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3-Aminophenyl naphthalene-1-sulfonate

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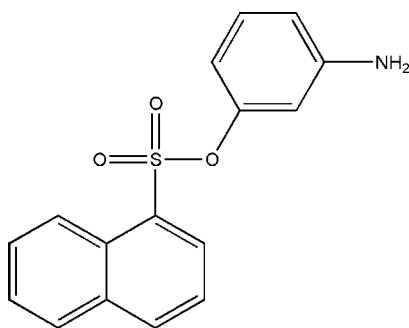
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Key indicators: single-crystal X-ray study; $T = 295$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.055; wR factor = 0.174; data-to-parameter ratio = 26.2.

In the title compound, $\text{C}_{16}\text{H}_{13}\text{NO}_3\text{S}$, the plane of the naphthalene ring system forms a dihedral angle of 64.66 (10)° with the benzene ring. The molecular structure is stabilized by weak intramolecular $\text{C}-\text{H}\cdots\text{O}$ interactions and the crystal packing is stabilized by weak intermolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ interactions and by $\pi-\pi$ stacking interactions of the inversion-related naphthalene units [centroid-centroid distance of 3.7373 (14) Å].

Related literature

For the structures of closely related compounds, see: Manivannan *et al.* (2005a,b); Ramachandran *et al.* (2007); Vennila *et al.* (2008). For applications, see: Spungin *et al.* (1984); Yachi *et al.* (1989).



Experimental

Crystal data

 $\text{C}_{16}\text{H}_{13}\text{NO}_3\text{S}$
 $M_r = 299.33$ Monoclinic, $P2_1/c$
 $a = 8.4558$ (2) Å $b = 8.6712$ (3) Å
 $c = 19.5915$ (6) Å
 $\beta = 100.321$ (2)°
 $V = 1413.24$ (7) Å³
 $Z = 4$ Mo $K\alpha$ radiation
 $\mu = 0.24$ mm⁻¹
 $T = 295$ (2) K
 $0.30 \times 0.25 \times 0.20$ mm

Data collection

Bruker Kappa APEXII diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.932$, $T_{\max} = 0.954$ 19808 measured reflections
4981 independent reflections
3126 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.055$
 $wR(F^2) = 0.174$
 $S = 1.05$
4981 reflections190 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.46$ e Å⁻³
 $\Delta\rho_{\min} = -0.44$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C2}-\text{H2}\cdots\text{O2}$	0.93	2.41	2.829 (3)	107
$\text{C9}-\text{H9}\cdots\text{O3}$	0.93	2.56	3.127 (3)	120
$\text{N1}-\text{H1B}\cdots\text{O3}^i$	0.86	2.43	3.246 (3)	158
$\text{C7}-\text{H7}\cdots\text{O2}^{ii}$	0.93	2.56	3.422 (3)	154

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

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT (Bruker, 2004); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97.

The authors acknowledge the Sophisticated Analytical Instrument Facility, Indian Institute of Technology, Madras, for the data collection.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2179).

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supporting information

Acta Cryst. (2009). E65, o72 [doi:10.1107/S1600536808041032]

3-Aminophenyl naphthalene-1-sulfonate

Jasmine P. Vennila, Helen P. Kavitha, D. John Thiruvadigal, B. R. Venkatraman and V. Manivannan

S1. Comment

Several compounds containing the *para*-toluene sulfonate moiety are used in the fields of biology and industry. The merging of lipids can be monitored using a derivative of *para*-toluene sulfonate (Yachi *et al.*, 1989). This method has been used in studying the membrane fusion during the acrosome reaction (Spungin *et al.*, 1984).

