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#### **Key indicators**

Single-crystal X-ray study T = 133 K Mean  $\sigma$ (C–C) = 0.002 Å R factor = 0.037 wR factor = 0.105 Data-to-parameter ratio = 23.8

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# 2,3-Dimethyl-2-(p-tolylsulfonyl)butyric acid

The title compound,  $C_{13}H_{18}O_4S$ , displays a  $C_{Ph}-S-C-C_{acid}$  torsion angle of 33.23 (8)°; it has been suggested that this angle should be approximately 60° for the molecule to act as an artificial sweetener. Molecules are connected by classical carboxylic acid hydrogen-bond dimers.

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# Comment

Arylsulfonylalkanoic acids have been synthesized as potential sweeteners (Polański *et al.*, 1997; Doleżych *et al.*, 1999; Ratajczak & Polański, 1991). Crystal structures have been reported for the compounds 2,3-dimethyl-2-phenylsulfonylbutyric acid [which differs from the title compound, (I), an effective artificial sweetener, only in the missing *p*-methyl group (Polański *et al.*, 1997)] and ethyl *p*-nitrophenylsulfonylacetate (Anulewicz *et al.*, 1990).



The structure of (I) is shown in Fig. 1. The molecule has a chiral centre at C1 (crystallographic numbering), which in Fig. 1 has the *R* configuration, but the overall structure is a racemate. Molecular dimensions may be regarded as normal. Atom O2 is approximately coplanar with the ring. The C11–S-C1-C3 torsion angle is 33.23 (8)°; Polański *et al.* (1997) suggested that this C<sub>Ph</sub>- $S-C-C_{acid}$  torsion angle should be approximately 60° for the artificial sweetener to be effective.

The packing involves the classical carboxylic acid hydrogenbond dimer over an inversion centre (Fig. 2). There are no unusually short  $C-H\cdots O$  or  $C-H\cdots \pi$  contacts.



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## **Experimental**

Compound (I) was prepared by methylation of 2-methylpropyl-*p*toluenesulfone (Yi *et al.*, 2004) according to the literature method of Ratajczak & Polański (1991) and subsequent carboxylation according to Cram & Ratajczak, (1971). Single crystals were obtained by slow evaporation in air of a solution in methanol (m.p. 417–419 K). Consistent analytical data were obtained.

Z = 8

 $D_x = 1.349 \text{ Mg m}^{-3}$ Mo K\alpha radiation  $\mu = 0.25 \text{ mm}^{-1}$ T = 133 (2) K Block, colourless 0.27 \times 0.24 \times 0.22 mm

#### Crystal data

M = 270.22
$M_r = 270.55$
Orthorhombic, Pbca
a = 10.6198 (11)  Å
b = 12.4497 (12) Å
c = 20.141 (2)  Å
$V = 2662.9 (5) \text{ Å}^3$

#### Data collection

Bruker SMART 1000 CCD areadetector diffractometer4064 independent reflections $\varphi$  and  $\omega$  scans3280 reflections with  $I > 2\sigma(I)$  $\varphi$  and  $\omega$  scans $R_{int} = 0.103$ Absorption correction: none $\theta_{max} = 30.5^{\circ}$ 36385 measured reflections $\theta_{max} = 30.5^{\circ}$ 

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0595P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.037$	+ 0.5795P]
$wR(F^2) = 0.105$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} = 0.001$
4064 reflections	$\Delta \rho_{\rm max} = 0.51 \ {\rm e} \ {\rm \AA}^{-3}$
171 parameters	$\Delta \rho_{\rm min} = -0.30 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

### Table 1

Selected geometric parameters (Å, °).

S-01	1.4427 (9)	S-C11	1.7661 (12)	
S-O2	1.4443 (9)	S-C1	1.8557 (11)	
O1-S-O2	118.86 (5)	O1-S-C1	107.31 (5)	
O1-S-C11	108.59 (6)	O2-S-C1	108.77 (5)	
O2-S-C11	108.19 (5)	C11-S-C1	104.16 (5)	
C11-S-C1-C3	33.23 (8)	O2-S-C11-C12	18.67 (11)	

#### Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O4-H04···O3 <sup>i</sup>	0.90 (2)	1.70 (2)	2.6051 (12)	178 (2)

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





Packing of the title compound, viewed parallel to the a axis. Hydrogen bonds are shown as dashed lines. H atoms have been omitted, except for the acidic H atom.

The acidic H atom was refined freely. Methyl H atoms were identified in difference syntheses, idealized and then refined using a rigid methyl group (C–H = 0.98 Å and H–C–H = 109.5°) allowed to rotate but not tip. Other H atoms were included using a riding model, with C–H = 0.95 (aromatic) and 1.00 Å (*sp*<sup>3</sup>-CH).  $U_{iso}(H)$  values were fixed at 1.2 $U_{eq}(C)$  of the parent C atom.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1994); software used to prepare material for publication: *SHELXL97*.

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