

N-(2-Methylphenylsulfonyl)acetamide

 K. Shakuntala,^a Sabine Foro^b and B. Thimme Gowda^{a*}

^aDepartment of Chemistry, Mangalore University, Mangalagangothri 574 199, Mangalore, India, and ^bInstitute of Materials Science, Darmstadt University of Technology, Petersenstrasse 23, D-64287 Darmstadt, Germany
Correspondence e-mail: gowdabt@yahoo.com

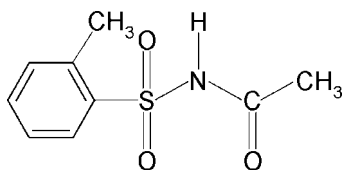
Received 12 April 2011; accepted 15 April 2011

Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; R factor = 0.035; wR factor = 0.086; data-to-parameter ratio = 14.7.

In the molecular structure of the title compound, $\text{C}_9\text{H}_{11}\text{NO}_3\text{S}$, the N—H and C=O bonds are *anti* to each other, while the amide H atom is *syn* with respect to the *ortho*-methyl group in the benzene ring. The C—S—N—C torsion angle is -58.2 (2)°, indicating a twist in the molecule. In the crystal, N—H...O hydrogen bonds link the molecules into chains along the c axis.

Related literature

For the sulfanilamide moiety in sulfonamide drugs, see: Maren (1976). For hydrogen bonding modes of sulfonamides, see: Adson & Grant (2001). For our study of the effect of substituents on the structures of *N*-(aryl)-amides, see: Gowda *et al.* (2004). For background to the structures of *N*-(substituted phenylsulfonyl)-substituted-amides, see: Gowda *et al.* (2010); Shakuntala *et al.* (2011) and for the oxidative strengths of *N*-chloro, *N*-arylsulfonamides, see: Gowda & Kumar (2003).



Experimental

Crystal data

$\text{C}_9\text{H}_{11}\text{NO}_3\text{S}$
 $M_r = 213.25$
Tetragonal, $P4_3$
 $a = 7.9804$ (5) Å

$c = 16.749$ (1) Å
 $V = 1066.69$ (11) Å³
 $Z = 4$
Mo $K\alpha$ radiation

$\mu = 0.29$ mm⁻¹
 $T = 293$ K

0.40 × 0.18 × 0.12 mm

Data collection

Oxford Diffraction Xcalibur diffractometer with a Sapphire CCD detector
Absorption correction: multi-scan (*CrysAlis RED*; Oxford)

Diffraction, 2009)
 $T_{\min} = 0.895$, $T_{\max} = 0.967$
4245 measured reflections
1944 independent reflections
1690 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.016$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.086$
 $S = 1.08$
1944 reflections
132 parameters
2 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.14$ e Å⁻³
 $\Delta\rho_{\min} = -0.17$ e Å⁻³
Absolute structure: Flack (1983), 824 Friedel pairs
Flack parameter: 0.03 (9)

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1N}\cdots\text{O3}^i$	0.86 (2)	1.95 (2)	2.770 (3)	162 (3)

 Symmetry code: (i) $-y + 1, x, z - \frac{1}{4}$

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2009); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2009); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL97*.

KS thanks the University Grants Commission, Government of India, New Delhi, for the award of a research fellowship under its faculty improvement program.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2737).

References

- Adson, D. A. & Grant, D. J. W. (2001). *J. Pharm. Sci.* **90**, 2058–2077.
Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
Gowda, B. T., Foro, S., Nirmala, P. G. & Fuess, H. (2010). *Acta Cryst.* **E66**, o1284.
Gowda, B. T. & Kumar, B. H. A. (2003). *Oxid. Commun.* **26**, 403–425.
Gowda, B. T., Svoboda, I. & Fuess, H. (2004). *Z. Naturforsch. Teil A*, **55**, 845–852.
Maren, T. H. (1976). *Annu. Rev. Pharmacol Toxicol.* **16**, 309–327.
Oxford Diffraction (2009). *CrysAlis CCD* and *CrysAlis RED*. Oxford Diffraction Ltd, Yarnton, England.
Shakuntala, K., Foro, S. & Gowda, B. T. (2011). *Acta Cryst.* **E67**, o1097.
Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.

supplementary materials

Acta Cryst. (2011). E67, o1188 [doi:10.1107/S1600536811014218]

