

1-(4-Methylphenyl)-2-(phenylsulfonyl)-ethanone

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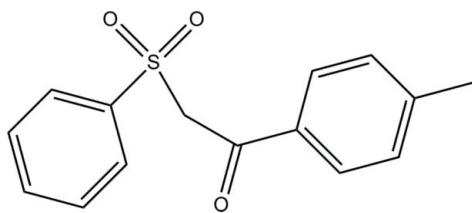
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; R factor = 0.045; wR factor = 0.124; data-to-parameter ratio = 14.3.

In the title compound, $\text{C}_{15}\text{H}_{14}\text{O}_3\text{S}$, the benzene and phenyl rings make a dihedral angle of 33.56 (16)°. In the crystal, molecules are linked by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds into a layer parallel to the ab plane.

Related literature

For background to the chemistry of sulfones, see: Xiang *et al.* (2007); Abdel-Aziz & Mekawey (2009); Abdel-Aziz *et al.* (2010). For a related structure, see: Abdel-Aziz *et al.* (2011). For reference bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{15}\text{H}_{14}\text{O}_3\text{S}$
 $M_r = 274.32$
Orthorhombic, $Pbca$
 $a = 11.5555$ (12) Å
 $b = 10.1981$ (11) Å
 $c = 22.843$ (2) Å

$V = 2692.0$ (5) Å³
 $Z = 8$
Cu $K\alpha$ radiation
 $\mu = 2.15$ mm⁻¹
 $T = 296$ K
 $0.55 \times 0.12 \times 0.04$ mm

Data collection

Bruker SMART APEXII CCD
area-detector diffractometer
Absorption correction: multi-scan
(*SADABS*; Bruker, 2009)
 $T_{\min} = 0.386$, $T_{\max} = 0.919$

10168 measured reflections
2487 independent reflections
1446 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.068$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$
 $wR(F^2) = 0.124$
 $S = 0.89$
2487 reflections

174 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.19$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.25$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C7}-\text{H7B}\cdots\text{O1}^i$	0.97	2.42	3.366 (3)	164
$\text{C10}-\text{H10A}\cdots\text{O3}^{ii}$	0.93	2.48	3.342 (4)	154

Symmetry codes: (i) $-x + \frac{3}{2}, y + \frac{1}{2}, z$; (ii) $x - \frac{1}{2}, y, -z + \frac{1}{2}$.

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; 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: IS5087).

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supplementary materials

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Comment

In the title compound (Fig. 1), the two aromatic rings (C1–C6 and C9–C14) form a dihedral angle of 33.56 (16)° with each other. Bond lengths (Allen *et al.*, 1987) and angles are within normal ranges and are comparable to the related structure (Abdel-Aziz *et al.*, 2011). In the crystal packing (Fig. 2), the molecules are linked by intermolecular C7—H7B···O1 and C10—H10A···O3 hydrogen bonds (Table 1) into two-dimensional networks parallel to the *ab* plane.

Experimental

The title compound was prepared according to the reported method (Xiang *et al.*, 2007). Single crystals of the title compound were obtained by slow evaporation from an ethanol solution at room temperature.

Refinement

All H atoms were positioned geometrically (C—H = 0.93, 0.96 and 0.97 Å) and refined using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2$ and $1.5U_{\text{eq}}(\text{C})$.

Computing details

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT* (Bruker, 2009); program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008) and *PLATON* (Spek, 2009).

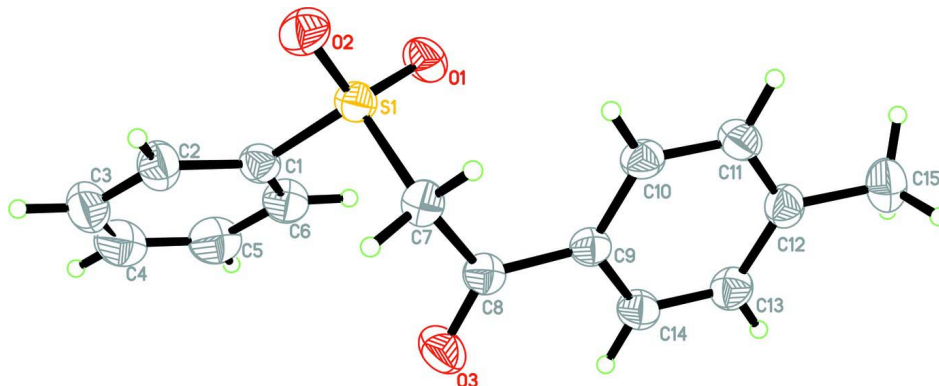


Figure 1

The molecular structure of the title compound with atom labels and 30% probability displacement ellipsoids.

