

Benzenesulfonamide

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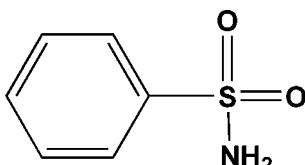
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Key indicators: single-crystal X-ray study; $T = 295$ K; mean $\sigma(C-C) = 0.016$ Å; R factor = 0.074; wR factor = 0.260; data-to-parameter ratio = 11.3.

The structure of the title compound (BSA), $C_6H_7NO_2S$, closely resembles those of other aryl sulfonamides. The geometric parameters in BSA are similar except for some difference in the $S=O$ bond lengths. Molecules are connected by $N-H \cdots O$ hydrogen bonds into layers parallel to the bc plane, with an interlayer distance of 7.734 (2) Å.

Related literature

For related literature, see: Gowda & Shetty (2004); Gowda *et al.* (2002, 2003, 2005, 2007); Jones & Weinkauf (1993); Kumar *et al.* (1992); O'Connor & Maslen (1965).



Experimental

Crystal data

$C_6H_7NO_2S$	$V = 352.52 (15)$ Å ³
$M_r = 157.19$	$Z = 2$
Monoclinic, $P\bar{c}$	Mo $K\alpha$ radiation
$a = 8.304 (2)$ Å	$\mu = 0.39$ mm ⁻¹
$b = 5.534 (1)$ Å	$T = 295 (2)$ K
$c = 8.237 (2)$ Å	$0.52 \times 0.46 \times 0.09$ mm
$\beta = 111.36 (3)^\circ$	

Data collection

Oxford Diffraction Xcalibur diffractometer
Absorption correction: analytical (*CrysAlis RED*; Oxford Diffraction, 2006; Clark & Reid, 1995)
 $T_{\min} = 0.812$, $T_{\max} = 0.956$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.074$
 $wR(F^2) = 0.260$
 $S = 1.01$
1003 reflections
93 parameters
5 restraints

H-atom parameters constrained
 $\Delta\rho_{\max} = 0.19$ e Å⁻³
 $\Delta\rho_{\min} = -0.31$ e Å⁻³
Absolute structure: Flack (1983), with 356 Friedel pairs
Flack parameter: -0.1 (3)

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N1—H1A···O1 ⁱ	0.89	2.20	2.932 (14)	139
N1—H1B···O2 ⁱⁱ	0.89	2.17	3.016 (17)	158

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

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2006); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2006); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*, *PLATON* (Spek, 2003) and *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DN2171).

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Benzenesulfonamide

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Comment

The chemistry of sulfonamides is of interest as they show distinct physical, chemical and biological properties. Many aryl-sulfonamides and their N-halo compounds exhibit pharmacological, fungicidal and herbicidal activities due to their oxidizing action in aqueous, partial aqueous and non-aqueous media. Thus N-halo arylsulfonamides are of interest in synthetic, mechanistic, analytical and biological chemistry (Gowda *et al.*, 2002; 2003; 2005; 2007; Gowda & Shetty, 2004). In the present work, the structure of benzenesulfonamide (BSA) has been determined to explore the substituent effects on the solid state structures of sulfonamides and N-halo arylsulfonamides (Gowda *et al.*, 2003, 2007). The structure of BSA (Fig. 1) closely resembles those of other aryl sulfonamides (Gowda *et al.*, 2003; Jones & Weinkauf, 1993; Kumar *et al.*, 1992; O'Connor & Maslen, 1965). The parent sulphonamide, BSA crystallizes in monoclinic *Pc* space group in contrast to orthorhombic *Pbca* space group observed with 4-fluorobenzenesulfonamide (Jones & Weinkauf, 1993) and 4-aminobenzenesulfonamide (O'Connor & Maslen, 1965) and monoclinic *P21/n* space group observed with 4-chlorobenzenesulfonamide and 4-bromobenzenesulfonamide (Gowda *et al.*, 2003), and 4-methylbenzenesulfonamide (Kumar *et al.*, 1992). The bond parameters in BSA are similar except for some slight differences in the S—O bond lengths.

Molecules are connected by N—H···O hydrogen bonds into layers parallel to the *bc*-plane (Fig. 2), with interlayer distance 7.734 (2) Å.

