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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.006 Å R factor = 0.035 wR factor = 0.069 Data-to-parameter ratio = 14.0

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

5-Bromo-10-oxa-3-thiatricyclo[5.2.1.0^{1,5}]dec-8-ene 3,3-dioxide

In the title compound, $C_8H_9BrO_3S$, the six-membered ring has a boat conformation, the two O-containing five-membered rings have envelope conformations and the tetrahydrothiophene ring has a twist conformation. The molecules are linked only by weak van der Waals interactions.

Comment

Sulfones are a core functional group in both organic and medicinal chemistry because of their versatile synthetic utility and their use as inhibitors of various types of enzymatic processes (Supuran et al., 2003). Alkenyl sulfones are well known for their ability to inhibit many types of cysteine proteases (Roush et al., 1998; Palmer et al., 1995). Alkenyl sulfones are reversible inhibitors of these enzymes through conjugated addition of the thiol of the active-site cysteine residue. In the synthetic sense, alkenyl sulfones have come to play an important role, acting as efficient Michael acceptors and as π donors in cycloadditon reactions (Simpkins, 1990). We have been studying the intramolecular Diels-Alder reaction of furan-cored compounds in which the tether connecting the diene and dienophile consists of three atoms and contains a heteroatom (Demircan & Parsons, 2002; Büyükgüngör et al., 2005). The title compound, (2), can be derived simply from bromofuranylthioether, (1) (see scheme).



The Br–C bond distance, 1.961 (3) Å, is not significantly different from the value reported for a pure Br–C single bond (1.94 Å; Toprak *et al.*, 2001). The six-membered ring is in a boat conformation. The two O-containing five-membered rings adopt envelope conformations, whereas the tetrahydro-thiophene ring adopts a twist conformation. The molecules are linked only by weak van der Waals interactions.

Experimental

To a solution of *meta*-chloroperbenzoic acid (*m*-CPBA) (300 mg, 1.2 mmol), which had previously been purified and recrystallized from dry diethyl ether, in dichloromethane (10 ml), cooled to 273 K,

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Received 2 August 2006 Accepted 7 August 2006 was added dropwise a solution of 2-(2-bromoallylsulfanylmethyl)furan (0.6 mmol) in dichloromethane (10 ml) over a period of 3 min. The reaction mixture was stirred at room temperature for 4 h and then diluted with cold 4% sodium bicarbonate solution (4 ml). The organic layer was separated, washed with water (20 ml) and concentrated in vacuo. The crude residue was heated in toluene (10 ml) for a further 24 h at 383 K. The solvent was then removed under reduced pressure and the resulting solid was subjected to flash column chromatography to afford (2).

Crystal data

C ₈ H ₉ BrO ₃ S	Z = 8
$M_r = 265.12$	$D_x = 1.899 \text{ Mg m}^{-3}$
Orthorhombic, Pbca	Mo $K\alpha$ radiation
a = 10.1723 (6) Å	$\mu = 4.63 \text{ mm}^{-1}$
b = 10.3446 (9) Å	T = 293 (2) K
c = 17.6278 (10) Å	Plate, colorless
V = 1854.9 (2) Å ³	$0.50\times0.27\times0.03$ mm

Data collection

Stoe IPDS-2 diffractometer (i) scans Absorption correction: integration (X-RED32; Stoe & Cie, 2002) $T_{\min} = 0.232, \ T_{\max} = 0.870$

Refinement

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Refinement on F^2
R[F^2 > 2\sigma(F^2)] = 0.035
wR(F<sup>2</sup>) = 0.069
S = 0.95
1823 reflections
                                                        \Delta \rho_{\rm min} = -0.34 \text{ e } \text{\AA}^{-3}
130 parameters
H atoms treated by a mixture of
   independent and constrained
   refinement
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 $w = 1/[\sigma^2(F_o^2) + (0.024P)^2]$ + 1.8112P] where $P = (F_0^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.30 \text{ e } \text{\AA}^{-3}$

12215 measured reflections

 $R_{\rm int} = 0.102$

 $\theta_{\rm max} = 26.0^\circ$

1823 independent reflections

1370 reflections with $I > 2\sigma(I)$

Atoms H3, H4 and H5, attached to C3, C4 and C5, respectively, were refined freely [C-H = 0.91 (6)-0.98 (5) Å]. The other H atoms were refined using a riding model, with C-H = 0.97 Å and $U_{iso}(H)$ = $1.2U_{eq}(C).$

Data collection: X-AREA (Stoe & Cie, 2002); cell refinement: X-AREA; data reduction: X-RED32 (Stoe & Cie, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).



Figure 1

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

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