***N*-(2-Methylphenylsulfonyl)acetamide**

K. Shakuntala, S. Foro and B. T. Gowda

Comment

The structures of sulfonamide drugs contain the sulfanilamide moiety (Maren, 1976). The hydrogen bonding preferences of sulfonamides has been investigated (Adsmund & Grant, 2001). The nature and position of the substituents play a significant role on their crystal structures and other aspects of *N*-(aryl)-amides and *N*-(aryl)-sulfonamides (Gowda *et al.*, 2003, 2004; Shakuntala *et al.*, 2011). As a part of a study of the effects of substituents on the structures of this class of compounds, the structure of *N*-(2-methylphenylsulfonyl)-acetamide (I) has been determined (Fig. 1). The conformation of the N—C bond in the C—SO₂—NH—C(O) segment has *gauche* torsions with respect to the S=O bonds, the torsion angles being C7—N1—S1—O2 = 57.5 (3) ° and C7—N1—S1—O1 = -174.5 (2) °.

The N—H and C=O bonds are *anti* to each other, similar to that observed in *N*-(phenylsulfonyl)-acetamide (II) (Gowda *et al.*, 2010) and *N*-(2-chlorophenylsulfonyl)-acetamide (III) (Shakuntala *et al.*, 2011). Further, the conformation of the amide-H atom is *syn* to the *ortho*-methyl group in the benzene ring, similar to that observed between the amide-H atom and the *ortho*-chloro group in (III). The molecule of (I) is bent at the *S*-atom with a C—S—N—C torsion angle of -58.2 (2) °, compared to the values of -58.8 (4) ° in (II), and -71.7 (3) and 61.2 (3) ° in the two independent molecules of (III).

In the crystal structure, intermolecular N—H···O hydrogen bonds (Table 1) link the molecules into chains along the *c* axis; part of the crystal structure is shown in Fig. 2.

Experimental

The title compound was prepared by refluxing 2-methylbenzenesulfonamide (0.10 mole) with an excess of acetyl chloride (0.20 mole) for one hour on a water bath. The reaction mixture was cooled and poured into ice cold water. The resulting solid was separated, washed thoroughly with water and dissolved in warm dilute sodium hydrogen carbonate solution. The title compound was re-precipitated by acidifying the filtered solution with glacial acetic acid. It was filtered, dried and recrystallized from ethanol. Colourless rods of the title compound were obtained from a slow evaporation of its ethanol solution.

Refinement

The H atom of the NH group was located in a difference map and later restrained to the distance N—H = 0.86±0.02 Å. The other H atoms were positioned with idealized geometry using a riding model with the aromatic C—H distance = 0.93 Å and methyl C—H = 0.96 Å. All H atoms were refined with isotropic displacement parameters set to 1.2 times of the U_{eq} of the parent atom.

Figures

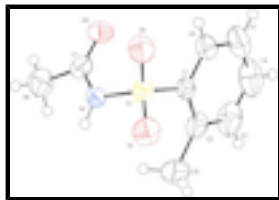


Fig. 1. Molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.

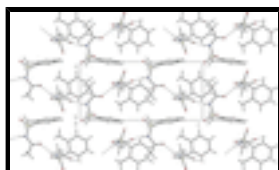


Fig. 2. View of the crystal packing in (I). Hydrogen bonds are shown as dashed lines.

N-(2-Methylphenylsulfonyl)acetamide

Crystal data

$C_9H_{11}NO_3S$

$M_r = 213.25$

Tetragonal, $P4_3$

Hall symbol: P 4cw

$a = 7.9804 (5) \text{ \AA}$

$c = 16.749 (1) \text{ \AA}$

$V = 1066.69 (11) \text{ \AA}^3$

$Z = 4$

$F(000) = 448$

$D_x = 1.328 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1894 reflections

$\theta = 2.5\text{--}27.8^\circ$

$\mu = 0.29 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Prism, colourless

$0.40 \times 0.18 \times 0.12 \text{ mm}$

Data collection

Oxford Diffraction Xcalibur diffractometer with a Sapphire CCD detector

Radiation source: fine-focus sealed tube graphite

Rotation method data acquisition using ω and φ scans $\theta_{\max} = 26.4^\circ$, $\theta_{\min} = 2.6^\circ$

Absorption correction: multi-scan (*CrysAlis RED*; Oxford Diffraction, 2009)

$T_{\min} = 0.895$, $T_{\max} = 0.967$

4245 measured reflections

1944 independent reflections

1690 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.016$

$h = -9 \rightarrow 5$

$k = -6 \rightarrow 9$

$l = -17 \rightarrow 20$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.035$

