# organic compounds

Acta Crystallographica Section E **Structure Reports** Online

ISSN 1600-5368

# 2-[4-(Methylsulfonyl)phenyl]acetonitrile

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Received 26 January 2011; accepted 30 January 2011

Key indicators: single-crystal X-ray study; T = 296 K; mean  $\sigma$ (C–C) = 0.002 Å; R factor = 0.038; wR factor = 0.110; data-to-parameter ratio = 15.3.

In the title compound,  $C_{9}H_{9}NO_{2}S$ , the benzene ring and the acetonitrile group are approximately coplanar, with a C-C-C-C torsion angle of  $1.1 (3)^{\circ}$  between them. In the crystal, molecules are linked via intermolecular C-H···O hydrogen bonds into layers parallel to (001).

#### **Related literature**

For general background to and the biological activity of COX-2 inhibitors, see: Orjales et al. (2008); Zarghi et al. (2008); Shah et al. (2010); Arico et al. (2002); Davies et al. (2002); Sawaoka et al. (1998); Liu et al. (2000); Pasinetti (2001); Norman et al. (1995). For a related structure, see: Charlier et al. (2004). For bond-length data, see: Allen et al. (1987).



#### **Experimental**

Crystal data

C<sub>9</sub>H<sub>9</sub>NO<sub>2</sub>S  $M_{\rm m} = 195.23$ Triclinic, P1 a = 5.5599 (2) Å b = 8.0942 (3) Å c = 10.9006 (4) Å  $\alpha = 81.162 (2)^{\circ}$  $\beta = 85.347 \ (2)^{\circ}$ 

$\gamma = 74.458 \ (2)^{\circ}$
V = 466.60 (3) Å <sup>3</sup>
Z = 2
Mo $K\alpha$ radiation
$\mu = 0.31 \text{ mm}^{-1}$
T = 296  K
$0.51$ $\times$ $0.28$ $\times$ 0.14 mm

