

3-Fluoro-4-nitrophenyl 4-methylbenzenesulfonate

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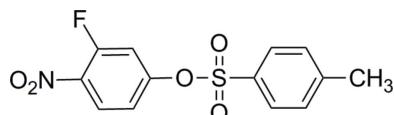
Received 16 December 2010; accepted 17 February 2011

Key indicators: single-crystal X-ray study; $T = 293\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$; R factor = 0.034; wR factor = 0.083; data-to-parameter ratio = 11.8.

In the title compound, $\text{C}_{13}\text{H}_{10}\text{FNO}_5\text{S}$, the dihedral angle between the benzene rings is $47.63(14)^\circ$. In the crystal, $\pi-\pi$ stacking occurs between nearly parallel benzene rings of adjacent molecules, the centroid–centroid distance being $3.7806(16)\text{ \AA}$. Weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonding is also present in the crystal structure.

Related literature

For related compounds and their biological activity, see: Cho *et al.* (2003); Marson *et al.* (2007).



Experimental

Crystal data

$\text{C}_{13}\text{H}_{10}\text{FNO}_5\text{S}$	$V = 1368.57(7)\text{ \AA}^3$
$M_r = 311.28$	$Z = 4$
Orthorhombic, $Pna2_1$	Mo $K\alpha$ radiation
$a = 14.2596(5)\text{ \AA}$	$\mu = 0.27\text{ mm}^{-1}$
$b = 11.4800(3)\text{ \AA}$	$T = 293\text{ K}$
$c = 8.3602(2)\text{ \AA}$	$0.30 \times 0.30 \times 0.20\text{ mm}$

Data collection

Oxford Diffraction Xcalibur Eos diffractometer
Absorption correction: multi-scan (*CrysAlis PRO*; Oxford Diffraction, 2006)
 $T_{\min} = 0.979$, $T_{\max} = 1.0$

10802 measured reflections
2251 independent reflections
1855 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.034$
 $wR(F^2) = 0.083$
 $S = 1.05$
2251 reflections
191 parameters
1 restraint

H-atom parameters constrained
 $\Delta\rho_{\max} = 0.12\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.23\text{ e \AA}^{-3}$
Absolute structure: Flack (1983),
752 Friedel pairs
Flack parameter: $-0.06(9)$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C6—H6 \cdots O2 ⁱ	0.93	2.55	3.224 (4)	129
Symmetry code: (i) $-x + 1, -y, z + \frac{1}{2}$.				

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2006); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *OLEX2*.

We thank the Analytical and Testing Center of Sichuan University for the X-ray measurements.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5126).

References

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Acta Cryst. (2011). E67, o750 [doi:10.1107/S1600536811005903]

3-Fluoro-4-nitrophenyl 4-methylbenzenesulfonate

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Comment

Aryloxyalkanoic acid hydroxyamides are potent inhibitors of histone deacetylase (Marson *et al.*, 2007; Cho *et al.*, 2003). 3-Fluoro-4-nitrophenyl 4-methylbenzenesulfonate is one of the key intermediates to synthesize the aryloxyalkanoic acid hydroxyamides derivatives. We report here its crystal structure. In the title compound (Fig. 1), the dihedral angle between the 3-fluoro-4-nitrophenyl ring and the 4-methylbenzene ring is 47.63 (14)°. In the crystal, intermolecular π – π stacking [centroid–centroid distance = 3.7806 (16) Å] stabilizes the structure (Fig. 2). Weak C—H···O hydrogen bonding is present in the crystal structure (Table 1).

Experimental

To the 3-fluoro-4-nitrophenol (19.10 mmol) in chloroform (20 ml) at 273 K were added pyridine (3.70 ml, 45.84 mmol) dropwise over a period of 20 min and *p*-toluenesulfonyl chloride (22.92 mmol) in small portions. This reaction mixture was stirred at room temperature for 12 h and diluted with dichloromethane and then 10% aqueous HCl. The separated organic layer was washed with 10% aqueous HCl, water and saturated aqueous NaCl; dried over NaSO₄; and concentrated *in vacuo*. The crude 3-fluoro-4-nitrophenyl 4-methylbenzenesulfonate were purified by recrystallization. Crystals suitable for X-ray analysis were obtained by slow evaporation from a solution of ethanol.

Refinement

H atoms were positioned geometrically and refined using a riding model with C—H = 0.93 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aromatic H atoms and C—H = 0.96 Å and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms.

Figures

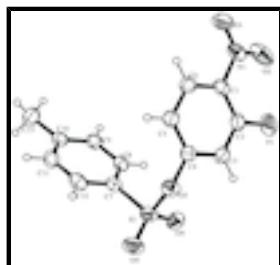


Fig. 1. The molecular structure of the title compound, with displacement ellipsoids drawn at the 30% probability level.