Experimental

The title compound was prepared according to the literature method (Gowda *et al.*, 2002, 2003). The purity of the compound was checked by determining its melting point. It was characterized by recording its infrared and NMR spectra (Gowda *et al.*, 2002). Single crystals of the title compound were obtained from a slow evaporation of its ethanolic solution and used for X-ray diffraction studies at room temperature.

Refinement

Monoclinic crystal (1) was refined as twinned, with two twin domains (fractional contributions 84 and 16 percent). The non-merohedral twinning was analysed using TwinRotMat routine within *WinGX* package. It was found that 22 percent of total reflections were overlapped with rotation matrix (1.000 0.000 0.734) (0.000 – 1.000 0.000) (0.000 0.000 – 1.000) Using the above twin matrix a HKLF5 file was generated which was subsequently used in the *SHELXL97* refinement of the structure. The BASF parameter was refined to final value 0.154.

Hydrogen atoms attached to carbons were positioned geometrically and treated as riding with C—H = 0.93 Å. H atoms attached to N1 atom were placed in positions with N—H bond distance restrained to 0.89 (2) Å and H—H restrained to 1.50 (3) Å. In the last stage of refinement, these H were treated as riding on their parent N atom with $U_{\text{iso}}(\text{H})=1.2 U_{\text{eq}}(\text{N})$.

Although the N atom has an elongated ellipsoids, no reasonable disordered model could be defined.

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Owing to the poor quality of the data, the absolute structure couldn't be reliably defined and any references to the Flack parameter have been omitted. The Friedel pairs were merged.

Figures

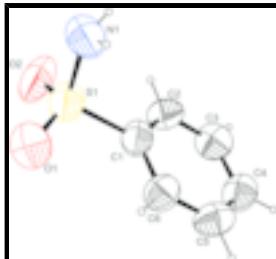


Fig. 1. *ORTEP* view of the title compound showing the atom labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are represented as small spheres of arbitrary radii.

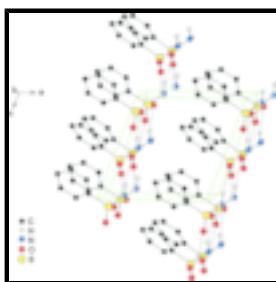


Fig. 2. Packing of the title compound viewed down the *b* axis, showing molecular layers parallel to the *bc*-plane. Within a layer the molecules are connected by N—H···O hydrogen bonds (dashed lines). H atoms not involved in hydrogen bonding have been omitted for clarity.

Benzenesulfonamide

Crystal data

C ₆ H ₇ NO ₂ S	$F_{000} = 164$
$M_r = 157.19$	$D_x = 1.481 \text{ Mg m}^{-3}$
Monoclinic, <i>Pc</i>	Mo <i>K</i> α radiation
Hall symbol: P -2yc	$\lambda = 0.71073 \text{ \AA}$
$a = 8.304 (2) \text{ \AA}$	Cell parameters from 706 reflections
$b = 5.534 (1) \text{ \AA}$	$\theta = 2.0\text{--}30.0^\circ$
$c = 8.237 (2) \text{ \AA}$	$\mu = 0.39 \text{ mm}^{-1}$
$\beta = 111.36 (3)^\circ$	$T = 295 (2) \text{ K}$
$V = 352.52 (15) \text{ \AA}^3$	Plate, colourless
$Z = 2$	$0.52 \times 0.46 \times 0.09 \text{ mm}$

Data collection

Oxford Diffraction Xcalibur diffractometer	621 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\text{int}} = 0.086$
$T = 295(2) \text{ K}$	$\theta_{\max} = 25.7^\circ$
Rotation method data acquisition using ω and phi scans	$\theta_{\min} = 4.5^\circ$
Absorption correction: analytical	$h = -10 \rightarrow 10$

(CrysAlis RED; Oxford Diffraction, 2006; Clark & Reid, 1995)

$T_{\min} = 0.812$, $T_{\max} = 0.956$

2613 measured reflections

1003 independent reflections

$k = -6 \rightarrow 6$

$l = -10 \rightarrow 8$

Refinement

Refinement on F^2

Hydrogen site location: inferred from neighbouring sites

Least-squares matrix: full

H-atom parameters constrained

$R[F^2 > 2\sigma(F^2)] = 0.074$

$$w = 1/[\sigma^2(F_o^2) + (0.1777P)^2]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$wR(F^2) = 0.260$

$$(\Delta/\sigma)_{\max} < 0.001$$

$S = 1.01$

$$\Delta\rho_{\max} = 0.19 \text{ e \AA}^{-3}$$

1003 reflections

$$\Delta\rho_{\min} = -0.31 \text{ e \AA}^{-3}$$

93 parameters

Extinction correction: none

5 restraints

Absolute structure: Flack (1983), with 356 Friedel pairs

Primary atom site location: structure-invariant direct methods

Flack parameter: -0.1 (3)

