

N-[2-(Phenylsulfonyl)ethyl]benzylamine¹

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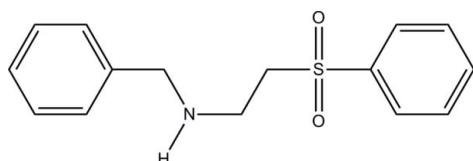
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Key indicators: single-crystal X-ray study; $T = 298\text{ K}$, $P = 0.0\text{ kPa}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.029; wR factor = 0.086; data-to-parameter ratio = 18.4.

The title compound, $C_{15}H_{17}NO_2S$, exhibits intramolecular hydrogen bonding between the amine H atom and a sulfonyl O atom. The conformation of the molecule is described by the four $\text{PhCH}_2-\text{NH}-\text{CH}_2-\text{CH}_2-\text{SO}_2\text{Ph}$ torsion angles of $79.6(2)$, $-166.21(14)$, $-70.29(17)$ and $-58.93(13)^\circ$.

Related literature

For the synthesis, see: Bandini *et al.* (2008). For reactions of benzylamine and vinylsulfonylbenzene, see: Makosza *et al.* (2008); Ni *et al.* (2003). For the determination of absolute structure from Bijvoet pairs, see: Hooft *et al.* (2008). For intermolecular interactions, see: Steiner (1996).

**Experimental***Crystal data*

$C_{15}H_{17}NO_2S$

$M_r = 275.36$

Monoclinic, $P2_1$

$a = 5.7428(11)\text{ \AA}$

$b = 10.170(2)\text{ \AA}$

$c = 12.486(2)\text{ \AA}$

$\beta = 102.09(3)^\circ$

$V = 713.1(2)\text{ \AA}^3$

$Z = 2$

$Mo K\alpha$ radiation

$\mu = 0.22\text{ mm}^{-1}$

$T = 298\text{ K}$

$0.50 \times 0.45 \times 0.32\text{ mm}$

Data collection

Enraf–Nonius CAD-4
diffractometer

Absorption correction: ψ scan
(North *et al.*, 1968)

$T_{\min} = 0.896$, $T_{\max} = 0.932$
5320 measured reflections

3264 independent reflections

2990 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.021$

3 standard reflections every 60 min
intensity decay: 4.0%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.029$

$wR(F^2) = 0.086$

$S = 0.99$

3264 reflections

177 parameters

1 restraint

H atoms treated by a mixture of
independent and constrained
refinement

$\Delta\rho_{\text{max}} = 0.15\text{ e \AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.22\text{ e \AA}^{-3}$

Absolute structure: Flack (1983),

2081 Bijvoet pairs

Flack parameter: 0.07 (6)

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1 \cdots O1	0.87 (2)	2.52 (2)	3.071 (2)	121.9 (17)

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); cell refinement: *CAD-4 EXPRESS*; data reduction: *XCAD4* (Farrugia, 1999); program(s) used to solve structure: *SIR2002* (Burla *et al.*, 2003); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

The purchase of the diffractometer was made possible by a National Science Foundation chemical instrumentation grant, which we gratefully acknowledge. Improvements to the LSU X-ray Crystallography Facility were supported by grant No. LEQSF(1196-97)-ENH-TR-10, administered by the Louisiana Board of Regents. We thank Dr J. Gabriel Garcia for providing the sample.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BG2477).

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¹ CAS 550350-59-9.

supplementary materials

Acta Cryst. (2012). E68, o2940 [doi:10.1107/S160053681203886X]

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Comment

The most striking feature of title compound (**I**), a secondary amine, is the intramolecular hydrogen bond between a sulfonyl oxygen and the amine hydrogen. The complete geometry of this system is: N—H = 0.87 (2) Å, H···O = 2.52 (2) Å, N—H···O = 122 (2)°, S=O···H = 97 (2)°. The two six-rings are flat: C1—C6, $\delta_{\text{r.m.s.}} = 0.0049$ Å, C10—C15, $\delta_{\text{r.m.s.}} = 0.0024$ Å. The absolute structure was determined by analysis of 2081 Bijvoet pairs.