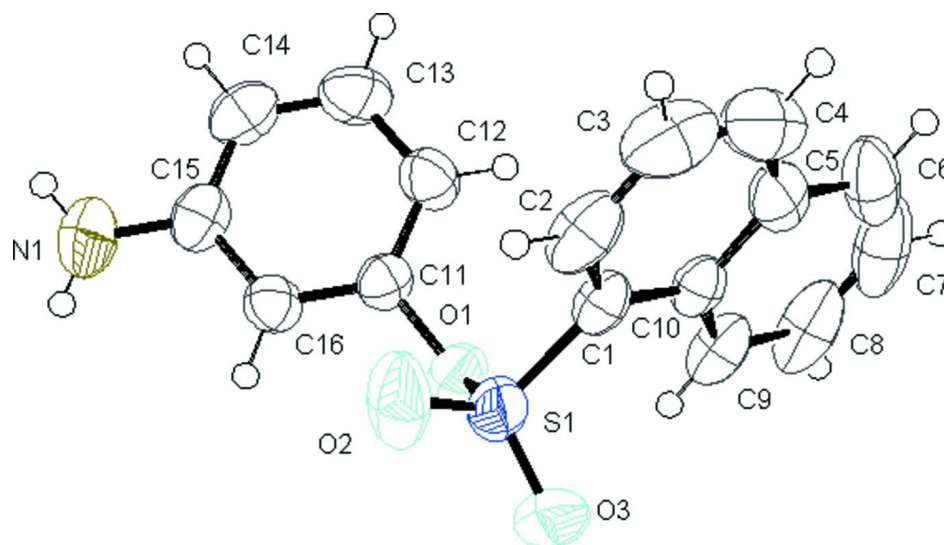
The plane of the benzene ring forms a dihedral angle of 64.66 (10) ° with the naphthalene ring system. The torsion angles O2—S1—C1—C2 and O3—S1—C1—C10 [5.58 (17) ° and 52.09 (16) °, respectively] indicate the *syn* conformation of sulfonyl moiety. The molecular structure is stabilized by weak intramolecular C—H···O interactions and the crystal packing is stabilized by weak intermolecular C—H···O interactions, N—H···O interactions and π - π stacking interactions of the naphthalene fragments related by inversion center

S2. Experimental

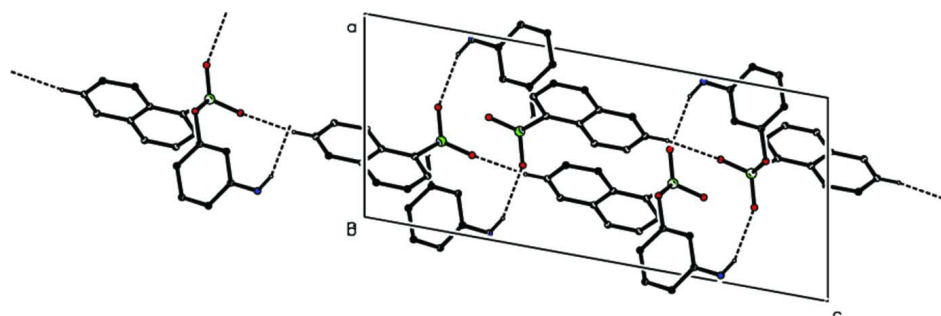
1-Naphthalene sulfonyl chloride (5 mmol) dissolved in acetone (4 ml) was added dropwise to 3-amino phenol (5 mmol) in aqueous NaOH (4 ml, 5%) with constant shaking. The precipitated compound (3 mmol, yield 60%) was recrystlized from ethanol to get diffraction quality brown colored crystals.

S3. Refinement

H atoms were positioned geometrically and refined using riding model with C—H = 0.93 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aromatic C—H and N—H = 0.86 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$ for N—H.

**Figure 1**

The molecular structure of the title compound, with atom labels and 50% probability displacement ellipsoids for non-H atoms.

**Figure 2**

The packing viewed down the *b* axis. Hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bonding have been omitted.

3-Aminophenyl naphthalene-1-sulfonate

Crystal data

$C_{16}H_{13}NO_3S$

$M_r = 299.33$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2_1/c$

$a = 8.4558(2)\ \text{\AA}$

$b = 8.6712(3)\ \text{\AA}$

$c = 19.5915(6)\ \text{\AA}$

$\beta = 100.321(2)^\circ$

$V = 1413.24(7)\ \text{\AA}^3$

$Z = 4$

$F(000) = 624$

$D_x = 1.407\ \text{Mg m}^{-3}$

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

Cell parameters from 4818 reflections

$\theta = 2.2\text{--}25.4^\circ$

$\mu = 0.24\ \text{mm}^{-1}$

$T = 295\ \text{K}$

Block, brown

$0.30 \times 0.25 \times 0.20\ \text{mm}$

Data collection

Bruker Kappa APEX2
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω and φ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
 $T_{\min} = 0.932$, $T_{\max} = 0.954$