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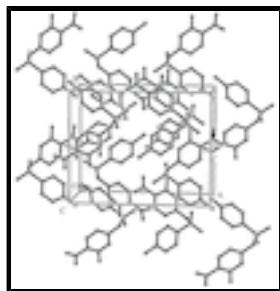


Fig. 2. The cell packing of the title compound.

3-Fluoro-4-nitrophenyl 4-methylbenzenesulfonate

Crystal data

C ₁₃ H ₁₀ FNO ₅ S	<i>F</i> (000) = 640
<i>M_r</i> = 311.28	<i>D_x</i> = 1.511 Mg m ⁻³
Orthorhombic, <i>Pna</i> 2 ₁	Mo <i>Kα</i> radiation, λ = 0.7107 Å
Hall symbol: P 2c -2n	Cell parameters from 4234 reflections
<i>a</i> = 14.2596 (5) Å	θ = 2.9–29.1°
<i>b</i> = 11.4800 (3) Å	μ = 0.27 mm ⁻¹
<i>c</i> = 8.3602 (2) Å	<i>T</i> = 293 K
<i>V</i> = 1368.57 (7) Å ³	Block, colorless
<i>Z</i> = 4	0.30 × 0.30 × 0.20 mm

Data collection

Oxford Diffraction Xcalibur Eos diffractometer	2251 independent reflections
Radiation source: fine-focus sealed tube graphite	1855 reflections with $I > 2\sigma(I)$
Detector resolution: 16.0874 pixels mm ⁻¹	R_{int} = 0.025
ω scans	$\theta_{\text{max}} = 26.4^\circ$, $\theta_{\text{min}} = 2.9^\circ$
Absorption correction: multi-scan (<i>CrysAlis PRO</i> ; Oxford Diffraction, 2006)	$h = -17 \rightarrow 17$
$T_{\text{min}} = 0.979$, $T_{\text{max}} = 1.0$	$k = -14 \rightarrow 14$
10802 measured reflections	$l = -7 \rightarrow 10$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.034$	H-atom parameters constrained
$wR(F^2) = 0.083$	$w = 1/[\sigma^2(F_o^2) + (0.039P)^2 + 0.1799P]$
$S = 1.05$	where $P = (F_o^2 + 2F_c^2)/3$
2251 reflections	$(\Delta/\sigma)_{\text{max}} = 0.025$
	$\Delta\rho_{\text{max}} = 0.12 \text{ e \AA}^{-3}$

191 parameters $\Delta\rho_{\min} = -0.23 \text{ e } \text{\AA}^{-3}$
 1 restraint Absolute structure: Flack (1983), 752 Friedel pairs
 Primary atom site location: structure-invariant direct Flack parameter: -0.06 (9)
 methods

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>	<i>U</i> _{iso} */* <i>U</i> _{eq}
S1	0.16636 (4)	0.27076 (5)	0.58998 (10)	0.05567 (19)
F1	0.34171 (15)	-0.14722 (14)	0.6292 (3)	0.1007 (7)
O1	0.56870 (17)	-0.0441 (2)	0.8899 (4)	0.0906 (8)
O2	0.51366 (19)	-0.1649 (2)	0.7187 (3)	0.1041 (9)
O3	0.18775 (13)	0.20401 (17)	0.7566 (2)	0.0574 (5)
O4	0.16341 (14)	0.18606 (19)	0.4665 (3)	0.0675 (6)
O5	0.08578 (12)	0.33777 (18)	0.6260 (3)	0.0788 (7)
N1	0.5080 (2)	-0.0778 (2)	0.7989 (3)	0.0675 (7)
C1	0.4217 (2)	-0.0077 (2)	0.7863 (4)	0.0523 (7)
C2	0.3442 (2)	-0.0442 (2)	0.7026 (4)	0.0601 (8)
C3	0.2660 (2)	0.0248 (2)	0.6906 (4)	0.0580 (7)
H3	0.2136	-0.0003	0.6341	0.070*
C4	0.26682 (18)	0.1317 (2)	0.7636 (3)	0.0474 (6)
C5	0.3427 (2)	0.1698 (2)	0.8491 (4)	0.0548 (7)
H5	0.3417	0.2424	0.8987	0.066*
C6	0.4204 (2)	0.0995 (2)	0.8609 (4)	0.0566 (7)
H6	0.4722	0.1243	0.9192	0.068*
C7	0.26414 (17)	0.3597 (2)	0.5642 (3)	0.0468 (6)
C8	0.33831 (18)	0.3209 (2)	0.4704 (4)	0.0532 (7)
H8	0.3359	0.2483	0.4213	0.064*
C9	0.41530 (18)	0.3917 (2)	0.4512 (4)	0.0538 (7)
H9	0.4659	0.3650	0.3914	0.065*
C10	0.41940 (19)	0.5008 (2)	0.5179 (3)	0.0519 (7)
C11	0.34570 (19)	0.5373 (2)	0.6128 (4)	0.0584 (7)
H11	0.3483	0.6103	0.6609	0.070*
C12	0.26858 (19)	0.4679 (2)	0.6375 (3)	0.0551 (7)
H12	0.2198	0.4931	0.7028	0.066*
C13	0.5028 (2)	0.5786 (3)	0.4867 (5)	0.0752 (9)
H13A	0.4860	0.6370	0.4096	0.113*