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H atoms treated by a mixture of independent and constrained refinement

$wR(F^2) = 0.086$	$w = 1/[\sigma^2(F_o^2) + (0.0453P)^2 + 0.1227P]$
$S = 1.08$	where $P = (F_o^2 + 2F_c^2)/3$
1944 reflections	$(\Delta/\sigma)_{\max} = 0.016$
132 parameters	$\Delta\rho_{\max} = 0.14 \text{ e } \text{\AA}^{-3}$
2 restraints	$\Delta\rho_{\min} = -0.17 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Absolute structure: Flack (1983), 824 Friedel pairs Flack parameter: 0.03 (9)

Special details

Experimental. CrysAlis RED (Oxford Diffraction, 2009) Empirical absorption correction using spherical harmonics, implemented in SCALE3 ABSPACK scaling algorithm.

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	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.0401 (3)	0.5352 (3)	0.16636 (17)	0.0418 (6)
C2	0.0148 (3)	0.3636 (3)	0.15630 (19)	0.0534 (7)
C3	-0.0178 (4)	0.2719 (4)	0.2248 (3)	0.0758 (11)
H3	-0.0359	0.1571	0.2205	0.091*
C4	-0.0243 (5)	0.3456 (5)	0.2992 (2)	0.0844 (12)
H4	-0.0458	0.2805	0.3440	0.101*
C5	0.0008 (4)	0.5134 (5)	0.3071 (2)	0.0730 (10)
H5	-0.0034	0.5631	0.3573	0.088*
C6	0.0326 (4)	0.6095 (4)	0.24026 (19)	0.0571 (7)
H6	0.0490	0.7244	0.2452	0.069*
C7	0.4165 (3)	0.6071 (3)	0.10946 (17)	0.0462 (6)
C8	0.5725 (3)	0.5282 (5)	0.0766 (2)	0.0738 (10)
H8A	0.6102	0.5908	0.0310	0.089*
H8B	0.5489	0.4149	0.0608	0.089*
H8C	0.6582	0.5283	0.1168	0.089*
C9	0.0213 (5)	0.2751 (4)	0.0766 (3)	0.0830 (11)
H9A	0.1276	0.2961	0.0517	0.100*
H9B	-0.0669	0.3162	0.0429	0.100*
H9C	0.0075	0.1568	0.0845	0.100*
N1	0.2818 (3)	0.6051 (3)	0.05903 (12)	0.0413 (5)
H1N	0.278 (3)	0.556 (3)	0.0138 (12)	0.050*
O1	-0.0096 (3)	0.6269 (3)	0.01708 (12)	0.0644 (6)

supplementary materials

O2	0.0971 (3)	0.8346 (2)	0.11214 (13)	0.0632 (6)
O3	0.4067 (2)	0.6678 (3)	0.17544 (11)	0.0588 (5)
S1	0.09057 (8)	0.66608 (8)	0.08490 (5)	0.04393 (18)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0313 (12)	0.0414 (14)	0.0526 (15)	0.0022 (11)	0.0085 (11)	-0.0017 (12)
C2	0.0416 (15)	0.0414 (15)	0.077 (2)	0.0004 (12)	0.0128 (14)	-0.0002 (14)
C3	0.068 (2)	0.0495 (18)	0.110 (3)	0.0058 (16)	0.027 (2)	0.017 (2)
C4	0.078 (2)	0.090 (3)	0.085 (3)	0.011 (2)	0.030 (2)	0.033 (2)
C5	0.069 (2)	0.098 (3)	0.0526 (19)	0.0076 (18)	0.0181 (17)	0.0058 (18)
C6	0.0480 (16)	0.0596 (17)	0.0637 (18)	0.0013 (13)	0.0148 (14)	-0.0082 (16)
C7	0.0392 (14)	0.0478 (14)	0.0515 (17)	-0.0071 (11)	-0.0043 (11)	-0.0001 (13)
C8	0.0401 (15)	0.102 (3)	0.080 (2)	0.0051 (15)	0.0006 (17)	-0.012 (2)
C9	0.084 (2)	0.0521 (18)	0.112 (3)	-0.0123 (16)	0.020 (2)	-0.030 (2)
N1	0.0379 (11)	0.0493 (12)	0.0366 (12)	-0.0012 (9)	0.0021 (9)	-0.0061 (9)
O1	0.0510 (12)	0.0821 (16)	0.0601 (14)	0.0057 (10)	-0.0138 (10)	0.0055 (12)
O2	0.0717 (13)	0.0375 (10)	0.0806 (15)	0.0068 (9)	0.0170 (11)	0.0028 (10)
O3	0.0577 (12)	0.0760 (14)	0.0426 (11)	-0.0075 (10)	-0.0091 (9)	-0.0100 (10)
S1	0.0385 (3)	0.0424 (3)	0.0509 (4)	0.0049 (3)	0.0006 (3)	0.0015 (3)