In addition to the intramolecular hydrogen bond, weaker intermolecular C—H···O and C—H··· π interactions (Steiner, 1996) are present. The geometry of the C—H···O interaction is C7—H7A···O1 (at $x + 1, y, z$), C7···O1 3.3896 (18) Å, H7A···O1 2.44 Å, angle about H 166°, forming chains in the [100] direction. The C—H··· π interaction involves the phenyl C4—H as donor and the other phenyl ring C10—C15 (at $1 - x, y - 1/2, 1 - z$) as acceptor. The C4···Cg distance is 3.731 (2) Å, and the contact is fairly linear, with angle about H 174°. This interaction forms chains in the [010] direction, propagated by the screw axis.

Experimental

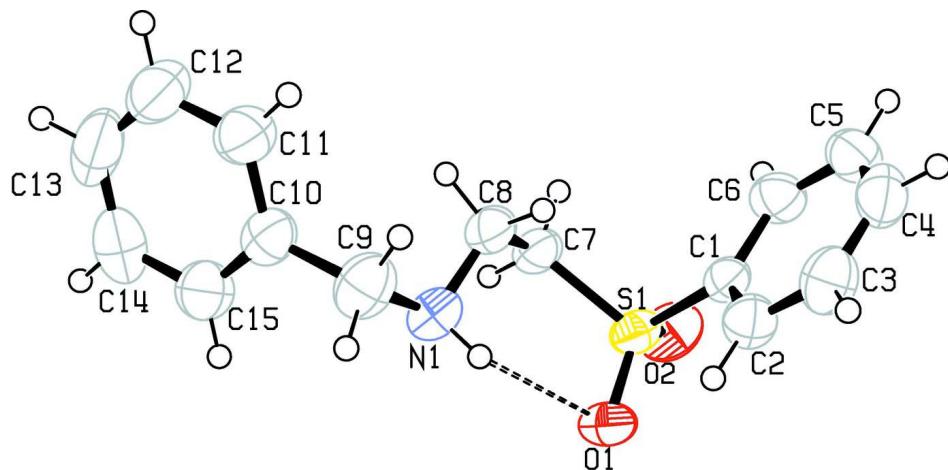
This compound was synthesized from benzylamine and vinylsulfonylbenzene by Dr. J. Gabriel Garcia. It was recrystallized by very slow cooling of an ethanol solution.

Refinement

The coordinates and isotropic displacement parameter of the amine hydrogen atom were refined without constraint. All other H atoms were placed in calculated positions with C(sp^2)—H = 0.930 Å, C(sp^3)—H = 0.970 Å, $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$, and thereafter allowed to ride the attached C atom. The absolute structure is supported by analysis of 2081 Bijvoet pairs, with Flack (1983) parameter $x = 0.07$ (6) and Hooft (Hooft *et al.*, 2008) parameter $y = 0.12$ (2).

Computing details

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); cell refinement: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); data reduction: *XCAD4* (Farrugia, 1999); program(s) used to solve structure: *SIR2002* (Burla *et al.*, 2003); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

**Figure 1**

View of (I) (50% probability displacement ellipsoids)

N*-[2-(Phenylsulfonyl)ethyl]benzylamineCrystal data*

$C_{15}H_{17}NO_2S$
 $M_r = 275.36$
Monoclinic, $P2_1$
Hall symbol: P 2yb
 $a = 5.7428 (11) \text{ \AA}$
 $b = 10.170 (2) \text{ \AA}$
 $c = 12.486 (2) \text{ \AA}$
 $\beta = 102.09 (3)^\circ$
 $V = 713.1 (2) \text{ \AA}^3$
 $Z = 2$

$F(000) = 292$
 $D_x = 1.282 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Cell parameters from 25 reflections
 $\theta = 2.6\text{--}27.5^\circ$
 $\mu = 0.22 \text{ mm}^{-1}$
 $T = 298 \text{ K}$
Prism, colorless
 $0.50 \times 0.45 \times 0.32 \text{ mm}$

