

S-Phenyl benzothioate

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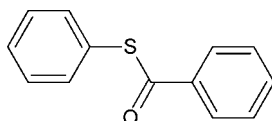
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 Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.032; wR factor = 0.082; data-to-parameter ratio = 12.9.

In the title compound, $\text{C}_{13}\text{H}_{10}\text{OS}$, the phenyl rings are inclined to one another by $51.12(8)^\circ$. There is a short $\text{C}-\text{H}\cdots\text{S}$ contact in the molecule. In the crystal, molecules are linked *via* $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds forming chains along the a axis. Molecules are also linked by $\text{C}-\text{H}\cdots\pi$ and weak $\pi-\pi$ interactions [centroid-centroid distance = $3.9543(10)$ Å].

Related literature

The title compound was obtained by the reaction of thio-phenolate and benzoyl chloride in an alkaline medium. For background to the title compound, see: Reddy *et al.* (2010); Katritzky *et al.* (2007). For details of the Cambridge Structural Database, see: Allen (2002).



Experimental

Crystal data

$\text{C}_{13}\text{H}_{10}\text{OS}$	$V = 1036.44(4)$ Å ³
$M_r = 214.27$	$Z = 4$
Monoclinic, $P2_1/c$	Cu $K\alpha$ radiation
$a = 5.7203(1)$ Å	$\mu = 2.49$ mm ⁻¹
$b = 15.1315(3)$ Å	$T = 100$ K
$c = 12.0606(3)$ Å	$0.25 \times 0.12 \times 0.12$ mm
$\beta = 96.867(1)^\circ$	

Data collection

Bruker APEX DUO 4K-CCD diffractometer	9164 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2008)	1759 independent reflections
$T_{\min} = 0.575$, $T_{\max} = 0.754$	1702 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.022$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$	136 parameters
$wR(F^2) = 0.082$	H-atom parameters constrained
$S = 1.04$	$\Delta\rho_{\text{max}} = 0.35$ e Å ⁻³
1759 reflections	$\Delta\rho_{\text{min}} = -0.26$ e Å ⁻³

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 and Cg2 are the centroids of the C2–C7 and C8–C13 rings, respectively.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C7}-\text{H7}\cdots\text{S1}$	0.95	2.52	2.9592 (16)	109
$\text{C13}-\text{H13}\cdots\text{O1}^{\text{i}}$	0.95	2.56	3.4889 (18)	167
$\text{C10}-\text{H10}\cdots\text{Cg1}^{\text{ii}}$	0.95	2.97	3.506 (2)	117
$\text{C5}-\text{H5}\cdots\text{Cg2}^{\text{iii}}$	0.95	2.73	3.5915 (19)	152

 Symmetry codes: (i) $x - 1, y, z$; (ii) $-x + 2, -y + 1, -z$; (iii) $-x + 2, y + \frac{1}{2}, -z + \frac{1}{2}$.

Data collection: APEX2 (Bruker, 2011); cell refinement: SAINT (Bruker, 2008); data reduction: SAINT and XPREP (Bruker, 2008); program(s) used to solve structure: SIR97 (Altomare *et al.*, 1999); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: DIAMOND (Brandenburg & Putz, 2005); software used to prepare material for publication: WinGX (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DS2217).

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supplementary materials

Acta Cryst. (2012). E68, o2825 [doi:10.1107/S1600536812037142]

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Comment

Reaction of thiophenolate and benzoyl chloride in alkaline medium was described previously by Reddy *et al.*, 2010. We have repeated the preparation of this compound to be used as starting material in some of our research. Benzoylation of thiophenol afforded colorless crystals of the title compound (see scheme and Figure 1) suitable for single crystal X-ray analysis of which the structure is reported herein. Molecules of the title compound crystallize in the $P2_1/c$ ($Z=4$) space group. All bond lengths are within their normal ranges (Allen, 2002). In the crystal packing several C—H \cdots O/ π interactions (see table 1, Fig. 2) as well as π - π stacking are observed (centroid to centroid distance = 3.9543 (10) Å, ring slippage = 1.366 Å).