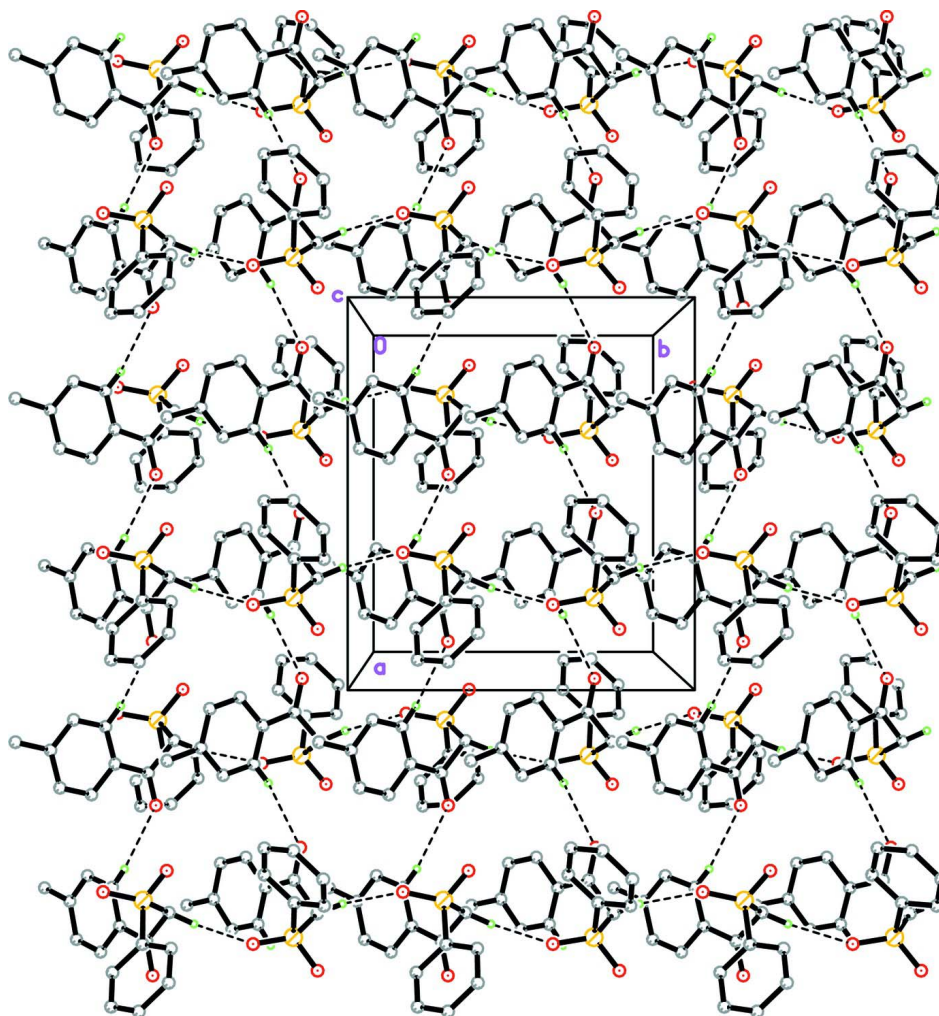


Figure 2

The crystal packing of the title compound. The dashed lines represent the hydrogen bonds. For clarity sake, hydrogen atoms not involved in hydrogen bonding have been omitted.

