
H19	−0.2089	−0.0879	0.0640	0.083*
C20	−0.0481 (4)	0.08113 (14)	0.13453 (14)	0.0505 (5)
H20	0.0756	0.0729	0.1020	0.061*
H1A	0.416 (3)	0.5170 (16)	0.3974 (17)	0.049 (5)*

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0816 (4)	0.0375 (2)	0.0398 (3)	−0.0063 (2)	−0.0094 (2)	0.02171 (19)
O1	0.0816 (11)	0.0378 (7)	0.1021 (12)	0.0228 (7)	0.0492 (9)	0.0372 (7)
O2	0.0428 (8)	0.0374 (7)	0.0779 (9)	0.0029 (6)	0.0069 (7)	0.0217 (6)
N1	0.0461 (9)	0.0274 (7)	0.0387 (7)	−0.0018 (6)	−0.0051 (7)	0.0153 (6)
N2	0.0446 (8)	0.0302 (7)	0.0328 (7)	0.0003 (6)	0.0040 (6)	0.0142 (5)
C6	0.0488 (11)	0.0418 (9)	0.0496 (10)	0.0113 (8)	0.0086 (8)	0.0238 (8)
C5	0.0430 (10)	0.0521 (11)	0.0630 (12)	0.0058 (9)	0.0131 (9)	0.0333 (9)
C4	0.0488 (11)	0.0375 (9)	0.0517 (10)	0.0015 (8)	0.0055 (8)	0.0253 (8)
C3	0.0518 (11)	0.0335 (9)	0.0469 (9)	0.0085 (8)	0.0110 (8)	0.0188 (7)
C2	0.0461 (10)	0.0367 (9)	0.0495 (10)	0.0072 (8)	0.0132 (8)	0.0218 (8)
C1	0.0430 (10)	0.0328 (8)	0.0361 (8)	0.0026 (7)	0.0026 (7)	0.0176 (7)
C21	0.0784 (15)	0.0376 (10)	0.0712 (13)	0.0151 (10)	0.0196 (11)	0.0313 (10)
C7	0.0489 (11)	0.0310 (8)	0.0379 (8)	0.0046 (7)	0.0078 (7)	0.0137 (7)
C8	0.0401 (9)	0.0309 (8)	0.0372 (8)	0.0020 (7)	0.0029 (7)	0.0147 (7)
C9	0.0448 (10)	0.0315 (8)	0.0334 (8)	0.0053 (7)	0.0072 (7)	0.0152 (6)
C14	0.0459 (11)	0.0524 (10)	0.0402 (9)	0.0039 (8)	0.0045 (8)	0.0205 (8)
C13	0.0669 (14)	0.0732 (13)	0.0408 (10)	0.0146 (11)	0.0008 (9)	0.0283 (10)
C12	0.0808 (16)	0.0711 (14)	0.0466 (11)	0.0179 (12)	0.0230 (11)	0.0353 (10)
C11	0.0609 (13)	0.0614 (12)	0.0640 (12)	0.0120 (10)	0.0276 (10)	0.0350 (10)
C10	0.0463 (11)	0.0460 (10)	0.0472 (10)	0.0078 (8)	0.0090 (8)	0.0223 (8)
C15	0.0432 (10)	0.0341 (8)	0.0340 (8)	−0.0015 (7)	0.0017 (7)	0.0162 (7)
C16	0.0512 (12)	0.0517 (11)	0.0542 (11)	0.0064 (9)	0.0130 (9)	0.0247 (9)
C17	0.0547 (13)	0.0831 (16)	0.0705 (14)	−0.0078 (12)	0.0141 (11)	0.0403 (13)
C18	0.0862 (17)	0.0643 (14)	0.0560 (12)	−0.0330 (12)	0.0002 (12)	0.0286 (11)
C19	0.1047 (19)	0.0375 (10)	0.0451 (11)	−0.0142 (11)	0.0102 (12)	0.0110 (8)

C20	0.0684 (13)	0.0364 (9)	0.0386 (9)	0.0007 (9)	0.0121 (9)	0.0118 (7)

Geometric parameters (Å, °)

