

5-Chloro-3-cyclohexylsulfinyl-2,7-dimethyl-1-benzofuran

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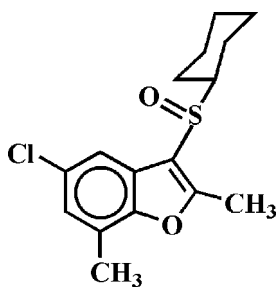
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 Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.044; wR factor = 0.116; data-to-parameter ratio = 19.1.

In the title compound, $\text{C}_{16}\text{H}_{19}\text{ClO}_2\text{S}$, the cyclohexyl ring adopts a chair conformation and the arylsulfinyl unit is positioned equatorial relative to the cyclohexyl group. The least-squares plane through all six C atoms of the cyclohexyl ring makes a dihedral angle of $74.80(6)^\circ$ with the mean plane of the benzofuran fragment. In the crystal, molecules are linked through intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For the pharmacological activity of benzofuran compounds, see: Aslam *et al.* (2009); Galal *et al.* (2009); Khan *et al.* (2005). For natural products with benzofuran rings, see: Akgul & Anil (2003); Soekamto *et al.* (2003). For structural studies of related 3-cyclohexylsulfinyl-5-halo-2-methyl-1-benzofuran derivatives, see: Choi *et al.* (2011a,b,c).



Experimental

Crystal data

 $\text{C}_{16}\text{H}_{19}\text{ClO}_2\text{S}$
 $M_r = 310.82$

Triclinic, $P\bar{1}$
 $a = 5.7480(3)$ Å
 $b = 11.6838(5)$ Å
 $c = 12.2551(5)$ Å
 $\alpha = 70.076(3)^\circ$
 $\beta = 77.244(2)^\circ$
 $\gamma = 83.413(3)^\circ$

$V = 753.98(6)$ Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.39$ mm⁻¹
 $T = 173$ K
 $0.33 \times 0.18 \times 0.14$ mm

Data collection

Bruker SMART APEXII CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2009)
 $T_{\min} = 0.883$, $T_{\max} = 0.948$

13573 measured reflections
 3487 independent reflections
 2752 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.049$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.116$
 $S = 1.06$
 3487 reflections

183 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.44$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.27$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C11}-\text{H11}\cdots\text{O2}^i$	1.00	2.31	3.276 (2)	163

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

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 (Farrugia, 1997) and DIAMOND (Brandenburg, 1998); software used to prepare material for publication: SHELXL97.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GW2105).

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supporting information

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5-Chloro-3-cyclohexylsulfinyl-2,7-dimethyl-1-benzofuran

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S1. Comment

Recently, many compounds containing a benzofuran skeleton have drawn much attention owing to their significant pharmacological properties such as antibacterial and antifungal, antitumor and antiviral, and antimicrobial activities (Aslam *et al.*, 2009, Galal *et al.*, 2009, Khan *et al.*, 2005). These compounds occur in a wide range of natural products (Akgul & Anil, 2003; Soekamto *et al.*, 2003). As a part of our ongoing study of the substituent effect on the solid state structures of 3-cyclohexylsulfinyl-5-halo-2-methyl-1-benzofuran analogues (Choi *et al.*, 2011*a,b,c*), we report herein the crystal structure of the title compound.