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H13B	0.5537	0.5327	0.4460	0.113*
H13C	0.5215	0.6155	0.5847	0.113*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0420 (3)	0.0668 (4)	0.0582 (4)	0.0045 (3)	-0.0005 (4)	0.0083 (4)
F1	0.1299 (17)	0.0508 (9)	0.1214 (18)	0.0063 (10)	-0.0321 (16)	-0.0197 (11)
O1	0.0640 (15)	0.0999 (18)	0.108 (2)	0.0104 (14)	-0.0161 (15)	0.0259 (17)
O2	0.135 (2)	0.0934 (17)	0.0836 (18)	0.0580 (16)	-0.0025 (17)	-0.0028 (15)
O3	0.0489 (11)	0.0689 (12)	0.0543 (12)	0.0047 (9)	0.0104 (10)	0.0087 (10)
O4	0.0635 (14)	0.0765 (13)	0.0625 (13)	-0.0099 (10)	-0.0104 (11)	-0.0038 (11)
O5	0.0452 (11)	0.0901 (14)	0.1011 (19)	0.0158 (9)	0.0081 (13)	0.0152 (14)
N1	0.0737 (19)	0.0709 (17)	0.0579 (17)	0.0151 (14)	0.0075 (15)	0.0231 (15)
C1	0.0533 (18)	0.0547 (16)	0.0487 (17)	0.0054 (12)	0.0020 (14)	0.0132 (14)
C2	0.078 (2)	0.0415 (14)	0.0605 (19)	-0.0031 (14)	-0.0059 (17)	0.0043 (14)
C3	0.0585 (17)	0.0529 (15)	0.0627 (19)	-0.0129 (13)	-0.0115 (15)	0.0035 (14)
C4	0.0457 (14)	0.0534 (14)	0.0430 (15)	-0.0017 (11)	0.0040 (13)	0.0085 (12)
C5	0.0580 (17)	0.0520 (15)	0.0543 (17)	-0.0027 (13)	-0.0022 (15)	0.0005 (13)
C6	0.0557 (18)	0.0606 (17)	0.0534 (18)	-0.0076 (14)	-0.0060 (15)	0.0069 (14)
C7	0.0431 (13)	0.0520 (13)	0.0453 (17)	0.0104 (10)	0.0012 (13)	0.0046 (13)
C8	0.0524 (16)	0.0467 (13)	0.0604 (17)	0.0088 (12)	0.0042 (15)	-0.0067 (13)
C9	0.0453 (15)	0.0563 (16)	0.0597 (18)	0.0090 (12)	0.0078 (14)	-0.0005 (14)
C10	0.0506 (16)	0.0555 (16)	0.0497 (17)	0.0051 (12)	-0.0056 (13)	0.0021 (12)
C11	0.0692 (18)	0.0491 (13)	0.0568 (18)	0.0042 (12)	-0.0026 (18)	-0.0081 (14)
C12	0.0601 (17)	0.0582 (15)	0.0471 (17)	0.0176 (12)	0.0093 (14)	-0.0015 (13)
C13	0.069 (2)	0.0714 (19)	0.085 (2)	-0.0119 (15)	0.001 (2)	-0.0069 (18)

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

S1—O3	1.619 (2)	C5—C6	1.374 (4)
S1—O4	1.419 (2)	C6—H6	0.9300
S1—O5	1.4152 (19)	C7—C8	1.390 (4)
S1—C7	1.741 (3)	C7—C12	1.386 (3)
F1—C2	1.332 (3)	C8—H8	0.9300
O1—N1	1.216 (4)	C8—C9	1.375 (3)
O2—N1	1.206 (3)	C9—H9	0.9300
O3—C4	1.401 (3)	C9—C10	1.373 (4)
N1—C1	1.475 (4)	C10—C11	1.382 (4)
C1—C2	1.374 (4)	C10—C13	1.509 (4)
C1—C6	1.380 (4)	C11—H11	0.9300
C2—C3	1.372 (4)	C11—C12	1.374 (4)
C3—H3	0.9300	C12—H12	0.9300
C3—C4	1.371 (4)	C13—H13A	0.9600
C4—C5	1.369 (4)	C13—H13B	0.9600
C5—H5	0.9300	C13—H13C	0.9600
F1—C2—C1	121.8 (3)	C6—C1—N1	117.8 (3)
F1—C2—C3	117.2 (3)	C6—C5—H5	120.4

