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## Structure Reports

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***N*-Phenyl-*tert*-butanesulfonamide**Mritunjoy Datta,<sup>a</sup> Alan J. Buglass<sup>a\*</sup> and Mark R. J. Elsegood<sup>b</sup><sup>a</sup>Department of Chemistry, Korea Advanced Institute of Science and Technology, Daejeon 305-701, Republic of Korea, and <sup>b</sup>Chemistry Department, Loughborough University, Loughborough LE11 3TU, EnglandCorrespondence e-mail: [ajbuglass@kaist.ac.kr](mailto:ajbuglass@kaist.ac.kr)

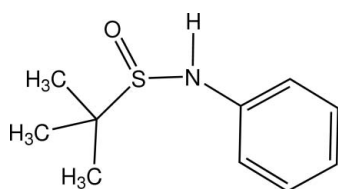
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Key indicators: single-crystal X-ray study;  $T = 150$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.034;  $wR$  factor = 0.092; data-to-parameter ratio = 25.2.

In the racemic title compound,  $\text{C}_{10}\text{H}_{15}\text{NOS}$ , the packing exhibits centrosymmetric pairs of molecules linked by  $\text{N}-\text{H}\cdots\text{O}=\text{S}$  hydrogen bonds in a head-to-tail fashion. The  $\text{N}-\text{C}_{\text{aryl}}$  bond [1.4083 (12) Å] is considerably shorter than the  $\text{N}-\text{C}_{\text{alkyl}}$  bonds typically found in *N*-alkylalkanesulfonamides (1.470–1.530 Å).

## Related literature

For *N*-arylalkanesulfonamides, see: Datta *et al.* (2008) and for cyclic *N*-arylalkanesulfonamides (sultims), see: Schulze *et al.* (2005). For *N*-alkylalkanesulfonamides, see: Sato *et al.* (1975); Schuckmann *et al.* (1978); Ferreira *et al.* (2005). For the synthesis, see: Stretter *et al.* (1969).



## Experimental

## Crystal data

 $\text{C}_{10}\text{H}_{15}\text{NOS}$   
 $M_r = 197.29$   
 Monoclinic,  $P2_1/c$ 
 $a = 7.4822$  (3) Å  
 $b = 15.7881$  (6) Å  
 $c = 8.8333$  (4) Å

 $\beta = 99.3865$  (6)°  
 $V = 1029.50$  (7) Å<sup>3</sup>  
 $Z = 4$   
 Mo  $K\alpha$  radiation

 $\mu = 0.28$  mm<sup>-1</sup>  
 $T = 150$  K  
 $0.54 \times 0.49 \times 0.39$  mm

## Data collection

 Bruker APEXII CCD  
 diffractometer  
 Absorption correction: multi-scan  
 (*SADABS*; Sheldrick, 2007)  
 $T_{\text{min}} = 0.866$ ,  $T_{\text{max}} = 0.900$ 

 12022 measured reflections  
 3150 independent reflections  
 2861 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.020$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.034$   
 $wR(F^2) = 0.092$   
 $S = 1.05$   
 3150 reflections  
 125 parameters  
 1 restraint

 H atoms treated by a mixture of  
 independent and constrained  
 refinement  
 $\Delta\rho_{\text{max}} = 0.38$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.36$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1}\cdots\text{O1}^i$	0.808 (12)	2.085 (12)	2.8882 (11)	173.1 (14)

Symmetry code: (i)  $-x, -y + 1, -z + 1$ .

Data collection: *APEX2* (Bruker, 2006); cell refinement: *SAINT* (Bruker, 2006); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* and local programs.

MD and AJB thank KAIST for financial support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: ZS2002).