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

C1—C6	1.374 (4)	C7—N1	1.367 (3)
C1—C2	1.395 (4)	C7—C8	1.500 (4)
C1—S1	1.765 (3)	C8—H8A	0.9600
C2—C3	1.386 (5)	C8—H8B	0.9600
C2—C9	1.511 (5)	C8—H8C	0.9600
C3—C4	1.378 (5)	C9—H9A	0.9600
C3—H3	0.9300	C9—H9B	0.9600
C4—C5	1.361 (6)	C9—H9C	0.9600
C4—H4	0.9300	N1—S1	1.659 (2)
C5—C6	1.380 (5)	N1—H1N	0.855 (17)
C5—H5	0.9300	O1—S1	1.424 (2)
C6—H6	0.9300	O2—S1	1.4213 (19)
C7—O3	1.209 (3)		
C6—C1—C2	121.8 (3)	C7—C8—H8A	109.5
C6—C1—S1	116.8 (2)	C7—C8—H8B	109.5
C2—C1—S1	121.4 (2)	H8A—C8—H8B	109.5
C3—C2—C1	116.5 (3)	C7—C8—H8C	109.5
C3—C2—C9	119.4 (3)	H8A—C8—H8C	109.5
C1—C2—C9	124.1 (3)	H8B—C8—H8C	109.5
C4—C3—C2	122.0 (3)	C2—C9—H9A	109.5
C4—C3—H3	119.0	C2—C9—H9B	109.5
C2—C3—H3	119.0	H9A—C9—H9B	109.5
C5—C4—C3	120.2 (3)	C2—C9—H9C	109.5
C5—C4—H4	119.9	H9A—C9—H9C	109.5

C3—C4—H4	119.9	H9B—C9—H9C	109.5
C4—C5—C6	119.6 (3)	C7—N1—S1	123.94 (18)
C4—C5—H5	120.2	C7—N1—H1N	125.3 (19)
C6—C5—H5	120.2	S1—N1—H1N	109.7 (19)
C1—C6—C5	119.9 (3)	O2—S1—O1	119.00 (13)
C1—C6—H6	120.0	O2—S1—N1	109.12 (11)
C5—C6—H6	120.0	O1—S1—N1	104.10 (12)
O3—C7—N1	121.2 (2)	O2—S1—C1	108.68 (13)
O3—C7—C8	123.9 (3)	O1—S1—C1	110.99 (13)
N1—C7—C8	114.9 (3)	N1—S1—C1	103.79 (11)
C6—C1—C2—C3	-0.2 (4)	O3—C7—N1—S1	-6.3 (4)
S1—C1—C2—C3	177.5 (2)	C8—C7—N1—S1	173.3 (2)
C6—C1—C2—C9	180.0 (3)	C7—N1—S1—O2	57.5 (3)
S1—C1—C2—C9	-2.3 (4)	C7—N1—S1—O1	-174.5 (2)
C1—C2—C3—C4	-0.3 (5)	C7—N1—S1—C1	-58.2 (2)
C9—C2—C3—C4	179.6 (3)	C6—C1—S1—O2	-6.2 (3)
C2—C3—C4—C5	0.3 (6)	C2—C1—S1—O2	176.0 (2)
C3—C4—C5—C6	0.1 (5)	C6—C1—S1—O1	-138.9 (2)
C2—C1—C6—C5	0.6 (4)	C2—C1—S1—O1	43.3 (2)
S1—C1—C6—C5	-177.3 (2)	C6—C1—S1—N1	109.8 (2)
C4—C5—C6—C1	-0.5 (5)	C2—C1—S1—N1	-68.0 (2)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N1—H1N \cdots O3 ⁱ	0.86 (2)	1.95 (2)	2.770 (3)	162 (3)

Symmetry codes: (i) $-y+1, x, z-1/4$.

Fig. 2