Secondary atom site location: difference Fourier map

Special details

Experimental. Analytical numeric absorption correction using a multifaceted crystal model based on expressions derived by Clark & Reid, 1995).

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 $F^2 > 2\sigma(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	1.1057 (12)	0.6751 (12)	0.7719 (12)	0.124 (2)
O2	1.0908 (12)	1.0728 (16)	0.6495 (18)	0.153 (4)
S1	1.0618 (2)	0.8225 (4)	0.6248 (2)	0.1057 (11)
C1	0.8412 (11)	0.7887 (19)	0.4996 (12)	0.095 (3)
C2	0.7597 (12)	0.9503 (16)	0.3760 (14)	0.096 (2)
H2	0.8191	1.0812	0.3544	0.115*
C3	0.5887 (15)	0.919 (2)	0.2829 (17)	0.111 (3)
H3	0.5293	1.0355	0.2016	0.134*

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C4	0.5006 (14)	0.717 (2)	0.3070 (17)	0.111 (3)
H4	0.3844	0.6934	0.2395	0.133*
C5	0.5888 (16)	0.5511 (17)	0.4325 (17)	0.109 (3)
H5	0.5307	0.4170	0.4519	0.130*
C6	0.7579 (12)	0.5802 (16)	0.5278 (13)	0.096 (2)
H6	0.8182	0.4656	0.6100	0.115*
N1	1.1731 (11)	0.734 (3)	0.5134 (11)	0.149 (4)
H1A	1.1665	0.5763	0.4933	0.179*
H1B	1.1408	0.8270	0.4185	0.179*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.131 (6)	0.116 (5)	0.115 (5)	0.014 (4)	0.035 (5)	-0.001 (4)
O2	0.092 (5)	0.168 (7)	0.185 (9)	-0.035 (5)	0.036 (6)	-0.078 (7)
S1	0.0844 (13)	0.126 (2)	0.1070 (17)	0.0067 (16)	0.0350 (11)	-0.0243 (18)
C1	0.071 (4)	0.122 (7)	0.094 (6)	0.000 (5)	0.032 (4)	-0.032 (5)
C2	0.083 (5)	0.098 (6)	0.114 (7)	0.003 (4)	0.045 (5)	0.011 (5)
C3	0.099 (6)	0.116 (7)	0.121 (7)	0.026 (6)	0.043 (6)	0.007 (7)
C4	0.082 (5)	0.126 (7)	0.129 (9)	0.000 (5)	0.044 (5)	-0.029 (7)
C5	0.114 (8)	0.088 (6)	0.137 (10)	-0.008 (5)	0.062 (7)	0.005 (6)
C6	0.092 (6)	0.087 (5)	0.109 (7)	-0.003 (4)	0.036 (5)	-0.006 (5)
N1	0.067 (4)	0.283 (13)	0.097 (6)	0.045 (7)	0.029 (4)	-0.020 (7)

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

O1—S1	1.394 (8)	C3—H3	0.9300
O2—S1	1.408 (9)	C4—C5	1.376 (16)
S1—N1	1.597 (8)	C4—H4	0.9300
S1—C1	1.755 (9)	C5—C6	1.346 (15)
C1—C2	1.339 (13)	C5—H5	0.9300
C1—C6	1.407 (15)	C6—H6	0.9300
C2—C3	1.356 (16)	N1—H1A	0.8885
C2—H2	0.9300	N1—H1B	0.8901
C3—C4	1.393 (18)		
O1—S1—O2	118.2 (7)	C4—C3—H3	119.4
O1—S1—N1	107.3 (6)	C5—C4—C3	118.7 (10)
O2—S1—N1	106.2 (8)	C5—C4—H4	120.7
O1—S1—C1	109.5 (5)	C3—C4—H4	120.7
O2—S1—C1	106.1 (6)	C6—C5—C4	121.0 (9)
N1—S1—C1	109.2 (4)	C6—C5—H5	119.5
C2—C1—C6	122.0 (9)	C4—C5—H5	119.5
C2—C1—S1	120.4 (8)	C5—C6—C1	118.3 (9)
C6—C1—S1	117.5 (8)	C5—C6—H6	120.9
C1—C2—C3	118.7 (9)	C1—C6—H6	120.9
C1—C2—H2	120.6	S1—N1—H1A	113.4
C3—C2—H2	120.6	S1—N1—H1B	106.0
C2—C3—C4	121.2 (10)	H1A—N1—H1B	115.1

