

(2*R*,4*R*)-(4-Iodo-5,5-dimethyl-2-hexylsulfonyl)benzene**Guido J. Reiß^{a*} and Marek Masnyk^b**^aInstitut für Anorganische Chemie und Strukturchemie, Heinrich-Heine-Universität Düsseldorf, Universitätsstraße, D-40225 Düsseldorf, Germany, and ^bInstitute of Organic Chemistry, Polish Academy of Sciences, Kasprzaka 44, 01-224 Warsaw, Poland

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Key indicators

Single-crystal X-ray study

T = 293 KMean $\sigma(\text{C}-\text{C}) = 0.006 \text{ \AA}$ *R* factor = 0.035*wR* factor = 0.066

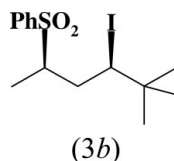
Data-to-parameter ratio = 15.4

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

Structure refinement of X-ray diffraction data in the polar space group $P2_1$ and the assignment of the absolute structure identified the crystal of the title compound, $\text{C}_{14}\text{H}_{21}\text{IO}_2\text{S}$, under investigation as the $2R,4R$ isomer. The packing is dominated by hydrophobic layers in the $a0c$ and $a\frac{1}{2}c$ planes and hydrophilic stacks along [100].

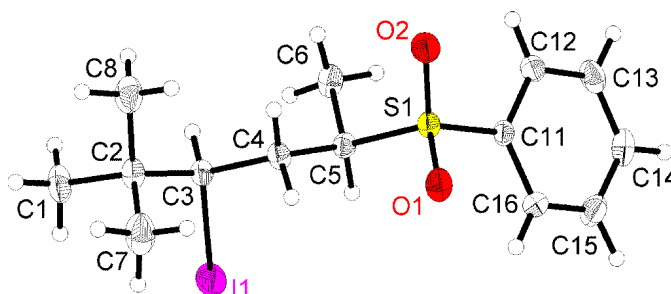
Comment

As we have reported (Reiß & Masnyk, 2001), (1-iodoethanesulfonyl)benzene reacts with 3,3-dimethylbut-1-ene under standard conditions (Jankowski *et al.*, 1995; Masnyk, 1991) to yield a chromatographically separable mixture of the RS,SR and the RR,SS diastereomers of (4-iodo-5,5-dimethyl-2-hexylsulfonyl)benzene. In this context, the crystal structure determination of the title compound, (3*b*), was undertaken to relate the molecular structure to the ^1H and ^{13}C NMR data. This should allow us to predict the stereochemistry of analogous compounds from NMR data.



Crystallization of the title compound from hexane yielded needle-shaped colourless crystals. The compound crystallizes in the polar space group $P2_1$. The crystal is composed of molecules having a $2R,4R$ -configuration at both chiral atoms C3 and C5 (Fig. 1).

All bond lengths, bond angles and torsion angles are as expected, including a significant distortion from tetrahedral geometry at the S atom caused by the $\text{S}=\text{O}$ double bonds (Table 1). Deviations of the ideal bond angles also occur at atoms C3 and C4 since the bulky substituents *tert*-butyl and

**Figure 1**

The molecular structure of the title compound. Displacement ellipsoids are shown at the 50% probability level and H atoms are drawn with an arbitrary radius.

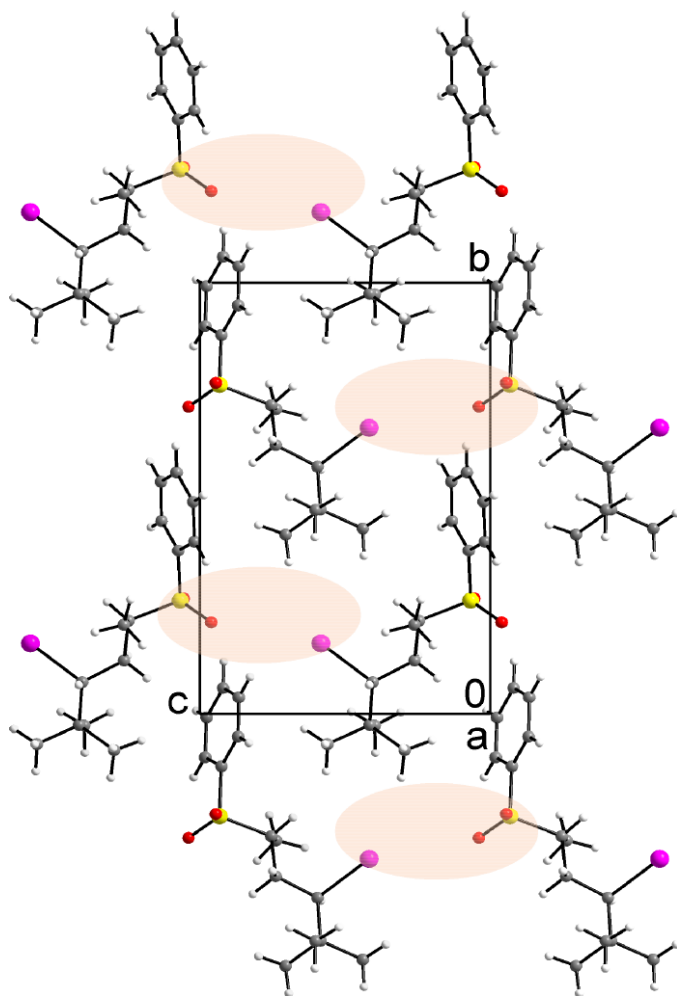


Figure 2
The packing of the title compound viewed down [100].

iodine cause a stretching of the aliphatic section of the molecule. A reason for the formation of enantiopure crystals from a racemic solution of the title compound may be that the molecules do have an excellent fit within the bc plane (Fig. 2). This arrangement shows separated hydrophobic regions in the $a0c$ and $a\frac{1}{2}c$ planes as well as hydrophilic regions perpendicular to the bc plane (shaded ellipses).