Experimental

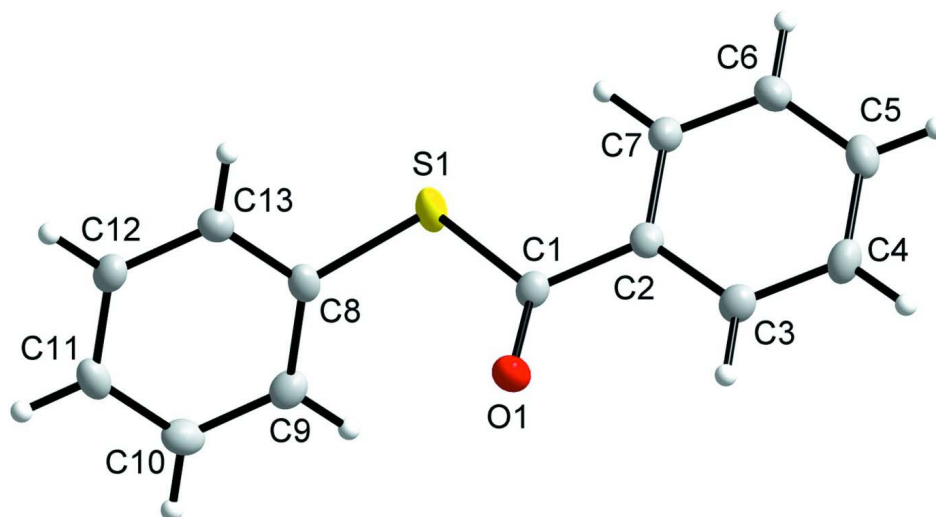
A mixture of sodium hydroxide (344 mg, 8.61 mmol) and thiophenol (0.9 ml, 8.61 mmol) were dissolved in methanol (22 ml) for about 10 minutes. Benzoyl chloride (1 ml) was added to it. The reaction mixture was stirred overnight and then poured into ice-cold water. Afterwards it was filtered and dried to afford the title compound as white crystals in 63% yield.

Refinement

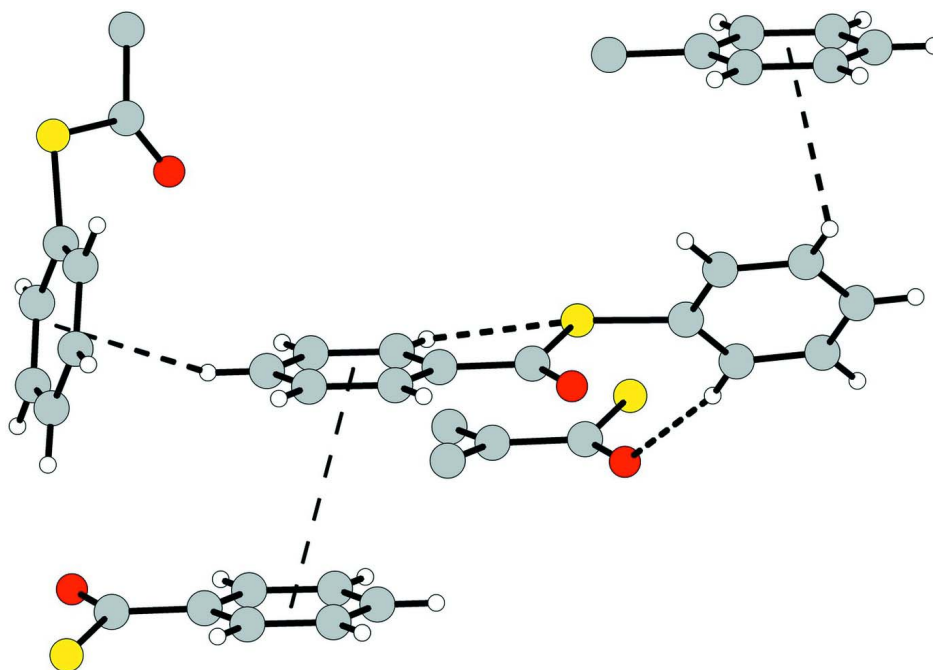
All hydrogen atoms were positioned in geometrically idealized positions with C—H = 0.95 Å and were allowed to ride on their parent atoms with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}$. A discrepant reflection (1 3 2) was removed in the final stages of refinement.