1-(4-Methylphenyl)-2-(phenylsulfonyl)ethanone

Crystal data

$C_{15}H_{14}O_3S$

$M_r = 274.32$

Orthorhombic, *Pbca*

Hall symbol: $-P\ 2ac\ 2ab$

$a = 11.5555\ (12)\ \text{\AA}$

$b = 10.1981\ (11)\ \text{\AA}$

$c = 22.843\ (2)\ \text{\AA}$

$V = 2692.0\ (5)\ \text{\AA}^3$

$Z = 8$

$F(000) = 1152$

$D_x = 1.354\ \text{Mg m}^{-3}$

Cu $K\alpha$ radiation, $\lambda = 1.54178\ \text{\AA}$

Cell parameters from 484 reflections

$\theta = 3.9\text{--}33.7^\circ$

$\mu = 2.15\ \text{mm}^{-1}$

$T = 296\ \text{K}$

Needle, colourless

$0.55 \times 0.12 \times 0.04\ \text{mm}$

Data collection

Bruker SMART APEXII CCD area-detector diffractometer	10168 measured reflections 2487 independent reflections
Radiation source: fine-focus sealed tube	1446 reflections with $I > 2\sigma(I)$
Graphite monochromator	$R_{\text{int}} = 0.068$
φ and ω scans	$\theta_{\text{max}} = 70.0^\circ$, $\theta_{\text{min}} = 3.9^\circ$
Absorption correction: multi-scan (SADABS; Bruker, 2009)	$h = -11 \rightarrow 14$
$T_{\text{min}} = 0.386$, $T_{\text{max}} = 0.919$	$k = -9 \rightarrow 11$ $l = -27 \rightarrow 17$

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.045$	$w = 1/[\sigma^2(F_o^2) + (0.0623P)^2]$
$wR(F^2) = 0.124$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.89$	$(\Delta/\sigma)_{\text{max}} < 0.001$
2487 reflections	$\Delta\rho_{\text{max}} = 0.19 \text{ e } \text{\AA}^{-3}$
174 parameters	$\Delta\rho_{\text{min}} = -0.25 \text{ e } \text{\AA}^{-3}$
0 restraints	Extinction correction: SHELXTL (Sheldrick, 2008), $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.00131 (15)
Secondary atom site location: difference Fourier map	

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.69612 (6)	0.25970 (7)	0.35841 (3)	0.0496 (2)
O1	0.67330 (18)	0.12398 (18)	0.34807 (10)	0.0602 (6)
O2	0.60372 (18)	0.3428 (2)	0.37858 (10)	0.0666 (6)
O3	0.94161 (18)	0.2723 (2)	0.27397 (11)	0.0727 (7)
C1	0.8099 (2)	0.2698 (3)	0.40947 (13)	0.0498 (7)
C2	0.8117 (3)	0.3717 (3)	0.44912 (14)	0.0652 (9)
H2A	0.7562	0.4377	0.4476	0.078*
C3	0.8977 (3)	0.3739 (4)	0.49129 (15)	0.0791 (11)
H3A	0.8998	0.4410	0.5188	0.095*
C4	0.9795 (4)	0.2774 (4)	0.49242 (17)	0.0835 (12)
H4A	1.0372	0.2799	0.5208	0.100*
C5	0.9782 (3)	0.1777 (4)	0.45296 (17)	0.0797 (11)
H5A	1.0354	0.1135	0.4542	0.096*
C6	0.8919 (3)	0.1715 (3)	0.41090 (14)	0.0636 (8)
H6A	0.8894	0.1026	0.3842	0.076*