Experimental

A mixture of (1-iodoethanesulfonyl)benzene (600 mg), 3,3-dimethylbut-1-ene (1.0 ml), benzene (1.5 ml) and benzoyl peroxide (40 mg) was heated for 6 h at 373 K in a sealed tube. The reaction mixture was then chromatographed on silica gel (hexane–ethyl acetate 93:7) yielding 280 mg of the *anti* isomer (2S4R and 2R4S) and 350 mg of the *syn* isomer (2R4R and 2S4S) (m.p. = 369–371 K). ^1H NMR (500 MHz, CDCl_3): δ = 1.08 (s, 9H), 1.25 (d, J = 6.8 Hz, 3H), 1.76 (ddd, J = 14.5, 11.2, 2.1 Hz, 1H), 2.35 (ddd, J = 14.5, 12.3, 2.6 Hz, 1H), 3.43 (dq, J = 11.2, 6.8, 2.6 Hz, 1H), 3.88 (dd, J = 12.3, 2.1 Hz, 1H), 7.57–7.61 (m, 2H), 7.68 (tt, J = 7.4, 1.2 Hz, 1H), 7.89–7.92 (m, 2H). ^{13}C NMR (125 MHz, CDCl_3): δ = 11.63, 28.49, 34.44, 35.90, 51.85, 61.00, 128.83, 129.17, 133.77, 137.29.

Crystal data

$\text{C}_{14}\text{H}_{21}\text{IO}_2\text{S}$
 M_r = 380.27
Monoclinic, $P2_1$
 a = 6.1329 (8) Å
 b = 14.3524 (13) Å
 c = 9.4381 (10) Å
 β = 103.919 (9)°
 V = 806.36 (15) Å³
 Z = 2

D_x = 1.566 Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 38 reflections
 θ = 7.0–12.5°
 μ = 2.11 mm⁻¹
 T = 293 (2) K
Needle, colourless
1.00 × 0.50 × 0.30 mm

Data collection

Siemens P4 diffractometer
 ω scans
Absorption correction: numerical (Stoe & Cie, 1996)
 T_{\min} = 0.459, T_{\max} = 0.623
3964 measured reflections
3189 independent reflections
2659 reflections with $I > 2\sigma(I)$

R_{int} = 0.012
 θ_{max} = 32.5°
 h = -1 → 9
 k = -1 → 21
 l = -14 → 14
3 standard reflections
every 100 reflections
intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)]$ = 0.035
 $wR(F^2)$ = 0.066
 S = 1.07
3189 reflections
207 parameters
H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.01P)^2 + 0.6P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}}$ = 0.001
 $\Delta\rho_{\text{max}}$ = 0.93 e Å⁻³
 $\Delta\rho_{\text{min}}$ = -0.73 e Å⁻³
Extinction correction: SHELXL97
Extinction coefficient: 0.0178 (10)
Absolute structure: Flack (1983)
Flack parameter = 0.01 (2)

Table 1

Selected geometric parameters (Å, °).

| | | | |
|-------------|-------------|--------------|-------------|
| I1–C3 | 2.191 (4) | C2–C7 | 1.518 (6) |
| S1–O2 | 1.428 (3) | C2–C3 | 1.551 (5) |
| S1–O1 | 1.440 (3) | C3–C4 | 1.521 (5) |
| S1–C11 | 1.771 (4) | C4–C5 | 1.531 (5) |
| S1–C5 | 1.807 (3) | C5–C6 | 1.522 (5) |
| C2–C8 | 1.525 (7) | | |
| O2–S1–O1 | 119.43 (19) | C11–S1–C5 | 104.03 (17) |
| O2–S1–C11 | 108.65 (17) | C4–C3–C2 | 115.5 (3) |
| O1–S1–C11 | 107.54 (18) | C4–C3–I1 | 108.7 (2) |
| O2–S1–C5 | 108.44 (18) | C2–C3–I1 | 111.9 (2) |
| O1–S1–C5 | 107.69 (17) | C3–C4–C5 | 114.6 (3) |
| C1–C2–C3–C4 | 175.4 (4) | C3–C4–C5–S1 | 179.4 (3) |
| C2–C3–C4–C5 | -172.7 (3) | C11–S1–C5–C4 | 168.3 (2) |

The needle-shaped crystal was mounted using a glass capillary aligned along the φ axis. This special orientation leads to maximum and minimum transmission factors – used for absorption correction – which are different from the theoretical values calculated from the crystal dimensions and the linear absorption coefficient. Atomic coordinates for all H atoms belonging to the CH₂ and the CH groups were included in the final stages of refinement, using a riding model. H atoms belonging to methyl groups were included in the refinement riding on their attached C atom and allowed to rotate about the C–C bond with one common U_{iso} value refined for each group. The absolute configuration assignment has been checked following some generally accepted guidelines (Flack & Bernardinelli, 2000). A comparable refinement of the inverted structure yielded a Flack parameter – calculated by a ‘hole in one’ method [x = 0.36 (2)] – indicating the wrong absolute structure assignment and significant higher R values ($wR2$ = 0.0731 and $R1$ = 0.0377).

Data collection: *XSCANS* (Siemens, 1996); cell refinement: *XSCANS*; data reduction: *XSCANS*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *DIAMOND* (Brandenburg, 1998); software used to prepare material for publication: *SHELXL97*.

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