Computing details

Data collection: *APEX2* (Bruker, 2011); cell refinement: *SAINT* (Bruker, 2008); data reduction: *SAINT* and *XPREP* (Bruker, 2008); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg & Putz, 2005); software used to prepare material for publication: *WinGX* (Farrugia, 1999).


Figure 1

A view of (1). Displacement ellipsoids are drawn at the 50% probability level.


Figure 2

Packing diagram of (1) showing the C—H...O/S/π interactions as well as the π-π stacking.

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Crystal data

$C_{13}H_{10}OS$

$M_r = 214.27$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2_1/c$

$a = 5.7203$ (1) Å

$b = 15.1315$ (3) Å

$c = 12.0606$ (3) Å

$\beta = 96.867$ (1)°

$V = 1036.44$ (4) Å³

$Z = 4$

$F(000) = 448$

$D_x = 1.373$ Mg m⁻³

Cu $K\alpha$ radiation, $\lambda = 1.54178 \text{ \AA}$
 Cell parameters from 6683 reflections
 $\theta = 4.7\text{--}65.8^\circ$
 $\mu = 2.49 \text{ mm}^{-1}$

$T = 100 \text{ K}$
 Rectangular, colourless
 $0.25 \times 0.12 \times 0.12 \text{ mm}$

Data collection

Bruker APEX DUO 4K-CCD
 diffractometer
 Radiation source: Incoatec I μ S microfocus X-
 ray source
 Incoatec Quazar Multilayer Mirror
 monochromator
 Detector resolution: $8.4 \text{ pixels mm}^{-1}$
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Bruker, 2008)

$T_{\min} = 0.575$, $T_{\max} = 0.754$
 9164 measured reflections
 1759 independent reflections
 1702 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.022$
 $\theta_{\max} = 66.4^\circ$, $\theta_{\min} = 4.7^\circ$
 $h = -2 \rightarrow 6$
 $k = -17 \rightarrow 17$
 $l = -14 \rightarrow 13$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.082$
 $S = 1.04$
 1759 reflections
 136 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0364P)^2 + 0.7558P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.35 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.26 \text{ e \AA}^{-3}$

Special details

Experimental. The intensity data was collected on a Bruker Apex DUO 4 K CCD diffractometer using an exposure time of 5 s/frame. A total of 2274 frames were collected with a frame width of 1° covering up to $\theta = 66.38^\circ$ with 96.3% completeness accomplished.

Analytical data: mp: $53\text{--}55^\circ\text{C}$ (Lit. $54\text{--}55^\circ\text{C}$; Katritzky *et al.*, 2007); ^1H NMR (CDCl_3 , 400 MHz): d 8.03 (d, $J = 0.8 \text{ Hz}$, 1H), 8.01 (d, $J = 1.2 \text{ Hz}$, 1H), 7.62–7.58 (m, 1H), 7.52–7.44 (m, 7H), ^{13}C NMR (CDCl_3 , 400 MHz): d 190.1, 136.6, 135.1, 133.6, 129.5, 129.2, 128.7, 127.5, 127.3.