## References

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## supporting information

*Acta Cryst.* (2009). E65, o2034 [doi:10.1107/S1600536809028633]

***N*-Phenyl-*tert*-butanesulfinamide****Mritunjoy Datta, Alan J. Buglass and Mark R. J. Elsegood****S1. Comment**

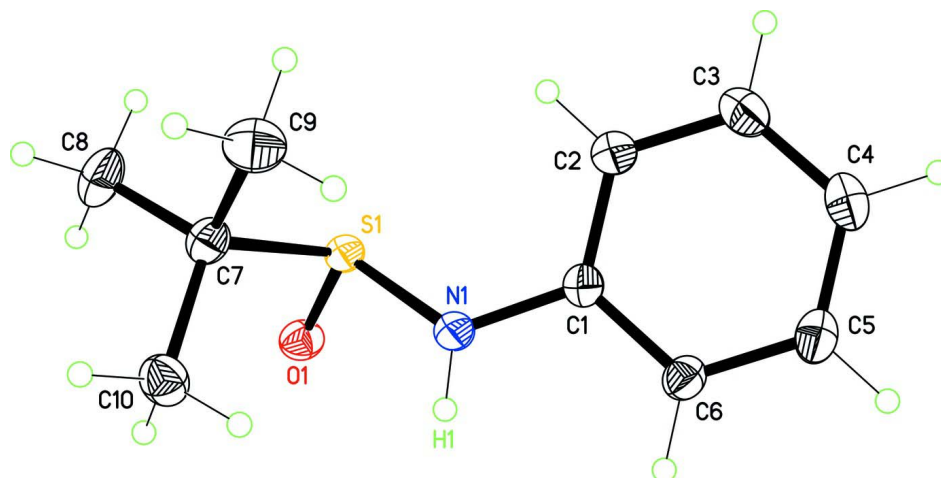
The molecular structure of (I) (Fig. 1) exhibits a short *N*—(aryl)C Bond (1.4083 (12) Å), like that in *N*-phenyladamantane-1-sulfinamide (1.409 (2) Å) (Datta *et al.*, 2008), and in contrast with *N*—(alkyl)C bonds in *N*-alkylalkanesulfinamides (1.470–1.530 Å) (Sato *et al.*, 1975; Schuckmann *et al.*, 1978; Ferreira *et al.*, 2005). Otherwise, molecular geometry is similar to other sulfinamides. The molecules of (I) in the crystal lattice are linked by pairs of N—H···O=S hydrogen bonds (Fig. 2 and Table 1). There is no evidence of weak intermolecular C—H···O=S hydrogen bonds, as in the packing of *N*-phenyladamantane-1-sulfinamide (Datta *et al.*, 2008). The crystal system and space group for (I) and *N*-phenyladamantane-1-sulfinamide are the same (namely monoclinic and *P*2<sub>1</sub>/*c*, respectively).

**S2. Experimental**

Compound (I) was prepared by the method of Stretter *et al.* (1969), using *tert*-butanesulfinyl chloride (702.5 mg, 5 mmol) and aniline (930 mg, 10 mmol) in dry chloroform (30 ml). After 6 h (with TLC monitoring) the white solid amine salt was filtered off and the solvent was removed under reduced pressure. Column chromatography (silica gel, dichloromethane) provided (I) as white crystals (950 mg, 96%), m.p. 376–377 K. Single crystals suitable for X-ray analysis were obtained by evaporation of a solution of (I) in dichloromethane at room temperature. Spectroscopic analysis: FTIR (KBr) (cm<sup>-1</sup>) 3015, 2599, 2330, 1496, 1469, 1420, 1370, 1274, 1068, 1026, 888, 859. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub> p.p.m. with respect to TMS) δ 7.26–7.22 (m, 2H), 7.00–6.98 (m, 3H), 5.48 (bs, 1H), 1.30 (s, 9H). <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub> p.p.m. with respect to TMS) δ 142.0, 129.4, 123.0, 118.4, 56.5, 22.4. EIMS *m/z* (%) 197 (*M*<sup>+</sup>, 18) 141 (100), 140 (*M*<sup>+</sup>-*t*Bu, 28), 105 (*M*<sup>+</sup>-PhNH, 86), 92 (*M*<sup>+</sup>-*t*BuSO, 72), 78 (77), 57 (93).