19808 measured reflections
4981 independent reflections
3126 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$
 $\theta_{\max} = 32.2^\circ$, $\theta_{\min} = 2.1^\circ$
 $h = -12 \rightarrow 10$
 $k = -12 \rightarrow 11$
 $l = -21 \rightarrow 29$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.055$
 $wR(F^2) = 0.174$
 $S = 1.05$
4981 reflections
190 parameters
0 restraints
Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0763P)^2 + 0.3485P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.46 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.44 \text{ e } \text{\AA}^{-3}$

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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.34222 (19)	0.8581 (2)	0.10826 (8)	0.0475 (4)
C2	0.2573 (3)	0.9698 (3)	0.13547 (12)	0.0688 (6)
H2	0.2543	0.9701	0.1827	0.083*
C3	0.1749 (3)	1.0837 (3)	0.09239 (19)	0.0876 (8)
H3	0.1173	1.1596	0.1109	0.105*
C4	0.1793 (3)	1.0828 (3)	0.02504 (18)	0.0858 (8)
H4	0.1241	1.1593	-0.0029	0.103*
C5	0.2639 (2)	0.9713 (2)	-0.00569 (11)	0.0634 (5)
C6	0.2687 (4)	0.9715 (4)	-0.07762 (13)	0.0910 (9)
H6	0.2122	1.0468	-0.1058	0.109*
C7	0.3515 (4)	0.8673 (4)	-0.10559 (13)	0.1015 (12)
H7	0.3537	0.8711	-0.1529	0.122*
C8	0.4342 (4)	0.7536 (3)	-0.06564 (14)	0.0872 (9)
H8	0.4913	0.6808	-0.0863	0.105*
C9	0.4342 (2)	0.7451 (2)	0.00414 (11)	0.0619 (5)
H9	0.4904	0.6666	0.0303	0.074*