C7	0.7484 (3)	0.3314 (3)	0.29256 (12)	0.0513 (7)
H7A	0.6841	0.3433	0.2658	0.062*
H7B	0.7799	0.4174	0.3013	0.062*
C8	0.8414 (2)	0.2493 (3)	0.26254 (12)	0.0503 (7)
C9	0.8077 (2)	0.1467 (3)	0.22005 (12)	0.0459 (6)
C10	0.6939 (2)	0.1183 (3)	0.20632 (13)	0.0520 (7)
H10A	0.6344	0.1646	0.2243	0.062*
C11	0.6681 (3)	0.0219 (3)	0.16615 (13)	0.0575 (8)
H11A	0.5910	0.0031	0.1580	0.069*
C12	0.7533 (3)	-0.0470 (3)	0.13781 (13)	0.0559 (7)
C13	0.8667 (3)	-0.0200 (3)	0.15236 (14)	0.0576 (8)
H13A	0.9258	-0.0669	0.1344	0.069*
C14	0.8942 (2)	0.0743 (3)	0.19262 (13)	0.0549 (7)
H14A	0.9714	0.0902	0.2017	0.066*
C15	0.7230 (3)	-0.1491 (3)	0.09252 (15)	0.0753 (10)
H15B	0.6483	-0.1854	0.1012	0.113*
H15C	0.7799	-0.2176	0.0931	0.113*
H15A	0.7214	-0.1093	0.0545	0.113*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0458 (4)	0.0490 (4)	0.0540 (4)	-0.0012 (3)	0.0010 (3)	-0.0013 (3)
O1	0.0659 (14)	0.0448 (12)	0.0700 (14)	-0.0143 (9)	-0.0041 (11)	-0.0014 (10)
O2	0.0541 (13)	0.0746 (16)	0.0710 (15)	0.0124 (10)	0.0071 (11)	-0.0079 (11)
O3	0.0480 (13)	0.0813 (17)	0.0887 (17)	-0.0041 (11)	-0.0105 (12)	-0.0201 (13)
C1	0.0506 (17)	0.0480 (17)	0.0509 (15)	-0.0049 (13)	0.0002 (13)	0.0035 (12)
C2	0.078 (2)	0.057 (2)	0.0599 (19)	-0.0065 (16)	-0.0027 (18)	-0.0040 (15)
C3	0.093 (3)	0.084 (3)	0.061 (2)	-0.024 (2)	-0.007 (2)	-0.0066 (19)
C4	0.074 (3)	0.114 (4)	0.062 (2)	-0.018 (2)	-0.0189 (19)	0.023 (2)
C5	0.067 (2)	0.094 (3)	0.078 (3)	0.006 (2)	-0.007 (2)	0.018 (2)
C6	0.0573 (19)	0.065 (2)	0.068 (2)	0.0057 (15)	-0.0047 (17)	0.0045 (16)
C7	0.0611 (18)	0.0432 (17)	0.0495 (16)	0.0030 (13)	-0.0005 (14)	0.0020 (12)
C8	0.0482 (17)	0.0517 (17)	0.0508 (16)	0.0008 (13)	-0.0023 (13)	0.0039 (13)
C9	0.0421 (15)	0.0466 (16)	0.0491 (15)	0.0022 (11)	0.0002 (13)	0.0034 (12)
C10	0.0408 (15)	0.0613 (18)	0.0540 (17)	0.0053 (13)	0.0030 (14)	0.0020 (13)
C11	0.0465 (18)	0.066 (2)	0.0599 (18)	-0.0071 (14)	-0.0064 (15)	0.0013 (15)
C12	0.0628 (19)	0.0579 (18)	0.0471 (15)	-0.0022 (14)	-0.0031 (16)	0.0031 (14)
C13	0.0544 (19)	0.0594 (19)	0.0590 (18)	0.0086 (13)	0.0040 (16)	-0.0045 (15)
C14	0.0412 (16)	0.0636 (19)	0.0599 (18)	0.0035 (13)	-0.0013 (14)	-0.0034 (14)
C15	0.093 (3)	0.071 (2)	0.061 (2)	-0.0040 (18)	-0.0110 (19)	-0.0085 (17)

Geometric parameters (Å, °)

S1—O1	1.4287 (19)	C7—H7A	0.9700
S1—O2	1.439 (2)	C7—H7B	0.9700
S1—C1	1.760 (3)	C8—C9	1.480 (4)
S1—C7	1.778 (3)	C9—C10	1.382 (4)
O3—C8	1.210 (3)	C9—C14	1.391 (4)
C1—C2	1.379 (4)	C10—C11	1.377 (4)