#### Data collection

```
Bruker SMART APEXII CCD
  area-detector diffractometer
Absorption correction: multi-scan
  (SADABS; Bruker, 2009)
  T_{\min} = 0.810, T_{\max} = 0.957
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#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.038$	119 parameters
$wR(F^2) = 0.110$	H-atom parameters constrained
S = 1.09	$\Delta \rho_{\rm max} = 0.29 \ {\rm e} \ {\rm \AA}^{-3}$
1826 reflections	$\Delta \rho_{\rm min} = -0.42 \text{ e } \text{\AA}^{-3}$

5970 measured reflections

 $R_{\rm int} = 0.023$ 

1826 independent reflections

1673 reflections with  $I > 2\sigma(I)$ 

#### Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C5-H5A\cdotsO1^{i}$	0.93	2.47	3.384 (2)	169
$C9-H9B\cdots O2^{ii}$	0.96	2.39	3.343 (3)	175

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

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).

HKF and CKQ thank Universiti Sains Malaysia for the Research University Grant (No. 1001/PFIZIK/811160).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IS2671).

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‡ Thomson Reuters ResearcherID: A-3561-2009. § Thomson Reuters ResearcherID: A-5525-2009.

# supporting information

Acta Cryst. (2011). E67, o574 [doi:10.1107/S1600536811003837]

# 2-[4-(Methylsulfonyl)phenyl]acetonitrile

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### S1. Comment

Compounds bearing the 4-methylsulfonylphenyl moiety are found to possess diverse biological properties. They are found to be highly potent and specific COX-2 inhibitors (Orjales *et al.*, 2008; Zarghi *et al.*, 2008; Shah *et al.*, 2010). Recent studies have shown that selective COX-2 inhibitors can induce apoptosis in colon, stomach, prostate, and breast cancer cell lines (Arico *et al.*, 2002; Davies *et al.*, 2002; Sawaoka *et al.*, 1998; Liu *et al.*, 2000). Selective COX-2 inhibitors offer potential treatment for the prophylactic prevention of inflammatory neurodegerative disorders such as Alzheimer's disease (Pasinetti, 2001). They are also found to be anti-inflammatory agents (Norman *et al.*, 1995). The crystal structure of a methylsulfonylphenyl derivative has been reported (Charlier *et al.*, 2004).

The molecular structure is shown in Fig. 1. Bond lengths (Allen *et al.*, 1987) and angles are within normal ranges. The benzene ring (C1–C6) and the acetonitrile group (C7/C8/N1) are approximately coplanar [torsion angles C1–C6–C7–C8 = 1.1 (3) and C5–C6–C7–C8 = -178.67 (16) °]. In the crystal packing (Fig. 2), the molecules are linked *via* intermolecular C5–H5A···O1 and C9–H9B···O2 (Table 1) hydrogen bonds into infinite two-dimensional planes parallel to (001).