C2—C3—H3 119.4

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>
N1—H1A···O1 ⁱ	0.89	2.20	2.932 (14)	139
N1—H1B···O2 ⁱⁱ	0.89	2.17	3.016 (17)	158

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

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Fig. 1

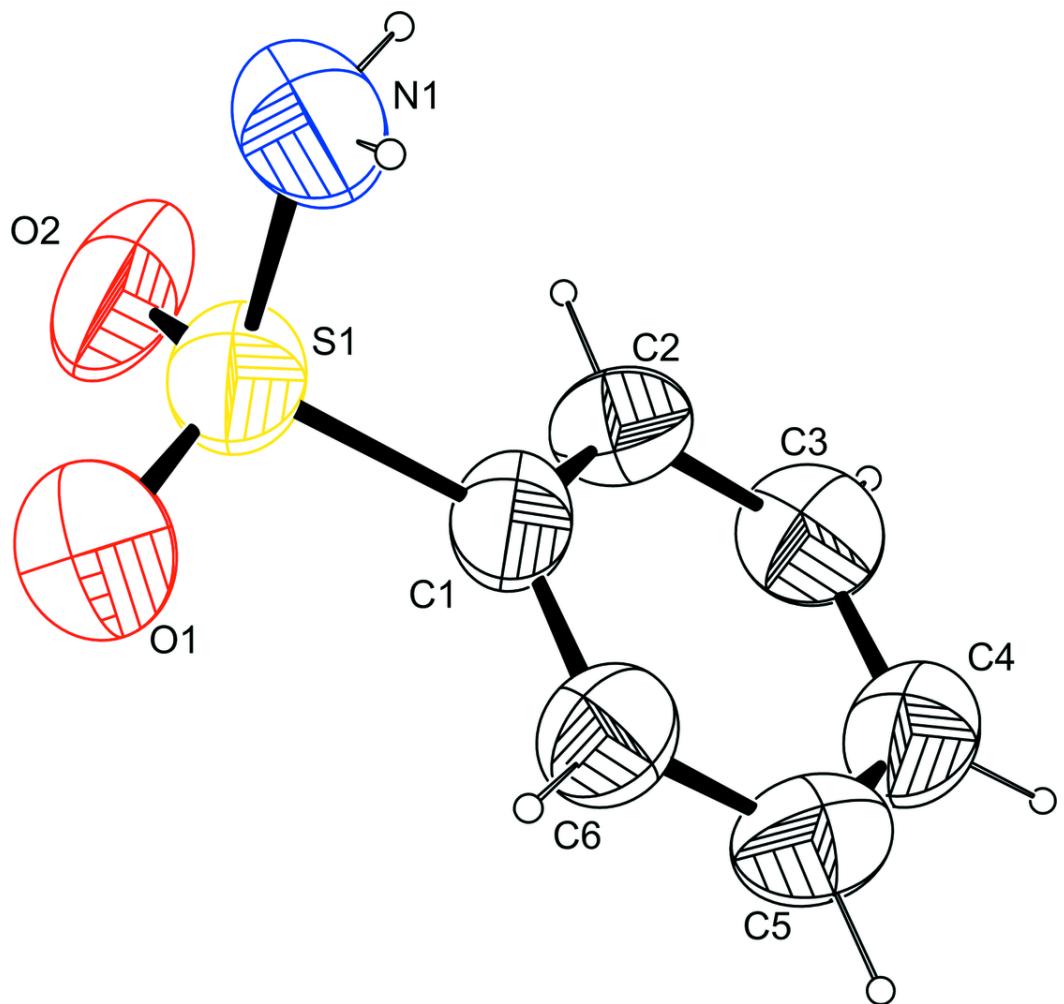


Fig. 2