C10	0.34948 (19)	0.85482 (19)	0.03643 (8)	0.0458 (4)
C11	0.21070 (19)	0.52112 (19)	0.14424 (9)	0.0466 (4)
C12	0.0822 (2)	0.5617 (2)	0.09445 (10)	0.0595 (5)
H12	0.0949	0.6176	0.0553	0.071*
C13	-0.0674 (2)	0.5150 (3)	0.10555 (12)	0.0696 (6)
H13	-0.1582	0.5414	0.0734	0.084*
C14	-0.0850 (2)	0.4309 (2)	0.16279 (11)	0.0637 (5)
H14	-0.1874	0.4025	0.1692	0.076*
C15	0.0471 (2)	0.3876 (2)	0.21112 (10)	0.0561 (4)
C16	0.1979 (2)	0.4357 (2)	0.20153 (9)	0.0511 (4)
H16	0.2890	0.4101	0.2336	0.061*
O1	0.36734 (15)	0.56180 (15)	0.13433 (7)	0.0580 (3)
O2	0.3993 (2)	0.7447 (2)	0.23185 (7)	0.0915 (6)
O3	0.60456 (18)	0.7106 (2)	0.15880 (10)	0.0903 (6)
N1	0.0316 (3)	0.3014 (3)	0.26868 (11)	0.0868 (6)
H1A	-0.0621	0.2733	0.2752	0.104*
H1B	0.1156	0.2758	0.2981	0.104*
S1	0.44040 (6)	0.72128 (7)	0.16539 (2)	0.06166 (18)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0464 (8)	0.0490 (9)	0.0497 (8)	-0.0131 (7)	0.0153 (6)	-0.0086 (7)
C2	0.0679 (12)	0.0677 (13)	0.0789 (13)	-0.0188 (10)	0.0346 (10)	-0.0268 (11)
C3	0.0697 (14)	0.0536 (13)	0.144 (3)	0.0011 (10)	0.0312 (15)	-0.0233 (15)
C4	0.0699 (14)	0.0511 (12)	0.130 (2)	-0.0040 (10)	0.0007 (14)	0.0086 (14)
C5	0.0621 (11)	0.0556 (11)	0.0683 (12)	-0.0236 (9)	0.0003 (9)	0.0073 (9)
C6	0.1060 (19)	0.0930 (19)	0.0634 (13)	-0.0531 (16)	-0.0135 (13)	0.0241 (13)
C7	0.136 (3)	0.118 (2)	0.0525 (13)	-0.082 (2)	0.0230 (15)	-0.0135 (15)
C8	0.1037 (19)	0.0958 (19)	0.0737 (14)	-0.0525 (16)	0.0469 (14)	-0.0426 (14)
C9	0.0619 (11)	0.0660 (12)	0.0638 (11)	-0.0206 (9)	0.0276 (9)	-0.0226 (9)
C10	0.0451 (8)	0.0451 (8)	0.0482 (8)	-0.0160 (6)	0.0112 (6)	-0.0062 (6)
C11	0.0461 (8)	0.0416 (8)	0.0523 (9)	-0.0039 (6)	0.0095 (6)	-0.0014 (7)
C12	0.0603 (10)	0.0606 (11)	0.0536 (10)	-0.0068 (9)	-0.0006 (8)	0.0050 (8)
C13	0.0519 (10)	0.0769 (14)	0.0725 (13)	-0.0074 (9)	-0.0091 (9)	-0.0047 (11)
C14	0.0522 (10)	0.0598 (11)	0.0798 (13)	-0.0175 (8)	0.0139 (9)	-0.0150 (10)
C15	0.0643 (10)	0.0438 (9)	0.0640 (11)	-0.0099 (8)	0.0214 (9)	-0.0058 (8)
C16	0.0524 (9)	0.0452 (9)	0.0546 (9)	-0.0002 (7)	0.0068 (7)	0.0035 (7)
O1	0.0501 (6)	0.0556 (7)	0.0704 (8)	-0.0023 (5)	0.0165 (6)	0.0074 (6)
O2	0.1164 (14)	0.1138 (14)	0.0419 (7)	-0.0494 (11)	0.0078 (8)	-0.0014 (8)
O3	0.0456 (8)	0.1086 (14)	0.1099 (14)	-0.0144 (8)	-0.0046 (8)	0.0307 (11)
N1	0.0905 (14)	0.0891 (15)	0.0865 (14)	-0.0159 (11)	0.0316 (11)	0.0238 (11)
S1	0.0536 (3)	0.0762 (4)	0.0524 (3)	-0.0199 (2)	0.00177 (19)	0.0072 (2)

Geometric parameters (Å, °)

C1—C2	1.370 (3)	C9—H9	0.9300
C1—C10	1.420 (2)	C11—C16	1.365 (2)

C1—S1	1.7368 (19)	C11—C12	1.370 (2)
C2—C3	1.401 (4)	C11—O1	1.4173 (19)
C2—H2	0.9300	C12—C13	1.382 (3)
C3—C4	1.327 (4)	C12—H12	0.9300
C3—H3	0.9300	C13—C14	1.368 (3)
C4—C5	1.401 (4)	C13—H13	0.9300
C4—H4	0.9300	C14—C15	1.381 (3)
C5—C6	1.417 (3)	C14—H14	0.9300
C5—C10	1.418 (3)	C15—N1	1.378 (3)
C6—C7	1.321 (5)	C15—C16	1.386 (2)
C6—H6	0.9300	C16—H16	0.9300
C7—C8	1.370 (5)	O1—S1	1.5905 (14)
C7—H7	0.9300	O2—S1	1.4212 (16)
C8—C9	1.369 (3)	O3—S1	1.4199 (16)
C8—H8	0.9300	N1—H1A	0.8600
C9—C10	1.408 (3)	N1—H1B	0.8600
C2—C1—C10	121.24 (18)	C5—C10—C1	117.02 (17)
C2—C1—S1	117.12 (15)	C16—C11—C12	123.72 (16)
C10—C1—S1	121.63 (13)	C16—C11—O1	117.47 (15)
C1—C2—C3	120.2 (2)	C12—C11—O1	118.71 (16)
C1—C2—H2	119.9	C11—C12—C13	116.40 (18)
C3—C2—H2	119.9	C11—C12—H12	121.8
C4—C3—C2	119.7 (2)	C13—C12—H12	121.8
C4—C3—H3	120.2	C14—C13—C12	121.48 (18)
C2—C3—H3	120.2	C14—C13—H13	119.3
C3—C4—C5	122.6 (2)	C12—C13—H13	119.3
C3—C4—H4	118.7	C13—C14—C15	120.92 (18)
C5—C4—H4	118.7	C13—C14—H14	119.5
C4—C5—C6	122.3 (3)	C15—C14—H14	119.5
C4—C5—C10	119.2 (2)	N1—C15—C14	121.64 (19)
C6—C5—C10	118.5 (2)	N1—C15—C16	119.91 (19)
C7—C6—C5	121.6 (3)	C14—C15—C16	118.43 (17)
C7—C6—H6	119.2	C11—C16—C15	119.01 (16)
C5—C6—H6	119.2	C11—C16—H16	120.5
C6—C7—C8	120.6 (2)	C15—C16—H16	120.5
C6—C7—H7	119.7	C11—O1—S1	118.20 (11)
C8—C7—H7	119.7	C15—N1—H1A	120.0
C9—C8—C7	121.2 (3)	C15—N1—H1B	120.0
C9—C8—H8	119.4	H1A—N1—H1B	120.0
C7—C8—H8	119.4	O3—S1—O2	119.75 (11)
C8—C9—C10	120.2 (2)	O3—S1—O1	103.16 (10)
C8—C9—H9	119.9	O2—S1—O1	109.45 (9)
C10—C9—H9	119.9	O3—S1—C1	110.36 (9)
C9—C10—C5	117.90 (18)	O2—S1—C1	108.99 (11)
C9—C10—C1	125.09 (18)	O1—S1—C1	103.85 (7)
C10—C1—C2—C3	0.2 (3)	C16—C11—C12—C13	1.8 (3)

