organic papers

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Key indicators

Single-crystal X-ray study T = 100 KMean σ (C–C) = 0.004 Å R factor = 0.040 wR factor = 0.107 Data-to-parameter ratio = 15.3

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

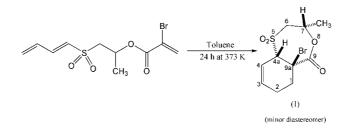
(4aSR,7SR,9aRS)-9a-Bromo-7-methyl-5,5dioxo-1,2,4a,6,7,9a-hexahydro-8-oxa-5 λ^{6} thia-1,4-benzocyclohepten-9-one

The title compound, $C_{10}H_{13}BrO_4S$, crystallizes as a racemic mixture in space group $P\overline{1}$. The seven-membered ring exhibits a boat conformation; the methyl group points away from the bromine substituent.

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Comment

The use of temporary tethers to control the regio- and stereoselectivity of Diels-Alder cycloaddition reactions has received considerable attention (Gauthier et al., 1998; Bear et al., 2001). For example, a variety of silicon-containing tethers (Gauthier et al., 1998; Bear et al., 2001; Coelho & Blanco, 2001; Brosius et al., 1999), and a more limited range of boron-(Gauthier et al., 1998; Bear et al., 2001) and metal alkoxidebased (Ward & Abaee, 2000; Barriault et al., 2003; Bertozzi et al., 2000) tethers have been described. Interestingly, only two types of tethers based on sulfur-containing functional groups have been reported, involving the use of sulfonate esters (Gauthier et al., 1998) or sulfonamides (Brosius et al., 1999). Two of us (NC and PS) have begun to explore the potential utility of sulfone-based tethers for such cycloaddition reactions. We have found that β -acyloxysulfonyl tethers can facilitate highly regioselective and diastereoselective thermally mediated intramolecular Diels-Alder cycloaddition reactions between 1-sulfonylbutadienes and various acrylate ester dienophiles. These studies have resulted in the synthesis of a series of bicyclic β -acyloxysulfone cycloadducts. In several cases, we were able to crystallize the major and/or minor diastereomeric cycloadducts in a form suitable for analysis using single-crystal X-ray diffraction. Details of the synthesis of the cycloadducts will be reported elsewhere (Chumachenko et al., 2004). In the present paper, the solid-state structure of the title cycloadduct, (I), as established by single-crystal X-ray diffraction methods, is described.



Compound (I) crystallizes as a racemic mixture of both enantiomers. The six-membered ring exhibits the half-chair conformation expected for cyclohexenes, and the angles at the unsaturated C atoms are $122.9 (2)^{\circ}$ (for C4–C3–C2) and 124.4 (2)° (for C3–C4–C4A; Fig. 1). The seven-membered

© 2004 International Union of Crystallography Printed in Great Britain – all rights reserved ring displays a boat-like conformation, with the carbonyl and SO₂ groups below the plane defined by the cyclohexene ring and the other groups pointing upwards. The methyl group on atom C7 is located in the equatorial position, pointing away from the bromine substituent on the bridgehead atom C9A.

Experimental

Compound (I) was isolated as the minor diastereomer from an intramolecular Diels-Alder cycloaddition reaction of 1-[(E)-buta-1,3-dienylsulfonyl]propan-2-yl 2-bromoacrylate in toluene at 373 K. Although it partially decomposed during silica column chromatography, we were able to obtain a mixed fraction containing cycloadduct (I) along with decomposition products. Crystals of (I) suitable for X-ray diffraction analysis were grown from this mixture via slow evaporation of a CDCl₃ solution.