#### **S2. Experimental**

4-Methylthiophenylacetonitrile (0.1 mol) was taken in 3 mL of acetic anhydride and cooled to 5°C. To the reaction mixture sodium tungstate (0.02 mol) was added followed by 30% hydrogen peroxide (0.2 mol) in 1.2 mL of acetic acid and water mixture (in 2:1 ratio). The temperature of the reaction mixture was slowly brought to room temperature. The completion of reaction was monitored by TLC. The solid precipitate was filtered and washed with water until the pH became neutral. The product was dried at 65 °C for 10-12 h. The product was then recrystallized in methanol (*m. p.*: 120–124 °C).

#### **S3. Refinement**

All H atoms were positioned geometrically and refined using a riding model with C—H = 0.93–0.97 Å and  $U_{iso}(H) = 1.2$  or 1.5  $U_{eq}(C)$ . The highest residual electron density peak is located at 0.88 Å from C3 and the deepest hole is located at 0.74 Å from S1. A rotating-group model was applied for the methyl group.



#### Figure 1

The molecular structure of the title compound showing 30% probability displacement ellipsoids for non-H atoms and the atom-numbering scheme.



## Figure 2

The crystal structure of the title compound, viewed along the c axis. H atoms not involved in hydrogen bonds (dashed lines) have been omitted for clarity.

#### 2-[4-(Methylsulfonyl)phenyl]acetonitrile

Crystal data	
$C_9H_9NO_2S$	Hall symbol: -P 1
$M_r = 195.23$	<i>a</i> = 5.5599 (2) Å
Triclinic, P1	b = 8.0942 (3) Å

Mo *K* $\alpha$  radiation,  $\lambda = 0.71073$  Å

 $\theta = 2.6 - 33.0^{\circ}$ 

 $\mu = 0.31 \text{ mm}^{-1}$ T = 296 K

 $R_{\rm int} = 0.023$ 

 $h = -6 \rightarrow 6$ 

 $k = -9 \rightarrow 9$ 

 $l = -13 \rightarrow 13$ 

Block, colourless

 $0.51\times0.28\times0.14~mm$ 

5970 measured reflections 1826 independent reflections

 $\theta_{\rm max} = 26.0^{\circ}, \ \theta_{\rm min} = 1.9^{\circ}$ 

1673 reflections with  $I > 2\sigma(I)$ 

Cell parameters from 4240 reflections

c = 10.9006 (4) Å  $\alpha = 81.162 (2)^{\circ}$   $\beta = 85.347 (2)^{\circ}$   $\gamma = 74.458 (2)^{\circ}$   $V = 466.60 (3) \text{ Å}^{3}$  Z = 2 F(000) = 204 $D_{x} = 1.390 \text{ Mg m}^{-3}$ 

#### Data collection

Bruker SMART APEXII CCD area-detector diffractometer Radiation source: fine-focus sealed tube Graphite monochromator  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Bruker, 2009)  $T_{\min} = 0.810, T_{\max} = 0.957$ 

#### Refinement

Refinement on $F^2$	Secondary atom site location: difference Fourier
Least-squares matrix: full	map
$R[F^2 > 2\sigma(F^2)] = 0.038$	Hydrogen site location: inferred from
$wR(F^2) = 0.110$	neighbouring sites
S = 1.09	H-atom parameters constrained
1826 reflections	$w = 1/[\sigma^2(F_o^2) + (0.054P)^2 + 0.1771P]$
119 parameters	where $P = (F_o^2 + 2F_c^2)/3$
0 restraints	$(\Delta/\sigma)_{\rm max} = 0.001$
Primary atom site location: structure-invariant	$\Delta \rho_{\rm max} = 0.29 \text{ e } \text{\AA}^{-3}$
direct methods	$\Delta \rho_{\rm min} = -0.42 \text{ e} \text{ Å}^{-3}$