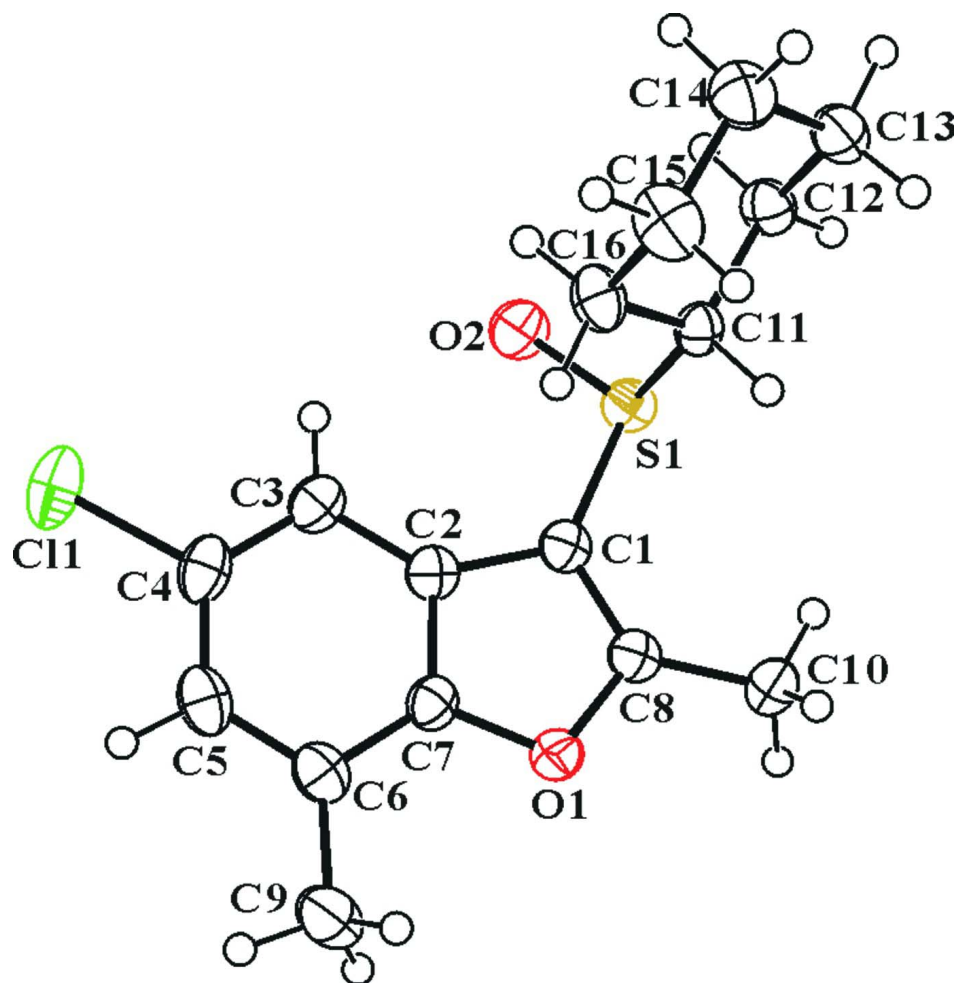
In the title molecule (Fig. 1), the benzofuran unit is essentially planar, with a mean deviation of 0.009 (1) Å from the least-squares plane defined by the nine constituent atoms. The cyclohexyl ring is in the chair form. The least-squares plane through all six C atoms of the cyclohexyl ring makes a dihedral angle of 74.80 (6)° with the mean plane of the benzofuran fragment. The molecular packing (Fig. 2) is stabilized by intermolecular C—H···O hydrogen bonds between a cyclohexyl H atom and the O atom of the sulfinyl group (Table 1; C11—H11···O2ⁱ).

