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(+)-Camphorsulfonylimine

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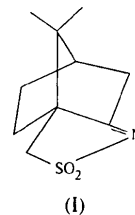
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Abstract

The norbornane ring system in the title molecule, 8,8-dimethyl-3,3a,4,5,6,7-hexahydro-3a,6-methanobenz[c]isothiazole *S,S*-dioxide, C₁₀H₁₅NO₂S, is regular with normal bond lengths and angles. The bridgehead bond angle is 92.5(2)°. The five-membered ring of the sulfonylimine moiety adopts a flattened envelope conformation. The crystal structure is stabilized by weak C—H...O hydrogen bonds.

Comment

Camphor derivatives are important chiral auxiliaries or catalysts for stereoselective synthesis (Oppolzer, 1987). The crystal structure determination of the title compound, (I), was carried out in order to elucidate the molecular conformation.



The bond lengths and angles of the camphor ring system are comparable to those in related molecules (Bear & Trotter, 1975; Garcia *et al.*, 1988). The six-membered ring of the norbornane ring system has a fairly symmetrical boat conformation, with atoms C3 and C7 displaced by $-0.852(3)$ and $-0.845(2)$ Å, respectively, from the best plane through atoms C1, C2, C5 and C6. The bridgehead bond angle (C3—C4—C7) of 92.5(2)° is typical of norbornane derivatives. The two five-membered rings formed by the bridging atom C4 (C1—C2—C3—C4—C7 and C3—C5—C6—C7—C4) adopt envelope conformations with $\Delta C_s(C4)$ asymmetry parameters (Nardelli, 1983a) equal to 0.007(2) and 0.015(2), respectively. The angles between the

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plane of the three-atom bridge C3—C4—C7 and each of the four-atom planes of the boat-shaped six-membered rings (C1, C2, C3, C7 and C3, C5, C6, C7) are 56.0 (2) and 125.8 (2)°, respectively. The five-membered ring of the sulfonylimine moiety adopts a flattened envelope conformation [$\Delta C_s(C8) = 0.007(2)$], with C8 deviating from the mean plane by $-0.290(5)$ Å.

In the crystal, the screw-related molecules are linked by C—H...O bifurcated hydrogen bonds involving O1. This hydrogen bond and other short C—H...O contacts are listed in Table 2.

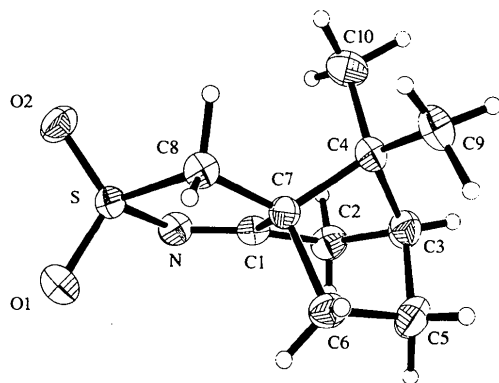


Fig. 1. The structure of title compound showing 50% probability displacement ellipsoids and the atom-numbering scheme.

Experimental

Formaldehyde dibenzylmercaptal (Ray & Chakraborty, 1997) (63 mg, 0.24 mmol) in 1 ml of chloroform was added to a stirred solution of (1R)-(-)-(10-camphorsulfonyl)oxaziridine (50 mg, 0.22 mmol; Aldrich) in 2 ml of chloroform under argon. After stirring for 1 h, (R)-camphorsulfonylimine and (R)-benzyl benzylthiomethyl sulfoxide were isolated by preparative TLC (silica gel, benzene/ethyl acetate, 9:1 then 4:1). Single crystals of (I) were obtained from dichloromethane.

Crystal data

C₁₀H₁₅NO₂S
M_r = 213.29
 Monoclinic
*P*2₁
a = 7.7208 (9) Å
b = 7.7501 (9) Å
c = 9.0531 (10) Å
 β = 94.818 (10)°
V = 539.79 (11) Å³
Z = 2
D_x = 1.312 Mg m⁻³
D_m not measured

Mo *K*α radiation
 λ = 0.71073 Å
 Cell parameters from 36 reflections
 θ = 4.96–11.05°
 μ = 0.275 mm⁻¹
T = 293 (2) K
 Rectangular slab
 0.64 × 0.26 × 0.06 mm
 Colourless

Data collection

Siemens P4 diffractometer
 $\theta/2\theta$ scans
 Absorption correction: none
 3014 measured reflections

θ_{\max} = 27.49°
h = -1 → 10
k = -10 → 10
l = -11 → 11

2468 independent reflections
 1852 reflections with
 $I > 2\sigma(I)$
*R*_{int} = 0.033

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.040
wR (*F*²) = 0.095
S = 0.905
 2468 reflections
 187 parameters
 All H atoms refined
 $w = 1/[\sigma^2(F_o^2) + (0.0502P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$

3 standard reflections
 every 97 reflections
 intensity decay: <3%

$\Delta\rho_{\max} = 0.158$ e Å⁻³
 $\Delta\rho_{\min} = -0.252$ e Å⁻³
 Extinction correction: none
 Scattering factors from
*International Tables for
 Crystallography* (Vol. C)
 Absolute configuration:
 Flack (1983)
 Flack parameter =
 -0.12 (10)

Table 1. Selected geometric parameters (Å, °)

S—N	1.676 (2)	S—C8	1.802 (2)
O1—S—O2	117.22 (11)	N—C1—C7	121.5 (2)
N—S—C8	98.29 (10)	C3—C4—C7	92.5 (2)
N—C1—C2	130.8 (2)	C8—C7—C4	122.3 (2)

Table 2. Hydrogen-bond geometry and short contacts (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C2—H2B...O1 ⁱ	1.12 (6)	2.33 (6)	3.411 (6)	160 (4)
C2—H2A...O2 ⁱⁱ	0.83 (4)	2.61 (4)	3.435 (6)	175 (3)
C8—H8A...O1 ⁱⁱⁱ	1.06 (5)	2.53 (5)	3.534 (5)	158 (4)
C8—H8B...O2 ^{iv}	0.82 (3)	2.65 (3)	3.421 (5)	157 (3)

Symmetry codes: (i) 1 - *x*, *y* - ½, 2 - *z*; (ii) 1 - *x*, ½ + *y*, 2 - *z*; (iii) 2 - *x*, *y* - ½, 2 - *z*; (iv) 2 - *x*, ½ + *y*, 2 - *z*.

Data collection, cell refinement and data reduction: XSCANS (Siemens, 1994). Structure solution and molecular graphics: SHELXTL/PC (Sheldrick, 1990). Structure refinement: SHELXL93 (Sheldrick, 1993). Geometric calculations: PARST (Nardelli, 1983b).

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