S1—C8	1.6503 (17)	C21—H21C	0.9600
O1—C3	1.351 (2)	C9—C14	1.378 (2)
O1—C21	1.421 (2)	C9—C10	1.381 (3)
O2—C7	1.210 (2)	C14—C13	1.381 (3)
N1—C8	1.384 (2)	C14—H14	0.9300
N1—C7	1.387 (2)	C13—C12	1.373 (3)
N1—H1A	0.87 (2)	C13—H13	0.9300
N2—C8	1.352 (2)	C12—C11	1.368 (3)
N2—C9	1.438 (2)	C12—H12	0.9300
N2—C15	1.439 (2)	C11—C10	1.375 (3)
C6—C5	1.382 (2)	C11—H11	0.9300
C6—C1	1.389 (3)	C10—H10	0.9300
C6—H6	0.9300	C15—C20	1.374 (3)
C5—C4	1.373 (3)	C15—C16	1.376 (3)
C5—H5	0.9300	C16—C17	1.377 (3)
C4—C3	1.378 (3)	C16—H16	0.9300
C4—H4	0.9300	C17—C18	1.365 (4)
C3—C2	1.387 (2)	C17—H17	0.9300
C2—C1	1.371 (3)	C18—C19	1.369 (4)
C2—H2	0.9300	C18—H18	0.9300
C1—C7	1.489 (2)	C19—C20	1.389 (3)
C21—H21A	0.9600	C19—H19	0.9300
C21—H21B	0.9600	C20—H20	0.9300
C3—O1—C21	118.49 (16)	C14—C9—C10	119.92 (16)
C8—N1—C7	122.68 (15)	C14—C9—N2	119.86 (15)
C8—N1—H1A	116.5 (13)	C10—C9—N2	120.21 (15)
C7—N1—H1A	117.1 (13)	C9—C14—C13	119.56 (18)
C8—N2—C9	122.38 (13)	C9—C14—H14	120.2
C8—N2—C15	119.92 (13)	C13—C14—H14	120.2
C9—N2—C15	117.62 (12)	C12—C13—C14	120.52 (19)
C5—C6—C1	118.54 (17)	C12—C13—H13	119.7
C5—C6—H6	120.7	C14—C13—H13	119.7
C1—C6—H6	120.7	C11—C12—C13	119.54 (18)
C4—C5—C6	121.22 (18)	C11—C12—H12	120.2
C4—C5—H5	119.4	C13—C12—H12	120.2
C6—C5—H5	119.4	C12—C11—C10	120.76 (19)
C5—C4—C3	119.88 (16)	C12—C11—H11	119.6
C5—C4—H4	120.1	C10—C11—H11	119.6
C3—C4—H4	120.1	C11—C10—C9	119.66 (18)
O1—C3—C4	124.89 (15)	C11—C10—H10	120.2
O1—C3—C2	115.53 (16)	C9—C10—H10	120.2
C4—C3—C2	119.57 (17)	C20—C15—C16	120.88 (16)
C1—C2—C3	120.20 (17)	C20—C15—N2	119.82 (16)
C1—C2—H2	119.9	C16—C15—N2	119.20 (16)
C3—C2—H2	119.9	C15—C16—C17	119.6 (2)

C2—C1—C6	120.58 (15)	C15—C16—H16	120.2
C2—C1—C7	117.72 (16)	C17—C16—H16	120.2
C6—C1—C7	121.53 (16)	C18—C17—C16	120.1 (2)
O1—C21—H21A	109.5	C18—C17—H17	120.0
O1—C21—H21B	109.5	C16—C17—H17	120.0
H21A—C21—H21B	109.5	C17—C18—C19	120.30 (19)
O1—C21—H21C	109.5	C17—C18—H18	119.9
H21A—C21—H21C	109.5	C19—C18—H18	119.9
H21B—C21—H21C	109.5	C18—C19—C20	120.5 (2)
O2—C7—N1	123.01 (15)	C18—C19—H19	119.8
O2—C7—C1	122.65 (16)	C20—C19—H19	119.8
N1—C7—C1	114.33 (15)	C15—C20—C19	118.6 (2)
N2—C8—N1	114.68 (14)	C15—C20—H20	120.7
N2—C8—S1	124.04 (12)	C19—C20—H20	120.7
N1—C8—S1	121.26 (12)		

Hydrogen-bond geometry (Å, °)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N1—H1A···S1^{i}	0.87 (2)	2.54 (2)	3.380 (2)	163.6 (17)

Symmetry codes: (i) −*x*+1, −*y*+1, −*z*+1.

Fig. 1

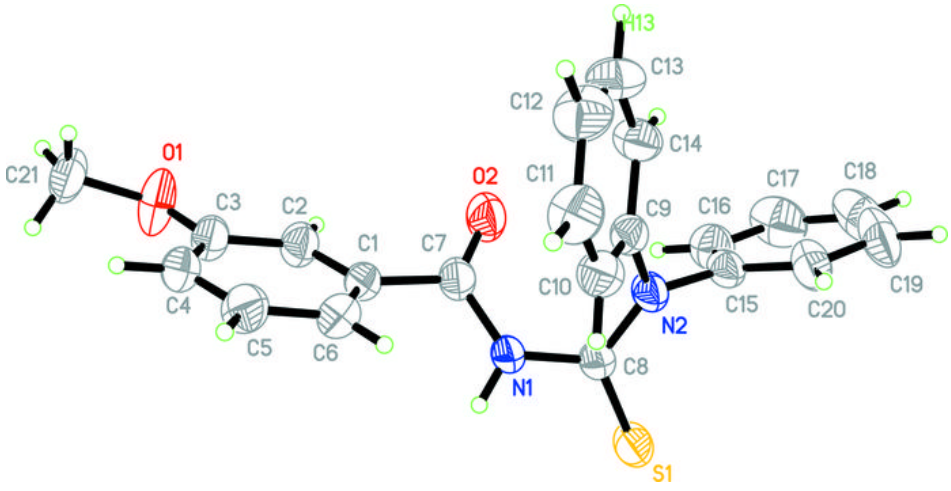
H13
C13
C12
C14
O1
C21
O2
C11
C17
C18
C3
C2
C9
C16
C1
C7
C10
C15
C19
C4
N2
C20
C5
C6
C8
N1
S1

Fig. 2