#### Special details

**Geometry**. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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  $F^2 > 2sigma(F^2)$  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	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
<b>S</b> 1	0.64766 (8)	0.30347 (6)	0.36428 (4)	0.04502 (19)	
01	0.6714 (4)	0.45775 (18)	0.28743 (16)	0.0728 (5)	
O2	0.4011 (3)	0.2889 (2)	0.40261 (19)	0.0773 (5)	
N1	1.5750 (4)	-0.2897 (2)	-0.02043 (19)	0.0638 (5)	
C1	1.1353 (3)	-0.0135 (2)	0.15700 (17)	0.0427 (4)	
H1A	1.2799	-0.0114	0.1081	0.051*	
C2	1.0162 (3)	0.1282 (2)	0.21644 (17)	0.0429 (4)	
H2A	1.0801	0.2246	0.2080	0.051*	
C3	0.8008 (3)	0.1237 (2)	0.28841 (15)	0.0365 (4)	

C4	0.7055 (3)	-0.0194 (2)	0.30170 (17)	0.0439 (4)
H4A	0.5607	-0.0212	0.3505	0.053*
C5	0.8260 (3)	-0.1598 (2)	0.24231 (18)	0.0445 (4)
H5A	0.7620	-0.2561	0.2512	0.053*
C6	1.0422 (3)	-0.1580 (2)	0.16938 (15)	0.0370 (4)
C7	1.1679 (4)	-0.3160 (2)	0.10645 (18)	0.0464 (4)
H7A	1.2073	-0.4167	0.1694	0.056*
H7B	1.0511	-0.3343	0.0517	0.056*
C8	1.3966 (4)	-0.3021 (2)	0.03441 (18)	0.0468 (4)
С9	0.8180 (4)	0.2777 (3)	0.4977 (2)	0.0586 (5)
H9A	0.7482	0.3732	0.5433	0.088*
H9B	0.9891	0.2744	0.4738	0.088*
H9C	0.8100	0.1716	0.5490	0.088*

Atomic displacement parameters  $(\mathring{A}^2)$ 

$U^{11}$	<i>U</i> <sup>22</sup>	$U^{33}$	$U^{12}$	U <sup>13</sup>	U <sup>23</sup>
0.0405 (3)	0.0414 (3)	0.0506 (3)	-0.00199 (19)	-0.00442 (19)	-0.0129 (2)
0.1054 (14)	0.0367 (8)	0.0691 (10)	-0.0059 (8)	-0.0101 (9)	-0.0042 (7)
0.0381 (8)	0.0876 (12)	0.1116 (14)	-0.0088 (7)	0.0091 (8)	-0.0496 (11)
0.0653 (12)	0.0591 (11)	0.0703 (12)	-0.0208 (9)	0.0182 (10)	-0.0217 (9)
0.0428 (9)	0.0427 (10)	0.0458 (10)	-0.0169 (7)	0.0069 (7)	-0.0099 (7)
0.0466 (10)	0.0382 (9)	0.0479 (10)	-0.0185 (7)	0.0028 (8)	-0.0079 (7)
0.0367 (8)	0.0361 (8)	0.0354 (8)	-0.0072 (7)	-0.0024 (6)	-0.0042 (6)
0.0390 (9)	0.0468 (10)	0.0474 (10)	-0.0153 (8)	0.0042 (7)	-0.0070 (8)
0.0471 (10)	0.0393 (9)	0.0519 (10)	-0.0200 (8)	-0.0002 (8)	-0.0062 (8)
0.0395 (9)	0.0366 (8)	0.0351 (8)	-0.0094 (7)	-0.0051 (7)	-0.0050 (7)
0.0500 (10)	0.0394 (9)	0.0516 (11)	-0.0119 (8)	-0.0004 (8)	-0.0123 (8)
0.0567 (12)	0.0381 (9)	0.0463 (10)	-0.0094 (8)	-0.0016 (9)	-0.0138 (8)
0.0550 (12)	0.0689 (14)	0.0505 (11)	-0.0037 (10)	-0.0073 (9)	-0.0231 (10)
	$U^{11}$ 0.0405 (3) 0.1054 (14) 0.0381 (8) 0.0653 (12) 0.0428 (9) 0.0466 (10) 0.0367 (8) 0.0390 (9) 0.0471 (10) 0.0395 (9) 0.0500 (10) 0.0567 (12) 0.0550 (12)	$\begin{array}{c cccc} U^{11} & U^{22} \\ \hline 0.0405 (3) & 0.0414 (3) \\ \hline 0.1054 (14) & 0.0367 (8) \\ \hline 0.0381 (8) & 0.0876 (12) \\ \hline 0.0653 (12) & 0.0591 (11) \\ \hline 0.0428 (9) & 0.0427 (10) \\ \hline 0.0466 (10) & 0.0382 (9) \\ \hline 0.0367 (8) & 0.0361 (8) \\ \hline 0.0390 (9) & 0.0468 (10) \\ \hline 0.0471 (10) & 0.0393 (9) \\ \hline 0.0395 (9) & 0.0366 (8) \\ \hline 0.0500 (10) & 0.0394 (9) \\ \hline 0.0550 (12) & 0.0689 (14) \\ \end{array}$	$U^{11}$ $U^{22}$ $U^{33}$ $0.0405(3)$ $0.0414(3)$ $0.0506(3)$ $0.1054(14)$ $0.0367(8)$ $0.0691(10)$ $0.0381(8)$ $0.0876(12)$ $0.1116(14)$ $0.0653(12)$ $0.0591(11)$ $0.0703(12)$ $0.0428(9)$ $0.0427(10)$ $0.0458(10)$ $0.0466(10)$ $0.0382(9)$ $0.0479(10)$ $0.0367(8)$ $0.0361(8)$ $0.0354(8)$ $0.0390(9)$ $0.0468(10)$ $0.0474(10)$ $0.0471(10)$ $0.0393(9)$ $0.0519(10)$ $0.0395(9)$ $0.0366(8)$ $0.0351(8)$ $0.0500(10)$ $0.0394(9)$ $0.0463(10)$ $0.0550(12)$ $0.0689(14)$ $0.0505(11)$	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	$U^{11}$ $U^{22}$ $U^{33}$ $U^{12}$ $U^{13}$ $0.0405(3)$ $0.0414(3)$ $0.0506(3)$ $-0.00199(19)$ $-0.00442(19)$ $0.1054(14)$ $0.0367(8)$ $0.0691(10)$ $-0.0059(8)$ $-0.0101(9)$ $0.0381(8)$ $0.0876(12)$ $0.1116(14)$ $-0.0088(7)$ $0.0091(8)$ $0.0653(12)$ $0.0591(11)$ $0.0703(12)$ $-0.0208(9)$ $0.0182(10)$ $0.0428(9)$ $0.0427(10)$ $0.0458(10)$ $-0.0169(7)$ $0.0069(7)$ $0.0466(10)$ $0.0382(9)$ $0.0479(10)$ $-0.0185(7)$ $0.0028(8)$ $0.0367(8)$ $0.0361(8)$ $0.0354(8)$ $-0.0072(7)$ $-0.0024(6)$ $0.0390(9)$ $0.0468(10)$ $0.0474(10)$ $-0.0153(8)$ $0.0042(7)$ $0.0471(10)$ $0.0393(9)$ $0.0519(10)$ $-0.0200(8)$ $-0.0002(8)$ $0.0395(9)$ $0.0366(8)$ $0.0351(8)$ $-0.0094(7)$ $-0.0051(7)$ $0.0500(10)$ $0.0394(9)$ $0.0463(10)$ $-0.0094(8)$ $-0.0016(9)$ $0.0550(12)$ $0.0689(14)$ $0.0505(11)$ $-0.0037(10)$ $-0.0073(9)$

Geometric parameters (Å, °)

<u>81—01</u>	1.4239 (16)	C4—C5	1.381 (3)
S1—O2	1.4306 (16)	C4—H4A	0.9300
S1—C9	1.755 (2)	С5—С6	1.388 (2)
S1—C3	1.7657 (17)	С5—Н5А	0.9300
N1—C8	1.136 (3)	C6—C7	1.519 (2)
C1—C6	1.385 (2)	С7—С8	1.461 (3)
C1—C2	1.387 (2)	С7—Н7А	0.9700
C1—H1A	0.9300	С7—Н7В	0.9700
C2—C3	1.383 (2)	С9—Н9А	0.9600
C2—H2A	0.9300	С9—Н9В	0.9600
C3—C4	1.382 (3)	С9—Н9С	0.9600
01—S1—O2	117.73 (12)	C4—C5—H5A	119.8
O1—S1—C9	108.35 (11)	C6—C5—H5A	119.8
O2—S1—C9	108.32 (12)	C1—C6—C5	119.12 (16)

O1—S1—C3	108.95 (9)	C1—C6—C7	122.56 (16)
O2—S1—C3	108.26 (9)	C5—C6—C7	118.32 (15)
C9—S1—C3	104.42 (9)	C8—C7—C6	113.81 (15)
C6—C1—C2	120.95 (16)	С8—С7—Н7А	108.8
C6—C1—H1A	119.5	С6—С7—Н7А	108.8
C2—C1—H1A	119.5	С8—С7—Н7В	108.8
C3—C2—C1	118.95 (16)	С6—С7—Н7В	108.8
C3—C2—H2A	120.5	H7A—C7—H7B	107.7
C1—C2—H2A	120.5	N1—C8—C7	179.0 (2)
C4—C3—C2	120.80 (16)	S1—C9—H9A	109.5
C4—C3—S1	119.91 (13)	S1—C9—H9B	109.5
C2—C3—S1	119.28 (13)	H9A—C9—H9B	109.5
C5—C4—C3	119.72 (16)	S1—C9—H9C	109.5
C5—C4—H4A	120.1	H9A—C9—H9C	109.5
C3—C4—H4A	120.1	H9B—C9—H9C	109.5
C4—C5—C6	120.45 (16)		
C6—C1—C2—C3	-0.3 (3)	C2—C3—C4—C5	-0.1 (3)
C1—C2—C3—C4	0.3 (3)	S1—C3—C4—C5	-179.93 (13)
C1—C2—C3—S1	-179.94 (13)	C3—C4—C5—C6	0.0 (3)
O1—S1—C3—C4	-145.12 (16)	C2-C1-C6-C5	0.1 (3)
O2—S1—C3—C4	-15.97 (18)	C2-C1-C6-C7	-179.65 (16)
C9—S1—C3—C4	99.29 (17)	C4—C5—C6—C1	0.0 (3)
O1—S1—C3—C2	35.06 (17)	C4—C5—C6—C7	179.80 (16)
O2—S1—C3—C2	164.22 (15)	C1—C6—C7—C8	1.1 (3)
C9—S1—C3—C2	-80.53 (17)	C5—C6—C7—C8	-178.67 (16)

## Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D····A	<i>D</i> —H··· <i>A</i>
C5—H5A…O1 <sup>i</sup>	0.93	2.47	3.384 (2)	169
C9—H9 <i>B</i> ···O2 <sup>ii</sup>	0.96	2.39	3.343 (3)	175

Symmetry codes: (i) *x*, *y*–1, *z*; (ii) *x*+1, *y*, *z*.