S2. Experimental

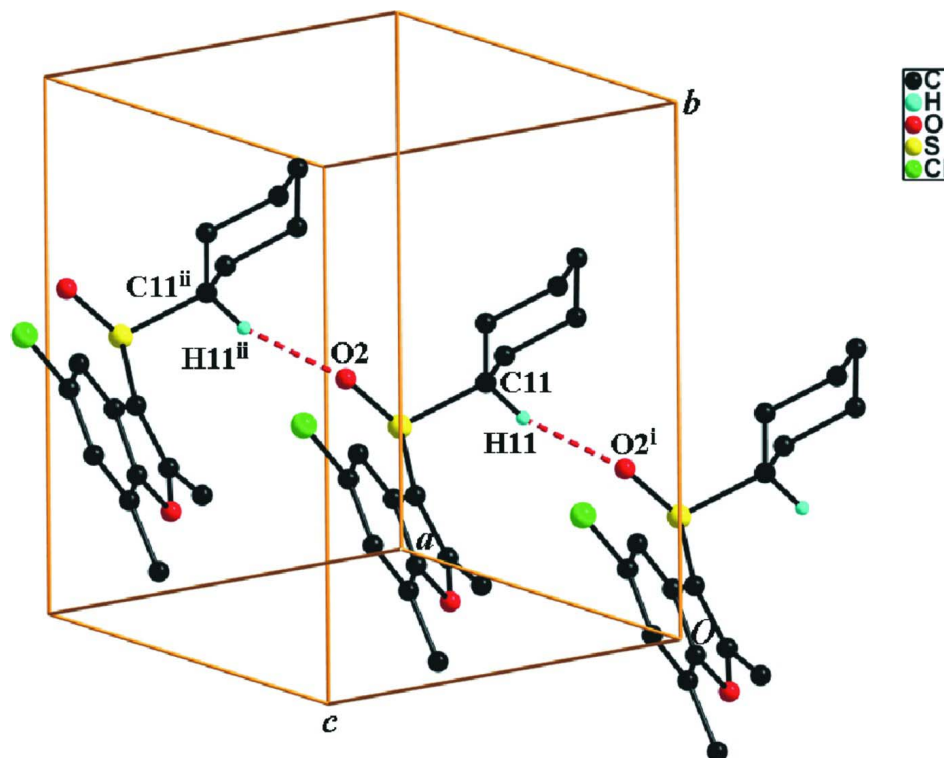
77% 3-chloroperoxybenzoic acid (247 mg, 1.1 mmol) was added in small portions to a stirred solution of 5-chloro-3-cyclohexylsulfonyl-2,7-dimethyl-1-benzofuran (324 mg, 1.1 mmol) in dichloromethane (40 mL) at 273 K. After being stirred at room temperature for 4h, the mixture was washed with saturated sodium bicarbonate solution and the organic layer was separated, dried over magnesium sulfate, filtered and concentrated at reduced pressure. The residue was purified by column chromatography (hexane–ethyl acetate, 1:1 v/v) to afford the title compound as a colorless solid [yield 73%, m.p. 405–406 K; R_f = 0.45 (hexane–ethyl acetate, 1:1 v/v)]. Single crystals suitable for X-ray diffraction were prepared by slow evaporation of a solution of the title compound in ethyl acetate at room temperature.

S3. Refinement

All H atoms were positioned geometrically and refined using a riding model, with C—H = 0.95 Å for aryl, 1.00 Å for methine, 0.99 Å for methylene and 0.98 Å for methyl H atoms, respectively. $U_{iso}(H) = 1.2U_{eq}(C)$ for aryl, methine, methylene, and $1.5U_{eq}(C)$ for methyl H atoms.

**Figure 1**

The molecular structure of the title compound with the atom numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are presented as a small spheres of arbitrary radius.

**Figure 2**

A view of the C—H···O hydrogen bonds (dotted lines) in the crystal structure of the title compound. [Symmetry codes: (i) $x - 1, y, z$; (ii) $x + 1, y, z$.]

5-Chloro-3-cyclohexylsulfinyl-2,7-dimethyl-1-benzofuran

Crystal data

$C_{16}H_{19}ClO_2S$

$M_r = 310.82$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

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$b = 11.6838$ (5) Å

$c = 12.2551$ (5) Å

$\alpha = 70.076$ (3)°

$\beta = 77.244$ (2)°

$\gamma = 83.413$ (3)°

$V = 753.98$ (6) Å³

$Z = 2$

$F(000) = 328$

$D_x = 1.369$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 4179 reflections

$\theta = 3.0$ – 27.4 °

$\mu = 0.39$ mm⁻¹

$T = 173$ K

Block, colourless

$0.33 \times 0.18 \times 0.14$ mm

Data collection

Bruker SMART APEXII CCD
diffractometer

Radiation source: rotating anode

Graphite multilayer monochromator

Detector resolution: 10.0 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2009)

$T_{\min} = 0.883$, $T_{\max} = 0.948$

13573 measured reflections

3487 independent reflections

2752 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.049$

$\theta_{\max} = 27.7$ °, $\theta_{\min} = 1.8$ °

$h = -7 \rightarrow 6$

$k = -15 \rightarrow 15$

$l = -15 \rightarrow 15$

*Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.044$ $wR(F^2) = 0.116$ $S = 1.06$

3487 reflections

183 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
map

Hydrogen site location: difference Fourier map

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0491P)^2 + 0.2498P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.44 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.27 \text{ e } \text{\AA}^{-3}$ *Special details*