C1—C6	1.380 (4)	C10—H10A	0.9300
C2—C3	1.385 (4)	C11—C12	1.373 (4)
C2—H2A	0.9300	C11—H11A	0.9300
C3—C4	1.364 (5)	C12—C13	1.379 (4)
C3—H3A	0.9300	C12—C15	1.509 (4)
C4—C5	1.359 (5)	C13—C14	1.368 (4)
C4—H4A	0.9300	C13—H13A	0.9300
C5—C6	1.386 (5)	C14—H14A	0.9300
C5—H5A	0.9300	C15—H15B	0.9600
C6—H6A	0.9300	C15—H15C	0.9600
C7—C8	1.525 (4)	C15—H15A	0.9600
O1—S1—O2	119.10 (13)	H7A—C7—H7B	107.7
O1—S1—C1	107.69 (13)	O3—C8—C9	122.0 (3)
O2—S1—C1	107.92 (14)	O3—C8—C7	118.1 (3)
O1—S1—C7	108.74 (14)	C9—C8—C7	119.9 (2)
O2—S1—C7	106.32 (14)	C10—C9—C14	118.0 (3)
C1—S1—C7	106.42 (14)	C10—C9—C8	123.2 (3)
C2—C1—C6	121.4 (3)	C14—C9—C8	118.8 (3)
C2—C1—S1	119.4 (2)	C11—C10—C9	120.4 (3)
C6—C1—S1	119.1 (2)	C11—C10—H10A	119.8
C1—C2—C3	118.7 (3)	C9—C10—H10A	119.8
C1—C2—H2A	120.7	C12—C11—C10	121.6 (3)
C3—C2—H2A	120.7	C12—C11—H11A	119.2
C4—C3—C2	119.9 (4)	C10—C11—H11A	119.2
C4—C3—H3A	120.0	C11—C12—C13	117.7 (3)
C2—C3—H3A	120.0	C11—C12—C15	120.7 (3)
C5—C4—C3	121.3 (4)	C13—C12—C15	121.6 (3)
C5—C4—H4A	119.3	C14—C13—C12	121.5 (3)
C3—C4—H4A	119.3	C14—C13—H13A	119.2
C4—C5—C6	120.1 (4)	C12—C13—H13A	119.2
C4—C5—H5A	120.0	C13—C14—C9	120.6 (3)
C6—C5—H5A	120.0	C13—C14—H14A	119.7
C1—C6—C5	118.5 (3)	C9—C14—H14A	119.7
C1—C6—H6A	120.7	C12—C15—H15B	109.5
C5—C6—H6A	120.7	C12—C15—H15C	109.5
C8—C7—S1	113.21 (19)	H15B—C15—H15C	109.5
C8—C7—H7A	108.9	C12—C15—H15A	109.5
S1—C7—H7A	108.9	H15B—C15—H15A	109.5
C8—C7—H7B	108.9	H15C—C15—H15A	109.5
S1—C7—H7B	108.9		
O1—S1—C1—C2	-146.6 (2)	S1—C7—C8—O3	-92.6 (3)
O2—S1—C1—C2	-16.8 (3)	S1—C7—C8—C9	88.4 (3)
C7—S1—C1—C2	97.0 (3)	O3—C8—C9—C10	179.8 (3)
O1—S1—C1—C6	30.0 (3)	C7—C8—C9—C10	-1.2 (4)
O2—S1—C1—C6	159.8 (2)	O3—C8—C9—C14	0.2 (4)
C7—S1—C1—C6	-86.4 (3)	C7—C8—C9—C14	179.1 (3)
C6—C1—C2—C3	-0.4 (5)	C14—C9—C10—C11	-0.6 (4)

S1—C1—C2—C3	176.1 (2)	C8—C9—C10—C11	179.7 (3)
C1—C2—C3—C4	1.0 (5)	C9—C10—C11—C12	-1.2 (5)
C2—C3—C4—C5	-0.4 (6)	C10—C11—C12—C13	2.3 (5)
C3—C4—C5—C6	-0.8 (6)	C10—C11—C12—C15	-178.0 (3)
C2—C1—C6—C5	-0.8 (5)	C11—C12—C13—C14	-1.5 (5)
S1—C1—C6—C5	-177.3 (3)	C15—C12—C13—C14	178.7 (3)
C4—C5—C6—C1	1.4 (5)	C12—C13—C14—C9	-0.3 (5)
O1—S1—C7—C8	-46.2 (3)	C10—C9—C14—C13	1.4 (4)
O2—S1—C7—C8	-175.5 (2)	C8—C9—C14—C13	-178.9 (3)
C1—S1—C7—C8	69.6 (2)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C7—H7B...O1 ⁱ	0.97	2.42	3.366 (3)	164
C10—H10A...O3 ⁱⁱ	0.93	2.48	3.342 (4)	154

Symmetry codes: (i) $-x+3/2, y+1/2, z$; (ii) $x-1/2, y, -z+1/2$.