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N-(3-Hydroxyphenyl)-*p*-toluenesulfonamide

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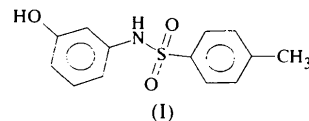
Abstract

In the title compound, C₁₃H₁₃NO₃S, the dihedral angle between the toluene and phenol moieties is 69.48 (5)°. The molecules form columns linked by N—H···O and O—H···O hydrogen bonds.

Comment

Molecular recognition is central to the understanding of biological activity and enzymatic systems (Goswami & Ghosh, 1997). The title compound, (I), contains a potentially active sulfonamide moiety. Sulfonamides have been extensively used against many common gram-positive bacterial infections like pneumonia, meningitis, dysentery and urinary-tract infection (Kar, 1993). Hydrogen bonding is expected to play an important role

in governing the crystal structure of the title compound (Aoyama *et al.*, 1995). The X-ray structure determination was carried out in order to elucidate the molecular conformation and hydrogen bonding.



Bond lengths agree with other reported values (Allen *et al.*, 1987). The sum of the bond angles around the N1 atom [349 (1)°] indicates that it is in a pyramidal configuration. The dihedral angle between the toluene and phenol moieties is 69.48 (5)°. Chains of molecules along **b** are linked by O—H···O hydrogen bonds, and glide-related chains are interlinked by N—H···O hydrogen bonds (see Table 2 for details). These two chains form a column-like structure along the *b* axis, with the phenol groups occupying a plane and the toluene groups projecting outwards from it on either side. The toluene groups of neighbouring columns are arranged close to each other, but are not stacked to allow any π – π interactions.

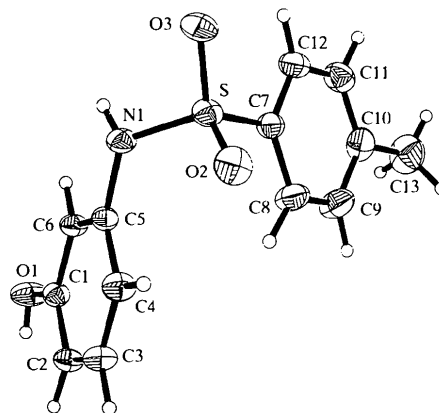


Fig. 1. The structure and atom-numbering scheme of the title compound, shown with 50% probability displacement ellipsoids.

Experimental

For the preparation of the title compound, *p*-toluenesulfonyl chloride (875 mg, 4.5 mmol) was added to a solution of *m*-aminophenol (500 mg, 4.5 mmol) and dry pyridine (0.5 ml) in dry CH₂Cl₂ solvent at 283 K. The reaction mixture was stirred for 15 min at this temperature and for a further 1 h at room temperature. After the usual work-up, the desired compound was obtained in 80% yield (Kurita, 1974). Single crystals were grown by slow evaporation of a 4:1 dichloro-methane–methanol solution of the compound.

Crystal data

C₁₃H₁₃NO₃S
M_r = 263.30

Mo *K*α radiation
λ = 0.71073 Å

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Monoclinic
C2/c
a = 23.9096 (7) Å
b = 9.5192 (3) Å
c = 11.9629 (3) Å
β = 112.136 (1)°
V = 2522.07 (13) Å³
Z = 8
D_x = 1.387 Mg m⁻³
D_m not measured

Cell parameters from 6375 reflections
θ = 1.5–33.0°
μ = 0.256 mm⁻¹
T = 293 (2) K
Block
0.52 × 0.36 × 0.30 mm
Colourless

Data collection: *SMART* (Siemens, 1996b). Cell refinement: *SAINT* (Siemens, 1996a). Data reduction: *SAINT*. Program(s) used to solve structure: *SHELXTL* (Sheldrick, 1996b). Program(s) used to refine structure: *SHELXTL*. Molecular graphics: *SHELXTL*. Software used to prepare material for publication: *SHELXTL* and *PARST* (Nardelli, 1995).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: HA1219). Services for accessing these data are described at the back of the journal.

Data collection

Siemens SMART CCD area-detector diffractometer
ω scans
Absorption correction: empirical (*SADABS*); Sheldrick, 1996a)
T_{min} = 0.871, T_{max} = 0.955
8060 measured reflections

2898 independent reflections
2474 reflections with I > 2σ(I)
R_{int} = 0.016
θ_{max} = 27.5°
h = -30 → 28
k = 0 → 12
l = 0 → 15

Refinement

Refinement on F²
R[F² > 2σ(F²)] = 0.036
wR(F²) = 0.102
S = 1.058
2897 reflections
215 parameters
All H atoms refined
w = 1/[σ²(F_o²) + (0.0496P)² + 1.4298P]
where P = (F_o² + 2F_c²)/3

(Δ/σ)_{max} < 0.001
Δρ_{max} = 0.218 e Å⁻³
Δρ_{min} = -0.280 e Å⁻³
Extinction correction: none
Scattering factors from *International Tables for Crystallography* (Vol. C)

Table 1. Selected geometric parameters (Å, °)

S—O2	1.4259 (13)	S—C7	1.760 (2)
S—O3	1.4347 (12)	O1—C1	1.371 (2)
S—N1	1.6214 (13)	N1—C5	1.432 (2)
O2—S—O3	119.55 (8)	O3—S—C7	107.62 (8)
O2—S—N1	108.65 (8)	N1—S—C7	107.96 (7)
O3—S—N1	104.59 (7)	C5—N1—S	123.62 (11)
O2—S—C7	107.97 (8)		
C4—C5—N1—S	52.0 (2)	N1—S—C7—C8	-74.3 (2)
C5—N1—S—C7	64.19 (14)		

Table 2. Hydrogen-bonding geometry (Å, °)

D—H...A	D—H	H...A	D...A	D—H...A
O1—H1O1...O3 ⁱ	0.79 (3)	1.95 (3)	2.727 (2)	171 (2)
N1—H1N1...O1 ⁱⁱ	0.85 (2)	2.14 (2)	2.983 (2)	174 (2)

Symmetry codes: (i) x, 1 + y, z; (ii) ½ - x, y - ½, ½ - z.

The data collection covered over a hemisphere of reciprocal space by a combination of three sets of exposures; each set had a different φ angle (0, 88 and 180°) for the crystal and each exposure of 30 s covered 0.3° in ω. The crystal-to-detector distance was 4 cm and the detector swing angle was -35°. Coverage of the unique set is over 99% complete. Crystal decay, monitored by repeating 30 initial frames at the end of data collection and analysing the duplicate reflections, was found to be negligible. The structure was solved by direct methods and refined by full-matrix least-squares techniques. All H atoms were located from a difference Fourier map and refined isotropically.

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Cyclodecanol

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Abstract

Cyclodecanol crystals, C₁₀H₂₀O, grown by slow sublimation, occur in the monoclinic system in space group