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F , and R-factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	1.26205 (10)	0.19475 (6)	0.06775 (5)	0.05045 (18)
S1	0.47811 (9)	0.36573 (4)	0.41383 (4)	0.03095 (15)
O1	0.3493 (2)	0.05713 (11)	0.37764 (11)	0.0314 (3)
O2	0.7264 (3)	0.40766 (12)	0.37025 (14)	0.0415 (4)
C1	0.4738 (4)	0.23324 (15)	0.37493 (16)	0.0288 (4)
C2	0.6529 (3)	0.18767 (16)	0.29578 (16)	0.0288 (4)
C3	0.8708 (4)	0.22616 (17)	0.22279 (17)	0.0333 (4)
H3	0.9363	0.2995	0.2171	0.040*
C4	0.9868 (4)	0.15224 (19)	0.15917 (17)	0.0365 (5)
C5	0.8937 (4)	0.04503 (18)	0.16450 (17)	0.0368 (5)
H5	0.9814	-0.0019	0.1180	0.044*
C6	0.6771 (4)	0.00549 (17)	0.23564 (17)	0.0338 (4)
C7	0.5656 (4)	0.07938 (16)	0.30115 (16)	0.0295 (4)
C8	0.2989 (4)	0.15212 (15)	0.42236 (16)	0.0296 (4)
C9	0.5649 (4)	-0.10708 (18)	0.2428 (2)	0.0449 (5)
H9A	0.6628	-0.1432	0.1855	0.067*
H9B	0.4045	-0.0858	0.2248	0.067*
H9C	0.5539	-0.1659	0.3228	0.067*
C10	0.0724 (4)	0.14949 (17)	0.50709 (17)	0.0347 (4)
H10A	0.0625	0.2173	0.5381	0.052*
H10B	0.0653	0.0722	0.5725	0.052*
H10C	-0.0614	0.1573	0.4671	0.052*
C11	0.2917 (3)	0.46699 (15)	0.31664 (15)	0.0271 (4)
H11	0.1319	0.4306	0.3381	0.033*
C12	0.3878 (4)	0.48526 (18)	0.18677 (17)	0.0365 (5)

H12A	0.3985	0.4062	0.1726	0.044*
H12B	0.5503	0.5168	0.1640	0.044*
C13	0.2246 (5)	0.5751 (2)	0.11125 (18)	0.0458 (6)
H13A	0.2956	0.5905	0.0265	0.055*
H13B	0.0680	0.5388	0.1275	0.055*
C14	0.1878 (5)	0.69445 (19)	0.13643 (18)	0.0436 (5)
H14A	0.0734	0.7484	0.0899	0.052*
H14B	0.3414	0.7354	0.1117	0.052*
C15	0.0929 (4)	0.67408 (17)	0.26706 (18)	0.0374 (5)
H15A	-0.0679	0.6405	0.2899	0.045*
H15B	0.0777	0.7531	0.2818	0.045*
C16	0.2574 (4)	0.58673 (16)	0.34240 (17)	0.0337 (4)
H16A	0.4139	0.6234	0.3250	0.040*
H16B	0.1876	0.5717	0.4272	0.040*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0289 (3)	0.0773 (4)	0.0411 (3)	-0.0036 (3)	-0.0002 (3)	-0.0178 (3)
S1	0.0310 (3)	0.0328 (2)	0.0332 (3)	-0.0023 (2)	-0.0108 (2)	-0.01293 (18)
O1	0.0281 (8)	0.0294 (6)	0.0369 (7)	-0.0023 (6)	-0.0058 (6)	-0.0110 (5)
O2	0.0292 (8)	0.0430 (7)	0.0580 (9)	-0.0037 (7)	-0.0144 (8)	-0.0194 (7)
C1	0.0275 (10)	0.0299 (8)	0.0300 (9)	0.0001 (8)	-0.0086 (9)	-0.0095 (7)
C2	0.0259 (10)	0.0321 (8)	0.0301 (9)	0.0006 (8)	-0.0111 (9)	-0.0092 (7)
C3	0.0290 (11)	0.0382 (9)	0.0328 (10)	-0.0039 (9)	-0.0088 (9)	-0.0092 (8)
C4	0.0253 (11)	0.0504 (11)	0.0307 (10)	0.0004 (9)	-0.0069 (9)	-0.0088 (8)
C5	0.0347 (12)	0.0458 (11)	0.0327 (10)	0.0108 (10)	-0.0116 (10)	-0.0169 (8)
C6	0.0338 (12)	0.0351 (9)	0.0351 (10)	0.0058 (9)	-0.0129 (10)	-0.0132 (8)
C7	0.0251 (10)	0.0323 (8)	0.0305 (9)	-0.0005 (8)	-0.0071 (9)	-0.0084 (7)
C8	0.0292 (11)	0.0296 (8)	0.0311 (9)	0.0002 (8)	-0.0095 (9)	-0.0097 (7)
C9	0.0470 (14)	0.0391 (10)	0.0557 (13)	0.0044 (10)	-0.0139 (12)	-0.0239 (10)
C10	0.0303 (11)	0.0391 (10)	0.0339 (10)	-0.0053 (9)	-0.0029 (9)	-0.0116 (8)
C11	0.0227 (10)	0.0315 (8)	0.0293 (9)	-0.0021 (8)	-0.0060 (8)	-0.0118 (7)
C12	0.0382 (12)	0.0432 (10)	0.0306 (9)	0.0054 (9)	-0.0069 (10)	-0.0174 (8)
C13	0.0556 (16)	0.0515 (12)	0.0327 (10)	0.0089 (11)	-0.0156 (11)	-0.0160 (9)
C14	0.0481 (15)	0.0425 (11)	0.0357 (11)	0.0048 (10)	-0.0107 (11)	-0.0074 (8)
C15	0.0389 (13)	0.0337 (9)	0.0399 (11)	0.0027 (9)	-0.0060 (10)	-0.0146 (8)
C16	0.0378 (12)	0.0337 (9)	0.0329 (10)	-0.0010 (9)	-0.0061 (9)	-0.0157 (8)