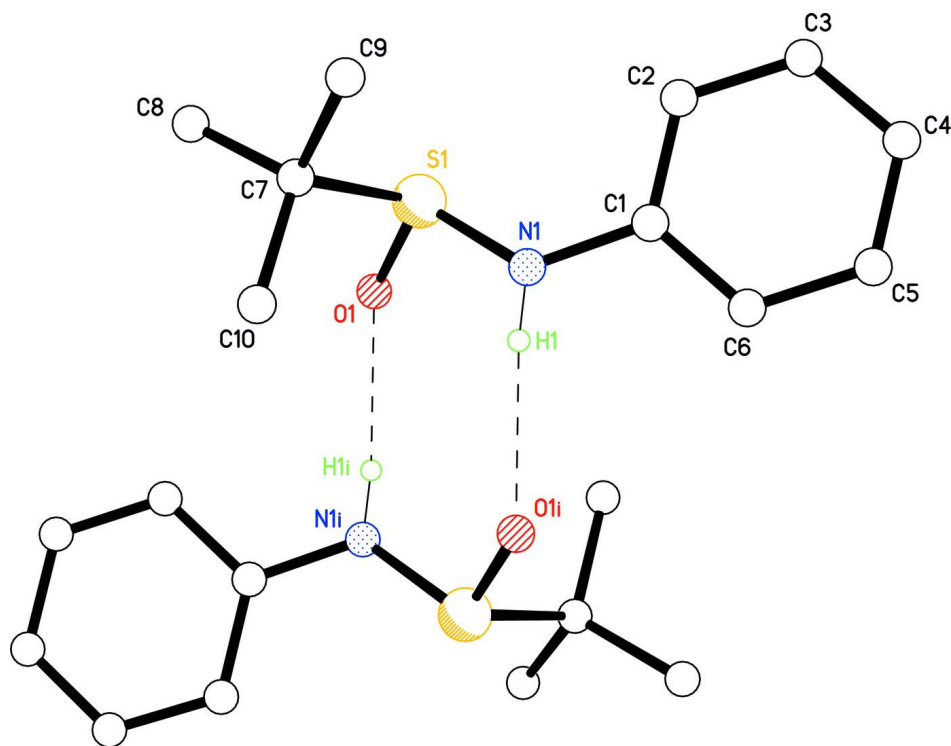
**S3. Refinement**

H atoms were located in a difference Fourier map and refined geometrically using a riding model except for NH for which the coordinates were freely refined. Bond lengths and displacement parameters were constrained as follows: C—H = 0.95–0.98 Å and  $U_{iso}(H) = 1.2$  (1.5 for CH<sub>3</sub>) times  $U_{eq}(C, N)$ .



**Figure 1**

Molecular structure of (I), with atom labels and 50% probability displacement ellipsoids for non-H atoms.



**Figure 2**

Intermolecular hydrogen bonding between a pair of opposite enantiomers of (I) in the crystal packing. Symmetry code  $i = -x, -y + 1, -z + 1$ .

### ***N*-Phenyl-*tert*-butanesulfinamide**

#### *Crystal data*

$C_{10}H_{15}NOS$

$M_r = 197.29$

Monoclinic,  $P2_1/c$

Hall symbol:  $-P 2_1bc$

$a = 7.4822 (3) \text{ \AA}$

$b = 15.7881 (6) \text{ \AA}$

$c = 8.8333$  (4) Å  
 $\beta = 99.3865$  (6)°  
 $V = 1029.50$  (7) Å<sup>3</sup>  
 $Z = 4$   
 $F(000) = 424$   
 $D_x = 1.273$  Mg m<sup>-3</sup>  
 Melting point = 376–377 K

Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å  
 Cell parameters from 6597 reflections  
 $\theta = 2.3$ – $30.5$ °  
 $\mu = 0.28$  mm<sup>-1</sup>  
 $T = 150$  K  
 Block, colourless  
 $0.54 \times 0.49 \times 0.39$  mm

*Data collection*

Bruker APEXII CCD  
 diffractometer  
 Radiation source: fine-focus sealed tube  
 Graphite monochromator  
 $\omega$  rotation with narrow frames scans  
 Absorption correction: multi-scan  
 (SADABS; Sheldrick, 2007)  
 $T_{\min} = 0.866$ ,  $T_{\max} = 0.900$

12022 measured reflections  
 3150 independent reflections  
 2861 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.020$   
 $\theta_{\text{max}} = 30.6$ °,  $\theta_{\text{min}} = 2.7$ °  
 $h = -10 \rightarrow 10$   
 $k = -22 \rightarrow 22$   
 $l = -12 \rightarrow 12$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.034$   
 $wR(F^2) = 0.092$   
 $S = 1.05$   
 3150 reflections  
 125 parameters  
 1 restraint  
 Primary atom site location: structure-invariant  
 direct methods  
 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0516P)^2 + 0.2529P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.002$   
 $\Delta\rho_{\text{max}} = 0.38$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.36$  e Å<sup>-3</sup>  
 Extinction correction: SHELXL97 (Sheldrick,  
 2008),  $F_c^* = kFc[1 + 0.001x\text{Fc}^2\lambda^3/\sin(2\theta)]^{-1/4}$   
 Extinction coefficient: 0.020 (3)

