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Key indicators

Single-crystal X-ray study T = 294 K Mean σ (C–C) = 0.004 Å R factor = 0.045 wR factor = 0.130 Data-to-parameter ratio = 16.6

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

endo-Bicyclo[2.2.1]hept-2-ene-5,6-dimethylene bis(p-toluenesulfonate)

The structure of the title compound, $C_{23}H_{26}O_6S_2$, contains three rings in a bicyclo[2.2.1] system (two five-membered rings and one six-membered ring). There are four $C-H\cdots O$ hydrogen bonds (three intramolecular hydrogen bonds and one intermolecular hydrogen bond) in the structure.

Comment

The structure determination of the title compound, (I), showed that, in the bicyclo[2.2.1] ring system, both fivemembered rings have envelope conformations (Table 1), atom C15 being the out-of-plane atom in both rings (C10/C11/C12/C13/C15) and C10/C9/C14/C13/C15). The six-membered ring



(C9–C14) displays a boat conformation. There are four C– $H \cdots O$ hydrogen bonds in (I), one of them being intermolecular (Table 2).



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Figure 1

The molecular structure of (I) showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 35% probability level.

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Figure 2 Intermolecular hydrogen bonds in (I), shown as dashed lines.

Experimental

5,6-Bis(endo-hydroxymethyl)bicyclo[2.2.1]hept-2-ene (2 g, 13 mmol) was added to NaH (1.78 g, 74 mmol) in 100 ml of dry tetrahydrofuran. The mixture was stirred at 273 K for 10 min. p-Toluenesulfonyl chloride (5.45 g, 28.6 mmol) was then added. The mixture was stirred at 273 K for another 6 h. The reaction mixture was poured into ice and extracted with dichloromethane. The extract was washed with water, dried over anhydrous sodium sulfate and evaporated in vacuo to give a solid. Colourless prisms of (I) were obtained after recrystallization from a methanol solution of the solid (yield: 3 g, 50 wt%; m.p. 359 K). IR (CHCl₃, cm⁻¹): 1600, 1361, 1173.

Z = 4

 $D_x = 1.339 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation $\mu = 0.27 \text{ mm}^{-1}$

T = 294 (2) K

Prism, colourless

 $0.24 \times 0.20 \times 0.14 \text{ mm}$

Crystal data

$C_{23}H_{26}O_6S_2$
$M_r = 462.56$
Monoclinic, $P2_1/c$
a = 14.139 (2) Å
b = 13.186 (2) Å
c = 13.626 (2) Å
$\beta = 115.424 \ (3)^{\circ}$
V = 2294.4 (6) Å ³

Data collection

Siemens SMART CCD	12673 measured reflections
diffractometer	4679 independent reflections
φ and ω scans	2574 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan	$R_{\rm int} = 0.040$
(SADABS; Sheldrick, 1996)	$\theta_{\rm max} = 26.4^{\circ}$
$T_{\min} = 0.927, \ T_{\max} = 0.963$	

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0547P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.045$	+ 0.5608P]
$wR(F^2) = 0.130$	where $P = (F_0^2 + 2F_c^2)/3$
S = 0.99	$(\Delta/\sigma)_{\rm max} = 0.002$
4679 reflections	$\Delta \rho_{\rm max} = 0.20 \ {\rm e} \ {\rm \AA}^{-3}$
282 parameters	$\Delta \rho_{\rm min} = -0.32 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1 Selected terrior angles (°)

Sciected torsion angles ().						
C10-C11-C12-C13	1.6 (3)	C10-C9-C14-C13	2.4 (3)			





Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
C6-H6···O3	0.93	2.54	2.915 (3)	104
$C8-H8B\cdots O4$	0.97	2.57	3.113 (3)	115
C9−H9···O5 ⁱ	0.98	2.56	3.529 (3)	169
C18−H18···O6	0.93	2.56	2.930 (3)	104

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

All H atoms were observable in a difference Fourier map, although the methyl H atoms were the least clear. Nevertheless, all H atoms were positioned geometrically and refined using a riding model, with C-H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ for any H atoms, C-H =0.98 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ for tertiary H atoms, C-H = 0.97 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ for methylene H atoms, and C-H = 0.96 Å and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms.

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

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