Data collection

Enraf–Nonius CAD-4
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 $\theta/2\theta$ scans
Absorption correction: ψ scan
(North *et al.*, 1968)
 $T_{\min} = 0.896$, $T_{\max} = 0.932$
5320 measured reflections

3264 independent reflections
2990 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.021$
 $\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 2.6^\circ$
 $h = -7 \rightarrow 7$
 $k = -13 \rightarrow 13$
 $l = -16 \rightarrow 16$
3 standard reflections every 60 min
intensity decay: 4.0%

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.029$
 $wR(F^2) = 0.086$
 $S = 0.99$
3264 reflections
177 parameters
1 restraint
0 constraints
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
 $w = 1/[\sigma^2(F_o^2) + (0.0562P)^2 + 0.0525P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.15 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.22 \text{ e \AA}^{-3}$

Absolute structure: Flack (1983), 2081 Bijvoet
pairs

Flack parameter: 0.07 (6)

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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.1957 (3)	0.01813 (15)	0.34724 (12)	0.0424 (3)
C2	0.0482 (3)	-0.08893 (17)	0.34717 (13)	0.0533 (4)
H2	-0.0657	-0.0884	0.3903	0.064*
C3	0.0698 (4)	-0.1963 (2)	0.28342 (15)	0.0657 (5)
H3	-0.0305	-0.2682	0.2826	0.079*
C4	0.2404 (4)	-0.1973 (2)	0.22054 (14)	0.0639 (5)
H4	0.254	-0.2699	0.177	0.077*
C5	0.3912 (4)	-0.0917 (2)	0.22157 (15)	0.0656 (5)
H5	0.5078	-0.094	0.1799	0.079*
C6	0.3693 (3)	0.01759 (19)	0.28437 (14)	0.0561 (4)
H6	0.469	0.0897	0.2847	0.067*
C7	0.4077 (2)	0.1536 (2)	0.53880 (12)	0.0522 (3)
H7A	0.5518	0.1629	0.5105	0.063*
H7B	0.3977	0.2293	0.5848	0.063*
C8	0.4311 (3)	0.03164 (18)	0.60956 (14)	0.0560 (4)
H8A	0.4215	-0.0452	0.5628	0.067*
H8B	0.587	0.0316	0.658	0.067*
C9	0.3093 (4)	-0.0788 (2)	0.76111 (17)	0.0691 (5)
H9A	0.374	-0.1558	0.732	0.083*
H9B	0.1651	-0.1049	0.7843	0.083*
C10	0.4876 (3)	-0.02798 (17)	0.85863 (14)	0.0554 (4)
C11	0.7095 (4)	-0.0864 (2)	0.88806 (17)	0.0698 (5)
H11	0.7488	-0.156	0.8469	0.084*
C12	0.8722 (4)	-0.0421 (3)	0.97776 (19)	0.0892 (8)
H12	1.021	-0.0819	0.9964	0.107*
C13	0.8185 (5)	0.0584 (3)	1.03905 (17)	0.0914 (8)
H13	0.9293	0.0868	1.1	0.11*
C14	0.6002 (5)	0.1187 (3)	1.01140 (17)	0.0836 (7)
H14	0.5637	0.1887	1.0529	0.1*
C15	0.4339 (4)	0.0746 (2)	0.92103 (16)	0.0670 (5)
H15	0.2854	0.1147	0.9027	0.08*
N1	0.2511 (3)	0.02105 (18)	0.67505 (12)	0.0610 (4)
H1	0.122 (4)	-0.001 (2)	0.6295 (17)	0.059 (5)*
O1	-0.05605 (19)	0.13963 (16)	0.46634 (10)	0.0646 (4)
O2	0.1828 (3)	0.27332 (14)	0.36372 (13)	0.0725 (4)
S1	0.16006 (6)	0.15714 (4)	0.42668 (3)	0.04876 (11)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0372 (7)	0.0472 (8)	0.0405 (6)	0.0003 (6)	0.0032 (5)	0.0037 (6)
C2	0.0498 (8)	0.0577 (9)	0.0529 (8)	-0.0103 (7)	0.0121 (7)	0.0007 (7)
C3	0.0720 (11)	0.0584 (10)	0.0625 (10)	-0.0135 (9)	0.0046 (9)	-0.0038 (8)
C4	0.0772 (12)	0.0615 (10)	0.0479 (8)	0.0123 (9)	0.0017 (8)	-0.0040 (8)
C5	0.0635 (10)	0.0825 (13)	0.0540 (9)	0.0153 (10)	0.0195 (8)	0.0043 (9)
C6	0.0481 (8)	0.0630 (10)	0.0598 (9)	-0.0047 (7)	0.0170 (7)	0.0065 (8)
C7	0.0383 (6)	0.0595 (8)	0.0546 (7)	0.0006 (9)	0.0004 (5)	-0.0105 (9)
C8	0.0480 (8)	0.0631 (10)	0.0521 (8)	0.0157 (7)	-0.0006 (7)	-0.0062 (7)
C9	0.0672 (11)	0.0664 (11)	0.0689 (11)	-0.0122 (9)	0.0035 (9)	0.0035 (9)
C10	0.0537 (9)	0.0578 (9)	0.0535 (8)	-0.0053 (7)	0.0082 (7)	0.0125 (7)
C11	0.0607 (10)	0.0811 (13)	0.0670 (11)	0.0066 (10)	0.0123 (9)	0.0108 (10)
C12	0.0620 (12)	0.129 (2)	0.0703 (13)	0.0054 (13)	0.0007 (10)	0.0235 (15)
C13	0.0837 (16)	0.133 (2)	0.0498 (10)	-0.0307 (16)	-0.0048 (10)	0.0129 (13)
C14	0.1094 (19)	0.0896 (16)	0.0552 (10)	-0.0127 (13)	0.0251 (11)	-0.0065 (10)
C15	0.0675 (12)	0.0706 (12)	0.0636 (10)	0.0038 (9)	0.0152 (9)	0.0075 (9)
N1	0.0474 (8)	0.0746 (10)	0.0560 (8)	0.0048 (7)	-0.0005 (6)	-0.0003 (7)
O1	0.0386 (5)	0.0823 (10)	0.0720 (7)	0.0089 (6)	0.0094 (5)	-0.0116 (7)
O2	0.0716 (9)	0.0513 (7)	0.0871 (9)	0.0056 (6)	-0.0010 (7)	0.0134 (7)
S1	0.03744 (16)	0.04880 (18)	0.05641 (19)	0.00474 (16)	0.00158 (12)	0.00009 (18)

