

(4a*SR*,7*RS*,9a*SR*)-7,9a-Dimethyl-5,5-dioxo-1,2,4a,6,7,9a-hexahydro-8-oxa-5 λ ⁶-thia-1,4-benzocyclohepten-9-one

Matthias Zeller,^a Allen D. Hunter,^{a*} Paul Sampson^b and Nataliya Chumachenko^b

^aDepartment of Chemistry, Youngstown State University, One University Plaza, Youngstown, OH 44555-3663, USA, and ^bDepartment of Chemistry, Kent State University, PO Box 5190, Kent, OH 44242-0001, USA

Correspondence e-mail: adhunter@ysu.edu

Key indicators

Single-crystal X-ray study

$T = 100$ K

Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å

R factor = 0.035

wR factor = 0.097

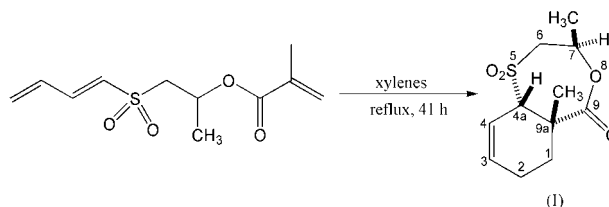
Data-to-parameter ratio = 14.7

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

The title compound, $\text{C}_{11}\text{H}_{16}\text{O}_4\text{S}$, crystallizes as a racemic mixture in space group $P2_1/n$. The seven-membered ring exhibits a chair conformation and the methyl substituents point away from each other.

Comment

As detailed in the first article in this series (Zeller *et al.*, 2004), two of us (NC and PS) have been exploring the potential utility of sulfone-based tethers in intramolecular Diels–Alder cycloaddition reactions. This has resulted in the preparation of a series of bicyclic β -acyloxy sulfone cycloadducts, several of which have been subjected to analysis using single-crystal X-ray diffraction. 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 in space group $P2_1/n$, with $Z = 4$, 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 123.93 (12)° (for $\text{C}4-\text{C}3-\text{C}2$) and 124.05 (12)° (for $\text{C}3-\text{C}4-\text{C}4\text{A}$; Fig. 1). The seven-membered ring displays a chair-like conformation. The methyl group on atom $\text{C}7$ is located in the equatorial position, pointing away from the methyl substituent on the bridgehead atom $\text{C}9\text{A}$.

Experimental

Compound (I) was isolated in 48% yield as the major diastereomer from an intramolecular Diels–Alder cycloaddition reaction of 1-[(*E*)-buta-1,3-dienylsulfonyl]propan-2-yl methacrylate in refluxing xylenes. The crude product yielded crystals of (I) suitable for X-ray diffraction analysis *via* crystallization from ethyl acetate/hexane (1:2) with slow cooling.

Crystal data

$\text{C}_{11}\text{H}_{16}\text{O}_4\text{S}$
 $M_r = 244.30$
 Monoclinic, $P2_1/n$
 $a = 10.0232$ (10) Å
 $b = 10.6255$ (10) Å
 $c = 11.6726$ (11) Å
 $\beta = 114.062$ (2)°
 $V = 1135.13$ (19) Å³
 $Z = 4$

$D_x = 1.430$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 9433 reflections
 $\theta = 2.2-28.3$ °
 $\mu = 0.28$ mm⁻¹
 $T = 100$ (2) K
 Block, colorless
 $0.4 \times 0.2 \times 0.2$ mm

Received 24 March 2004

Accepted 30 March 2004

Online 9 April 2004

Data collection

Bruker SMART APEX CCD
diffractometer

ω scans

Absorption correction: multi-scan
(*SADABS* in *SAINT-Plus*;
Bruker, 1997–1999)

$T_{\min} = 0.735$, $T_{\max} = 0.945$

11460 measured reflections

2829 independent reflections
2729 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.024$

$\theta_{\text{max}} = 28.3^\circ$

$h = -13 \rightarrow 13$

$k = -14 \rightarrow 14$

$l = -15 \rightarrow 15$

Refinement

Refinement on F^2

$R[F^2 > 2\sigma(F^2)] = 0.035$

$wR(F^2) = 0.097$

$S = 1.10$

2829 reflections

193 parameters

Only coordinates of H atoms
refined

$w = 1/[\sigma^2(F_o^2) + (0.0521P)^2 + 0.5113P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\text{max}} = 0.002$

$\Delta\rho_{\text{max}} = 0.44 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.32 \text{ e } \text{\AA}^{-3}$

All H atoms were positioned geometrically. Their coordinates were then refined freely, and their U_{iso} values were defined as 1.2 or 1.5 U_{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).

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*.

MZ was supported by the NSF (grant No. 0111511), and the diffractometer was funded by the NSF (grant No. 0087210), the Ohio Board of Regents (grant No. CAP-491) and YSU. Funds supporting the synthetic studies (NC and PS) were provided by the Department of Chemistry, Kent State University.

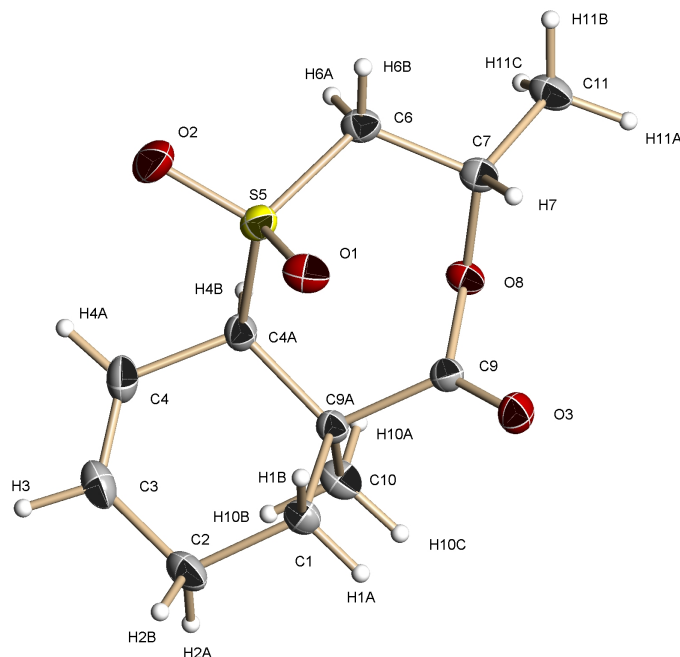


Figure 1
The molecular structure, showing 50% probability displacement ellipsoids.

References

- Bruker (1997–1999). *SAINT-Plus*. Version 6.02. Bruker AXS Inc., Madison, Wisconsin, USA.
 Bruker (1997–2000). *SMART*. Version 5.625 for WNT/2000. Bruker AXS Inc., Madison, Wisconsin, USA.
 Bruker (2000). *SHELXTL*. Version 6.10. Bruker AXS Inc., Madison, Wisconsin, USA.
 Herbstein, F. H. (2000). *Acta Cryst.* **B56**, 547–557.
 Sheldrick, G. M. (1990). *Acta Cryst.* **A46**, 467–473.
 Sheldrick, G. M. (1997). *SHELXL97*. University of Göttingen, Germany.
 Zeller, M., Hunter, A. D., Sampson, P. & Chumachenko, N. (2004). *Acta Cryst.* **E60**, o724–o726.