Z = 2

 $D_x = 1.769 \text{ Mg m}^{-3}$

Cell parameters from 4693

Mo $K\alpha$ radiation

reflections

 $\theta = 2.3 - 28.3^{\circ}$ $\mu=3.72~\mathrm{mm}^{-1}$

T = 100 (2) K

Block, colorless

 $0.60\,\times\,0.18\,\times\,0.18~\mathrm{mm}$

Crystal data

 $\mathrm{C}_{10}\mathrm{H}_{13}\mathrm{BrO}_{4}\mathrm{S}$ $M_r = 309.17$ Triclinic, P1 a = 6.2069 (10) Åb = 9.2067 (15) Åc = 10.4197 (17) Å $\alpha = 92.724 (3)^{\circ}$ $\beta = 94.191(3)^{\circ}$ $\gamma = 101.615(3)^{\circ}$ $V = 580.50 (16) \text{ Å}^3$

Data collection

Bruker SMART APEX CCD	2814 independent reflections
diffractometer	2627 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.038$
Absorption correction: multi scan	$\theta_{\rm max} = 28.3^{\circ}$
(SADABS in SAINT-Plus;	$h = -8 \rightarrow 8$
Bruker, 1997–1999)	$k = -12 \rightarrow 12$
$T_{\min} = 0.186, \ T_{\max} = 0.510$	$l = -13 \rightarrow 13$
5489 measured reflections	

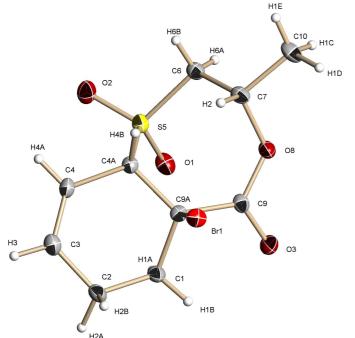


Figure 1

The molecular structure, showing 50% probability displacement ellipsoids.

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_a^2) + (0.0552P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.040$	+ 0.2933 <i>P</i>]
$wR(F^2) = 0.107$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} < 0.001$
2814 reflections	$\Delta \rho_{\rm max} = 1.59 \ {\rm e} \ {\rm \AA}^{-3}$
184 parameters	$\Delta \rho_{\rm min} = -1.25 \text{ e } \text{\AA}^{-3}$
Only coordinates of H atoms	
refined	

Table 1

Selected geometric parameters (Å, °).

Br1-C9A	1.992 (2)	C4A-C9A	1.551 (3)
C1-C2	1.523 (3)	C4A-S5	1.810(2)
C1-C9A	1.525 (3)	\$5-C6	1.777 (3)
O1-S5	1.4428 (19)	C6-C7	1.522 (4)
C2-C3	1.492 (4)	C7-O8	1.458 (3)
O2-S5	1.4381 (19)	C7-C10	1.511 (4)
C3-C4	1.327 (4)	O8-C9	1.342 (3)
O3-C9	1.200 (3)	C9-C9A	1.554 (3)
C4-C4A	1.509 (3)		
C2-C1-C9A	114.24 (19)	O8-C7-C10	105.4 (2)
C3-C2-C1	111.1 (2)	O8-C7-C6	110.17 (19)
C4-C3-C2	122.9 (2)	C10-C7-C6	110.9 (2)
C3-C4-C4A	124.4 (2)	C9-O8-C7	124.22 (18)
C4-C4A-C9A	114.47 (19)	03-C9-08	117.0 (2)
C4-C4A-S5	109.29 (16)	O3-C9-C9A	119.9 (2)
C9A-C4A-S5	110.78 (15)	O8-C9-C9A	123.0 (2)
O2-S5-O1	118.32 (12)	C1-C9A-C4A	111.64 (19)
O2-S5-C6	108.81 (12)	C1-C9A-C9	107.40 (18)
O1-S5-C6	109.25 (11)	C4A-C9A-C9	119.39 (19)
O2-S5-C4A	109.93 (11)	C1-C9A-Br1	109.00 (16)
O1-S5-C4A	107.43 (11)	C4A-C9A-Br1	105.72 (15)
C6-S5-C4A	101.83 (12)	C9-C9A-Br1	103.00 (15)
C7-C6-S5	113.61 (17)		

All H atoms were positioned geometrically. Their coordinates were then refined freely, and their U_{iso} values were defined as 1.2 or $1.5U_{eq}$ of the parent C atom. The s.u. values of the cell parameters are taken from the software, recognizing that the values are unreasonably small (Herbstein, 2000). The maximum and minimum electrondensity peaks are located 0.85 and 0.96 Å, respectively, from atom Br1.

Data collection: SMART (Bruker, 1997-2000); cell refinement: SAINT-Plus (Bruker, 1997–1999); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2000); software used to prepare material for publication: SHELXL97.

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