Geometric parameters (\AA , ^\circ)

C1—C2	1.379 (2)	C9—N1	1.465 (3)
C1—C6	1.392 (2)	C9—C10	1.508 (3)
C1—S1	1.7634 (16)	C9—H9A	0.97
C2—C3	1.372 (3)	C9—H9B	0.97
C2—H2	0.93	C10—C15	1.375 (3)
C3—C4	1.378 (3)	C10—C11	1.384 (3)
C3—H3	0.93	C11—C12	1.376 (3)
C4—C5	1.378 (3)	C11—H11	0.93
C4—H4	0.93	C12—C13	1.351 (4)
C5—C6	1.381 (3)	C12—H12	0.93
C5—H5	0.93	C13—C14	1.373 (4)
C6—H6	0.93	C13—H13	0.93
C7—C8	1.512 (3)	C14—C15	1.391 (3)
C7—S1	1.7748 (15)	C14—H14	0.93
C7—H7A	0.97	C15—H15	0.93
C7—H7B	0.97	N1—H1	0.87 (2)
C8—N1	1.450 (2)	O1—S1	1.4404 (12)
C8—H8A	0.97	O2—S1	1.4402 (15)
C8—H8B	0.97		
C2—C1—C6	120.57 (16)	C10—C9—H9A	109.3
C2—C1—S1	119.32 (12)	N1—C9—H9B	109.3
C6—C1—S1	120.10 (13)	C10—C9—H9B	109.3
C3—C2—C1	119.89 (16)	H9A—C9—H9B	108
C3—C2—H2	120.1	C15—C10—C11	118.58 (19)