*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 (Å<sup>2</sup>)*

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	-0.06049 (10)	0.60738 (4)	0.41648 (9)	0.02295 (16)
S1	0.10835 (3)	0.601595 (13)	0.34614 (3)	0.01896 (9)
N1	0.17155 (13)	0.50050 (5)	0.35165 (10)	0.02446 (18)
H1	0.1463 (19)	0.4727 (9)	0.4219 (15)	0.029*
C1	0.19492 (12)	0.45439 (6)	0.21979 (11)	0.01998 (18)
C2	0.24483 (16)	0.49296 (7)	0.09080 (12)	0.0281 (2)
H2	0.2589	0.5527	0.0877	0.034*
C3	0.27381 (17)	0.44344 (7)	-0.03327 (13)	0.0318 (2)

H3	0.3053	0.4700	-0.1218	0.038*
C4	0.25757 (15)	0.35617 (7)	-0.02991 (13)	0.0295 (2)
H4	0.2790	0.3230	-0.1148	0.035*
C5	0.20955 (14)	0.31769 (6)	0.09922 (12)	0.0257 (2)
H5	0.1990	0.2578	0.1029	0.031*
C6	0.17690 (13)	0.36629 (6)	0.22303 (11)	0.02134 (18)
H6	0.1422	0.3396	0.3101	0.026*
C7	0.29072 (13)	0.64579 (6)	0.48984 (11)	0.02172 (18)
C8	0.24368 (16)	0.73950 (7)	0.50283 (14)	0.0316 (2)
H8A	0.3444	0.7687	0.5668	0.047*
H8B	0.1342	0.7448	0.5497	0.047*
H8C	0.2223	0.7650	0.4003	0.047*
C9	0.46486 (15)	0.63537 (8)	0.42278 (14)	0.0312 (2)
H9A	0.5658	0.6613	0.4919	0.047*
H9B	0.4512	0.6632	0.3224	0.047*
H9C	0.4894	0.5750	0.4110	0.047*
C10	0.30068 (15)	0.60073 (6)	0.64357 (12)	0.0265 (2)
H10A	0.3290	0.5408	0.6314	0.040*
H10B	0.1839	0.6057	0.6794	0.040*
H10C	0.3956	0.6267	0.7188	0.040*

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0218 (3)	0.0215 (3)	0.0264 (4)	0.0011 (2)	0.0067 (3)	0.0021 (2)
S1	0.02263 (13)	0.01561 (12)	0.01895 (13)	0.00052 (7)	0.00429 (8)	0.00155 (7)
N1	0.0397 (5)	0.0153 (3)	0.0206 (4)	0.0013 (3)	0.0114 (3)	0.0012 (3)
C1	0.0214 (4)	0.0187 (4)	0.0202 (4)	0.0003 (3)	0.0045 (3)	-0.0011 (3)
C2	0.0397 (6)	0.0210 (4)	0.0254 (5)	-0.0014 (4)	0.0112 (4)	0.0010 (4)
C3	0.0422 (6)	0.0327 (6)	0.0224 (5)	0.0011 (4)	0.0111 (4)	0.0006 (4)
C4	0.0324 (5)	0.0316 (5)	0.0251 (5)	0.0014 (4)	0.0063 (4)	-0.0080 (4)
C5	0.0247 (5)	0.0218 (4)	0.0304 (5)	-0.0010 (3)	0.0043 (4)	-0.0060 (4)
C6	0.0207 (4)	0.0186 (4)	0.0252 (4)	-0.0012 (3)	0.0053 (3)	-0.0006 (3)
C7	0.0218 (4)	0.0186 (4)	0.0242 (4)	0.0006 (3)	0.0022 (3)	-0.0001 (3)
C8	0.0352 (5)	0.0183 (4)	0.0387 (6)	-0.0002 (4)	-0.0013 (4)	-0.0037 (4)
C9	0.0230 (5)	0.0366 (6)	0.0346 (6)	0.0004 (4)	0.0065 (4)	0.0042 (4)
C10	0.0290 (5)	0.0278 (5)	0.0220 (5)	0.0026 (4)	0.0020 (4)	0.0006 (3)