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.66521 (7)	0.44609 (2)	0.09450 (3)	0.02499 (16)
O1	1.05856 (19)	0.36323 (7)	0.17500 (9)	0.0242 (3)
C1	0.9304 (3)	0.42516 (10)	0.18525 (12)	0.0180 (3)
C2	0.9751 (3)	0.49420 (10)	0.27395 (12)	0.0174 (3)
C3	1.1950 (3)	0.49605 (10)	0.33797 (13)	0.0210 (3)
H3	1.3106	0.4531	0.3262	0.025*

C4	1.2443 (3)	0.56059 (11)	0.41871 (14)	0.0237 (4)
H4	1.3943	0.5619	0.462	0.028*
C5	1.0767 (3)	0.62332 (11)	0.43703 (13)	0.0234 (3)
H5	1.1117	0.6673	0.4928	0.028*
C6	0.8580 (3)	0.62174 (11)	0.37383 (13)	0.0239 (4)
H6	0.7429	0.6647	0.3863	0.029*
C7	0.8067 (3)	0.55781 (11)	0.29258 (13)	0.0214 (3)
H7	0.6566	0.5571	0.2493	0.026*
C8	0.6486 (3)	0.35922 (10)	-0.00504 (12)	0.0182 (3)
C9	0.8130 (3)	0.35150 (10)	-0.08087 (13)	0.0211 (3)
H9	0.9497	0.3877	-0.0737	0.025*
C10	0.7749 (3)	0.29047 (11)	-0.16688 (13)	0.0231 (4)
H10	0.8871	0.2843	-0.2184	0.028*
C11	0.5735 (3)	0.23835 (10)	-0.17811 (13)	0.0221 (3)
H11	0.5472	0.1972	-0.2378	0.027*
C12	0.4107 (3)	0.24634 (10)	-0.10217 (13)	0.0210 (3)
H12	0.2734	0.2105	-0.1097	0.025*
C13	0.4483 (3)	0.30674 (10)	-0.01499 (12)	0.0192 (3)
H13	0.3376	0.312	0.0374	0.023*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0287 (3)	0.0191 (2)	0.0243 (2)	0.00622 (15)	-0.00893 (16)	-0.00571 (15)
O1	0.0236 (6)	0.0242 (6)	0.0242 (6)	0.0056 (5)	0.0007 (4)	-0.0028 (5)
C1	0.0190 (7)	0.0164 (7)	0.0181 (8)	-0.0014 (6)	0.0004 (6)	0.0032 (6)
C2	0.0203 (7)	0.0174 (8)	0.0146 (7)	-0.0018 (6)	0.0027 (6)	0.0027 (6)
C3	0.0211 (8)	0.0189 (8)	0.0224 (8)	0.0007 (6)	0.0002 (6)	0.0032 (6)
C4	0.0235 (8)	0.0246 (8)	0.0215 (8)	-0.0034 (6)	-0.0043 (6)	0.0024 (6)
C5	0.0298 (8)	0.0226 (8)	0.0174 (7)	-0.0044 (6)	0.0015 (6)	-0.0019 (6)
C6	0.0245 (8)	0.0247 (8)	0.0233 (8)	0.0008 (6)	0.0055 (6)	-0.0040 (7)
C7	0.0188 (8)	0.0259 (9)	0.0191 (8)	0.0003 (6)	0.0010 (6)	-0.0019 (6)
C8	0.0223 (8)	0.0148 (7)	0.0161 (7)	0.0031 (6)	-0.0030 (6)	0.0010 (6)
C9	0.0203 (8)	0.0205 (8)	0.0220 (8)	-0.0013 (6)	0.0006 (6)	0.0060 (6)
C10	0.0248 (8)	0.0272 (9)	0.0181 (8)	0.0047 (6)	0.0054 (6)	0.0048 (6)
C11	0.0285 (8)	0.0196 (8)	0.0171 (8)	0.0045 (6)	-0.0021 (6)	-0.0021 (6)
C12	0.0195 (7)	0.0187 (8)	0.0242 (8)	-0.0010 (6)	-0.0005 (6)	-0.0006 (6)
C13	0.0202 (8)	0.0191 (8)	0.0184 (7)	0.0030 (6)	0.0024 (6)	0.0010 (6)