S1—C1—C2—C3	-179.87 (16)	O1—C11—C12—C13	178.00 (17)
C1—C2—C3—C4	0.1 (3)	C11—C12—C13—C14	-0.9 (3)
C2—C3—C4—C5	0.1 (4)	C12—C13—C14—C15	-1.0 (3)
C3—C4—C5—C6	-179.7 (2)	C13—C14—C15—N1	-179.3 (2)
C3—C4—C5—C10	-0.5 (3)	C13—C14—C15—C16	2.1 (3)
C4—C5—C6—C7	178.7 (2)	C12—C11—C16—C15	-0.7 (3)
C10—C5—C6—C7	-0.6 (3)	O1—C11—C16—C15	-176.96 (15)
C5—C6—C7—C8	0.9 (4)	N1—C15—C16—C11	-179.90 (18)
C6—C7—C8—C9	-0.4 (4)	C14—C15—C16—C11	-1.3 (3)
C7—C8—C9—C10	-0.5 (3)	C16—C11—O1—S1	-91.61 (17)
C8—C9—C10—C5	0.8 (3)	C12—C11—O1—S1	91.97 (18)
C8—C9—C10—C1	-179.44 (16)	C11—O1—S1—O3	168.69 (12)
C4—C5—C10—C9	-179.52 (17)	C11—O1—S1—O2	40.15 (16)
C6—C5—C10—C9	-0.3 (2)	C11—O1—S1—C1	-76.12 (13)
C4—C5—C10—C1	0.7 (2)	C2—C1—S1—O3	-127.86 (16)
C6—C5—C10—C1	179.93 (16)	C10—C1—S1—O3	52.09 (16)
C2—C1—C10—C9	179.68 (17)	C2—C1—S1—O2	5.58 (17)
S1—C1—C10—C9	-0.3 (2)	C10—C1—S1—O2	-174.48 (13)
C2—C1—C10—C5	-0.6 (2)	C2—C1—S1—O1	122.16 (14)
S1—C1—C10—C5	179.50 (12)	C10—C1—S1—O1	-57.89 (14)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C2—H2 \cdots O2	0.93	2.41	2.829 (3)	107
C9—H9 \cdots O3	0.93	2.56	3.127 (3)	120
N1—H1B \cdots O3 ⁱ	0.86	2.43	3.246 (3)	158
C7—H7 \cdots O2 ⁱⁱ	0.93	2.56	3.422 (3)	154

Symmetry codes: (i) $-x+1, y-1/2, -z+1/2$; (ii) $x, -y+3/2, z-1/2$.