*Geometric parameters (Å, °)*

O1—S1	1.4988 (7)	C5—H5	0.9500
S1—N1	1.6632 (9)	C6—H6	0.9500
S1—C7	1.8426 (10)	C7—C10	1.5241 (14)
N1—C1	1.4083 (12)	C7—C9	1.5255 (14)
N1—H1	0.808 (12)	C7—C8	1.5294 (14)
C1—C2	1.3954 (13)	C8—H8A	0.9800
C1—C6	1.3982 (12)	C8—H8B	0.9800
C2—C3	1.3918 (15)	C8—H8C	0.9800

C2—H2	0.9500	C9—H9A	0.9800
C3—C4	1.3838 (17)	C9—H9B	0.9800
C3—H3	0.9500	C9—H9C	0.9800
C4—C5	1.3903 (15)	C10—H10A	0.9800
C4—H4	0.9500	C10—H10B	0.9800
C5—C6	1.3898 (13)	C10—H10C	0.9800
O1—S1—N1	107.53 (4)	C10—C7—C9	112.02 (8)
O1—S1—C7	105.72 (4)	C10—C7—C8	111.29 (9)
N1—S1—C7	99.69 (5)	C9—C7—C8	110.80 (9)
C1—N1—S1	123.02 (7)	C10—C7—S1	111.11 (7)
C1—N1—H1	115.5 (10)	C9—C7—S1	105.97 (7)
S1—N1—H1	116.3 (10)	C8—C7—S1	105.33 (7)
C2—C1—C6	119.36 (9)	C7—C8—H8A	109.5
C2—C1—N1	122.39 (8)	C7—C8—H8B	109.5
C6—C1—N1	118.16 (8)	H8A—C8—H8B	109.5
C3—C2—C1	119.65 (10)	C7—C8—H8C	109.5
C3—C2—H2	120.2	H8A—C8—H8C	109.5
C1—C2—H2	120.2	H8B—C8—H8C	109.5
C4—C3—C2	121.12 (10)	C7—C9—H9A	109.5
C4—C3—H3	119.4	C7—C9—H9B	109.5
C2—C3—H3	119.4	H9A—C9—H9B	109.5
C3—C4—C5	119.21 (9)	C7—C9—H9C	109.5
C3—C4—H4	120.4	H9A—C9—H9C	109.5
C5—C4—H4	120.4	H9B—C9—H9C	109.5
C6—C5—C4	120.43 (9)	C7—C10—H10A	109.5
C6—C5—H5	119.8	C7—C10—H10B	109.5
C4—C5—H5	119.8	H10A—C10—H10B	109.5
C5—C6—C1	120.21 (9)	C7—C10—H10C	109.5
C5—C6—H6	119.9	H10A—C10—H10C	109.5
C1—C6—H6	119.9	H10B—C10—H10C	109.5
O1—S1—N1—C1	123.13 (8)	C4—C5—C6—C1	-1.02 (15)
C7—S1—N1—C1	-126.86 (8)	C2—C1—C6—C5	0.40 (14)
S1—N1—C1—C2	28.58 (14)	N1—C1—C6—C5	-176.09 (9)
S1—N1—C1—C6	-155.05 (8)	O1—S1—C7—C10	54.89 (8)
C6—C1—C2—C3	0.76 (16)	N1—S1—C7—C10	-56.55 (8)
N1—C1—C2—C3	177.09 (10)	O1—S1—C7—C9	176.79 (7)
C1—C2—C3—C4	-1.32 (18)	N1—S1—C7—C9	65.35 (7)
C2—C3—C4—C5	0.70 (17)	O1—S1—C7—C8	-65.73 (8)
C3—C4—C5—C6	0.48 (16)	N1—S1—C7—C8	-177.17 (7)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1 $\cdots$ O1 <sup>i</sup>	0.81 (1)	2.09 (1)	2.8882 (11)	173 (1)

Symmetry code: (i)  $-x, -y+1, -z+1$ .