Geometric parameters (\AA , $^\circ$)

S1—C8	1.7751 (15)	C6—H6	0.95
S1—C1	1.7894 (15)	C7—H7	0.95
O1—C1	1.2054 (19)	C8—C13	1.387 (2)
C1—C2	1.495 (2)	C8—C9	1.393 (2)
C2—C3	1.395 (2)	C9—C10	1.386 (2)
C2—C7	1.399 (2)	C9—H9	0.95
C3—C4	1.384 (2)	C10—C11	1.390 (2)
C3—H3	0.95	C10—H10	0.95
C4—C5	1.385 (2)	C11—C12	1.387 (2)

C4—H4	0.95	C11—H11	0.95
C5—C6	1.385 (2)	C12—C13	1.390 (2)
C5—H5	0.95	C12—H12	0.95
C6—C7	1.383 (2)	C13—H13	0.95
C8—S1—C1	104.81 (7)	C6—C7—H7	119.9
O1—C1—C2	124.23 (13)	C2—C7—H7	119.9
O1—C1—S1	123.81 (12)	C13—C8—C9	120.70 (14)
C2—C1—S1	111.96 (10)	C13—C8—S1	117.35 (12)
C3—C2—C7	119.34 (14)	C9—C8—S1	121.36 (12)
C3—C2—C1	118.40 (13)	C10—C9—C8	119.29 (14)
C7—C2—C1	122.24 (13)	C10—C9—H9	120.4
C4—C3—C2	119.85 (15)	C8—C9—H9	120.4
C4—C3—H3	120.1	C9—C10—C11	120.34 (14)
C2—C3—H3	120.1	C9—C10—H10	119.8
C3—C4—C5	120.60 (15)	C11—C10—H10	119.8
C3—C4—H4	119.7	C12—C11—C10	120.02 (15)
C5—C4—H4	119.7	C12—C11—H11	120
C6—C5—C4	119.80 (15)	C10—C11—H11	120
C6—C5—H5	120.1	C11—C12—C13	120.09 (14)
C4—C5—H5	120.1	C11—C12—H12	120
C7—C6—C5	120.21 (15)	C13—C12—H12	120
C7—C6—H6	119.9	C8—C13—C12	119.54 (14)
C5—C6—H6	119.9	C8—C13—H13	120.2
C6—C7—C2	120.20 (14)	C12—C13—H13	120.2
C8—S1—C1—O1	-0.46 (15)	C3—C2—C7—C6	-0.1 (2)
C8—S1—C1—C2	178.75 (10)	C1—C2—C7—C6	-178.42 (14)
O1—C1—C2—C3	10.8 (2)	C1—S1—C8—C13	122.78 (12)
S1—C1—C2—C3	-168.42 (11)	C1—S1—C8—C9	-66.00 (14)
O1—C1—C2—C7	-170.86 (15)	C13—C8—C9—C10	-0.1 (2)
S1—C1—C2—C7	9.93 (18)	S1—C8—C9—C10	-171.00 (11)
C7—C2—C3—C4	-0.1 (2)	C8—C9—C10—C11	0.8 (2)
C1—C2—C3—C4	178.26 (14)	C9—C10—C11—C12	-0.9 (2)
C2—C3—C4—C5	0.3 (2)	C10—C11—C12—C13	0.3 (2)
C3—C4—C5—C6	-0.2 (2)	C9—C8—C13—C12	-0.5 (2)
C4—C5—C6—C7	0.0 (2)	S1—C8—C13—C12	170.75 (11)
C5—C6—C7—C2	0.2 (2)	C11—C12—C13—C8	0.4 (2)

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

Cg1 and Cg2 are the centroids of the C2—C7 and C8—C13 rings, respectively.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C7—H7 \cdots S1	0.95	2.52	2.9592 (16)	109
C13—H13 \cdots O1 ⁱ	0.95	2.56	3.4889 (18)	167
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