Geometric parameters (Å, °)

C11—C4	1.739 (2)	C10—H10A	0.9800
S1—O2	1.4860 (15)	C10—H10B	0.9800
S1—C1	1.7723 (18)	C10—H10C	0.9800
S1—C11	1.8107 (19)	C11—C12	1.514 (3)
O1—C7	1.373 (2)	C11—C16	1.519 (2)
O1—C8	1.374 (2)	C11—H11	1.0000
C1—C8	1.354 (3)	C12—C13	1.528 (3)

C1—C2	1.438 (3)	C12—H12A	0.9900
C2—C7	1.390 (2)	C12—H12B	0.9900
C2—C3	1.390 (3)	C13—C14	1.510 (3)
C3—C4	1.377 (3)	C13—H13A	0.9900
C3—H3	0.9500	C13—H13B	0.9900
C4—C5	1.394 (3)	C14—C15	1.517 (3)
C5—C6	1.379 (3)	C14—H14A	0.9900
C5—H5	0.9500	C14—H14B	0.9900
C6—C7	1.385 (3)	C15—C16	1.520 (3)
C6—C9	1.498 (3)	C15—H15A	0.9900
C8—C10	1.471 (3)	C15—H15B	0.9900
C9—H9A	0.9800	C16—H16A	0.9900
C9—H9B	0.9800	C16—H16B	0.9900
C9—H9C	0.9800		
O2—S1—C1	106.29 (9)	H10A—C10—H10C	109.5
O2—S1—C11	108.27 (9)	H10B—C10—H10C	109.5
C1—S1—C11	98.88 (8)	C12—C11—C16	111.89 (15)
C7—O1—C8	106.36 (14)	C12—C11—S1	113.85 (13)
C8—C1—C2	107.62 (16)	C16—C11—S1	107.54 (12)
C8—C1—S1	124.13 (15)	C12—C11—H11	107.8
C2—C1—S1	128.16 (14)	C16—C11—H11	107.8
C7—C2—C3	119.49 (17)	S1—C11—H11	107.8
C7—C2—C1	104.58 (17)	C11—C12—C13	110.31 (17)
C3—C2—C1	135.92 (17)	C11—C12—H12A	109.6
C4—C3—C2	116.40 (18)	C13—C12—H12A	109.6
C4—C3—H3	121.8	C11—C12—H12B	109.6
C2—C3—H3	121.8	C13—C12—H12B	109.6
C3—C4—C5	123.1 (2)	H12A—C12—H12B	108.1
C3—C4—C11	118.46 (16)	C14—C13—C12	111.64 (18)
C5—C4—C11	118.48 (17)	C14—C13—H13A	109.3
C6—C5—C4	121.56 (19)	C12—C13—H13A	109.3
C6—C5—H5	119.2	C14—C13—H13B	109.3
C4—C5—H5	119.2	C12—C13—H13B	109.3
C5—C6—C7	114.61 (18)	H13A—C13—H13B	108.0
C5—C6—C9	124.04 (19)	C13—C14—C15	110.96 (17)
C7—C6—C9	121.34 (19)	C13—C14—H14A	109.4
O1—C7—C6	124.32 (17)	C15—C14—H14A	109.4
O1—C7—C2	110.81 (16)	C13—C14—H14B	109.4
C6—C7—C2	124.84 (19)	C15—C14—H14B	109.4
C1—C8—O1	110.61 (17)	H14A—C14—H14B	108.0
C1—C8—C10	132.59 (17)	C14—C15—C16	111.22 (17)
O1—C8—C10	116.78 (15)	C14—C15—H15A	109.4
C6—C9—H9A	109.5	C16—C15—H15A	109.4
C6—C9—H9B	109.5	C14—C15—H15B	109.4
H9A—C9—H9B	109.5	C16—C15—H15B	109.4
C6—C9—H9C	109.5	H15A—C15—H15B	108.0
H9A—C9—H9C	109.5	C11—C16—C15	110.08 (16)