C1—C2—H2	120.1	C15—C10—C9	121.49 (18)
C2—C3—C4	119.87 (18)	C11—C10—C9	119.93 (19)
C2—C3—H3	120.1	C12—C11—C10	120.4 (2)
C4—C3—H3	120.1	C12—C11—H11	119.8
C5—C4—C3	120.62 (18)	C10—C11—H11	119.8
C5—C4—H4	119.7	C13—C12—C11	120.8 (2)
C3—C4—H4	119.7	C13—C12—H12	119.6
C4—C5—C6	120.05 (17)	C11—C12—H12	119.6
C4—C5—H5	120	C12—C13—C14	120.0 (2)
C6—C5—H5	120	C12—C13—H13	120
C5—C6—C1	118.99 (16)	C14—C13—H13	120
C5—C6—H6	120.5	C13—C14—C15	119.7 (2)
C1—C6—H6	120.5	C13—C14—H14	120.2
C8—C7—S1	115.86 (13)	C15—C14—H14	120.2
C8—C7—H7A	108.3	C10—C15—C14	120.5 (2)
S1—C7—H7A	108.3	C10—C15—H15	119.8
C8—C7—H7B	108.3	C14—C15—H15	119.8
S1—C7—H7B	108.3	C8—N1—C9	112.65 (15)
H7A—C7—H7B	107.4	C8—N1—H1	105.3 (13)
N1—C8—C7	113.80 (13)	C9—N1—H1	109.5 (14)
N1—C8—H8A	108.8	O2—S1—O1	118.19 (9)
C7—C8—H8A	108.8	O2—S1—C1	108.44 (8)
N1—C8—H8B	108.8	O1—S1—C1	107.70 (8)
C7—C8—H8B	108.8	O2—S1—C7	107.33 (10)
H8A—C8—H8B	107.7	O1—S1—C7	109.39 (8)
N1—C9—C10	111.51 (16)	C1—S1—C7	105.02 (8)
N1—C9—H9A	109.3		
C6—C1—C2—C3	-1.0 (3)	C12—C13—C14—C15	0.9 (4)
S1—C1—C2—C3	178.24 (14)	C11—C10—C15—C14	0.3 (3)
C1—C2—C3—C4	0.7 (3)	C9—C10—C15—C14	179.22 (18)
C2—C3—C4—C5	0.5 (3)	C13—C14—C15—C10	-0.7 (3)
C3—C4—C5—C6	-1.2 (3)	C7—C8—N1—C9	-166.21 (14)
C4—C5—C6—C1	0.9 (3)	C10—C9—N1—C8	79.6 (2)
C2—C1—C6—C5	0.2 (2)	C2—C1—S1—O2	-138.39 (14)
S1—C1—C6—C5	-179.02 (13)	C6—C1—S1—O2	40.86 (14)
S1—C7—C8—N1	-70.29 (17)	C2—C1—S1—O1	-9.40 (15)
N1—C9—C10—C15	64.2 (2)	C6—C1—S1—O1	169.85 (13)
N1—C9—C10—C11	-116.9 (2)	C2—C1—S1—C7	107.12 (14)
C15—C10—C11—C12	-0.1 (3)	C6—C1—S1—C7	-73.64 (14)
C9—C10—C11—C12	-179.06 (19)	C8—C7—S1—O2	-174.19 (12)
C10—C11—C12—C13	0.4 (4)	C8—C7—S1—O1	56.42 (15)
C11—C12—C13—C14	-0.7 (4)	C8—C7—S1—C1	-58.93 (13)

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

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
N1—H1 \cdots O1	0.87 (2)	2.52 (2)	3.071 (2)	121.9 (17)