O1—N1—C1	117.7 (3)	C7—C8—H8	120.5
O2—N1—O1	124.3 (3)	C7—C12—H12	120.3
O2—N1—C1	118.0 (3)	C8—C7—S1	119.45 (19)
O3—S1—C7	103.48 (11)	C8—C9—H9	119.1
O4—S1—O3	107.89 (11)	C9—C8—C7	119.0 (2)
O4—S1—C7	109.59 (13)	C9—C8—H8	120.5
O5—S1—O3	103.14 (14)	C9—C10—C11	118.5 (2)
O5—S1—O4	120.23 (14)	C9—C10—C13	120.2 (3)
O5—S1—C7	110.97 (11)	C10—C9—C8	121.7 (2)
C1—C6—H6	119.9	C10—C9—H9	119.1
C2—C1—N1	122.7 (3)	C10—C11—H11	119.4
C2—C1—C6	119.5 (3)	C10—C13—H13A	109.5
C2—C3—H3	120.7	C10—C13—H13B	109.5
C3—C2—C1	121.0 (3)	C10—C13—H13C	109.5
C3—C4—O3	120.3 (2)	C11—C10—C13	121.3 (3)
C4—O3—S1	117.93 (16)	C11—C12—C7	119.3 (2)
C4—C3—C2	118.6 (3)	C11—C12—H12	120.3
C4—C3—H3	120.7	C12—C7—S1	120.46 (19)
C4—C5—H5	120.4	C12—C7—C8	120.1 (2)
C4—C5—C6	119.1 (3)	C12—C11—C10	121.3 (2)
C5—C4—O3	118.0 (2)	C12—C11—H11	119.4
C5—C4—C3	121.7 (2)	H13A—C13—H13B	109.5
C5—C6—C1	120.2 (3)	H13A—C13—H13C	109.5
C5—C6—H6	119.9	H13B—C13—H13C	109.5

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C6—H6···O2 ⁱ	0.93	2.55	3.224 (4)	129

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

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Fig. 1

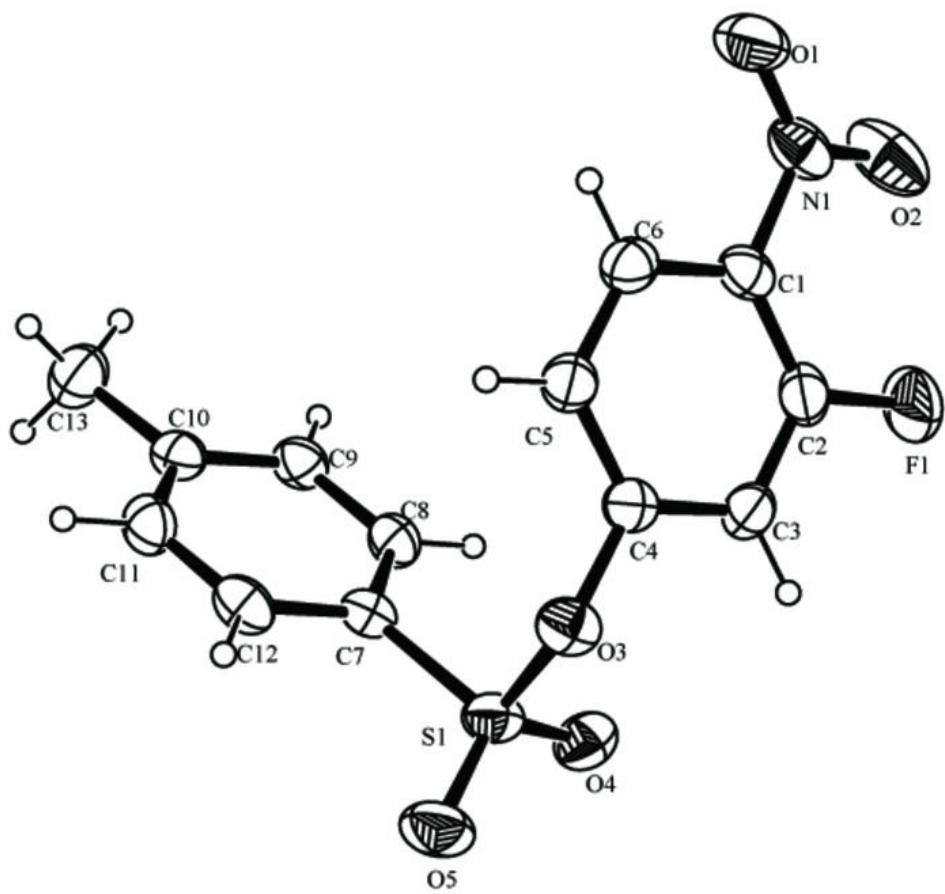


Fig. 2