H9B—C9—H9C	109.5	C11—C16—H16A	109.6
C8—C10—H10A	109.5	C15—C16—H16A	109.6
C8—C10—H10B	109.5	C11—C16—H16B	109.6
H10A—C10—H10B	109.5	C15—C16—H16B	109.6
C8—C10—H10C	109.5	H16A—C16—H16B	108.2
O2—S1—C1—C8	163.69 (16)	C3—C2—C7—O1	-179.98 (16)
C11—S1—C1—C8	-84.21 (17)	C1—C2—C7—O1	-0.2 (2)
O2—S1—C1—C2	-12.42 (19)	C3—C2—C7—C6	-1.9 (3)
C11—S1—C1—C2	99.67 (18)	C1—C2—C7—C6	177.87 (18)
C8—C1—C2—C7	0.8 (2)	C2—C1—C8—O1	-1.2 (2)
S1—C1—C2—C7	177.46 (14)	S1—C1—C8—O1	-177.97 (13)
C8—C1—C2—C3	-179.5 (2)	C2—C1—C8—C10	-179.5 (2)
S1—C1—C2—C3	-2.8 (3)	S1—C1—C8—C10	3.7 (3)
C7—C2—C3—C4	0.2 (3)	C7—O1—C8—C1	1.0 (2)
C1—C2—C3—C4	-179.4 (2)	C7—O1—C8—C10	179.65 (16)
C2—C3—C4—C5	0.9 (3)	O2—S1—C11—C12	51.25 (15)
C2—C3—C4—C11	-179.20 (14)	C1—S1—C11—C12	-59.27 (15)
C3—C4—C5—C6	-0.5 (3)	O2—S1—C11—C16	-73.30 (14)
C11—C4—C5—C6	179.60 (15)	C1—S1—C11—C16	176.18 (13)
C4—C5—C6—C7	-1.0 (3)	C16—C11—C12—C13	-55.8 (2)
C4—C5—C6—C9	178.37 (19)	S1—C11—C12—C13	-178.01 (14)
C8—O1—C7—C6	-178.57 (18)	C11—C12—C13—C14	55.1 (2)
C8—O1—C7—C2	-0.5 (2)	C12—C13—C14—C15	-55.6 (3)
C5—C6—C7—O1	-179.95 (17)	C13—C14—C15—C16	56.4 (3)
C9—C6—C7—O1	0.6 (3)	C12—C11—C16—C15	56.7 (2)
C5—C6—C7—C2	2.2 (3)	S1—C11—C16—C15	-177.55 (14)
C9—C6—C7—C2	-177.18 (19)	C14—C15—C16—C11	-56.5 (2)

Hydrogen-bond geometry (\AA , $^\circ$)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C11—H11 \cdots O2 ⁱ	1.00	2.31	3.276 (2)	163

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