

5-Iodo-2,7-dimethyl-3-(4-methylphenylsulfonyl)-1-benzofuran

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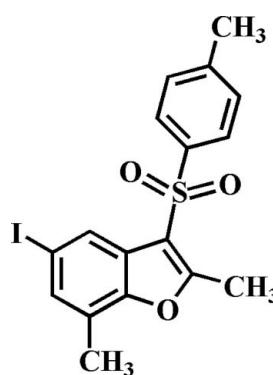
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Key indicators: single-crystal X-ray study; $T = 173\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.027; wR factor = 0.070; data-to-parameter ratio = 20.2.

In the title compound, $\text{C}_{17}\text{H}_{15}\text{IO}_3\text{S}$, the 4-methylphenyl ring makes a dihedral angle of $76.95(5)^\circ$ with the mean plane [r.m.s. deviation = $0.019(2)\text{ \AA}$] of the benzofuran fragment. In the crystal, molecules are linked via pairs of $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, forming inversion dimers. These dimers are connected by slipped $\pi-\pi$ interactions between the benzene rings of neighbouring molecules [centroid–centroid distance = $3.671(3)\text{ \AA}$ and slippage = $1.049(3)\text{ \AA}$].

Related literature

For background information and the crystal structures of related compounds, see: Choi *et al.* (2008); Seo *et al.* (2012).



Experimental

Crystal data

$\text{C}_{17}\text{H}_{15}\text{IO}_3\text{S}$
 $M_r = 426.25$
Monoclinic, $P2_1/n$
 $a = 11.5480(5)\text{ \AA}$
 $b = 9.9394(4)\text{ \AA}$
 $c = 14.8911(6)\text{ \AA}$
 $\beta = 107.611(1)^\circ$

$V = 1629.10(12)\text{ \AA}^3$
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 2.10\text{ mm}^{-1}$
 $T = 173\text{ K}$
 $0.33 \times 0.27 \times 0.22\text{ mm}$

Data collection

Bruker SMART APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2009)
 $T_{\min} = 0.634$, $T_{\max} = 0.746$

16012 measured reflections
4075 independent reflections
3619 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.027$
 $wR(F^2) = 0.070$
 $S = 1.05$
4075 reflections

202 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.49\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.91\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C16—H16···O3 ⁱ	0.95	2.58	3.246 (2)	127

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

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: IM2395).

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supporting information

Acta Cryst. (2012). E68, o2893 [https://doi.org/10.1107/S1600536812037932]

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S1. Comment

As a part of our ongoing study of 5-iodo-2,7-dimethyl-benzofuran derivatives containing either phenyl-sulfonyl (Choi *et al.*, 2008) or 4-fluorophenyl-sulfonyl (Seo *et al.*, 2012) substituents in 3-position, 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 0.019 (2) Å from the least-squares plane defined by the nine constituent atoms. The dihedral angle between the 4-methylphenyl ring and the mean plane of the benzofuran fragment is 76.95 (5)°. In the crystal structure, molecules are linked via pairs of C—H···O hydrogen bonds (Fig. 2 & Table 1), forming inversion dimers. These dimers are connected by slipped π – π interactions between the benzene rings of neighbouring molecules, with a Cg···Cgⁱⁱ distance of 3.671 (3) Å and an interplanar distance of 3.518 (2) Å resulting in a slippage of 1.049 (3) Å (Fig. 2, Cg is the centroid of the C2–C7 benzene ring).

S2. Experimental

3-Chloroperoxybenzoic acid (77%, 381 mg, 1.7 mmol) was added in small portions to a stirred solution of 5-iodo-2,7-dimethyl-3-(4-methylphenylsulfonyl)-benzofuran (315 mg, 0.8 mmol) in dichloromethane (50 mL) at 273 K. After being stirred at room temperature for 10 h, the mixture was washed with saturated sodium bicarbonate solution, the organic layer was separated and dried over magnesium sulfate. After filtration the solution was concentrated at reduced pressure. The residue was purified by column chromatography (benzene) to afford the title compound as a colorless solid [yield 68%, m.p. 483–484 K; R_f = 0.56 (benzene)]. Single crystals suitable for X-ray diffraction were prepared by slow evaporation of a solution of the title compound in acetone 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 and 0.98 Å for methyl H atoms. $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aryl and $1.5U_{\text{eq}}(\text{C})$ for methyl H atoms. The positions of methyl hydrogens were optimized rotationally.

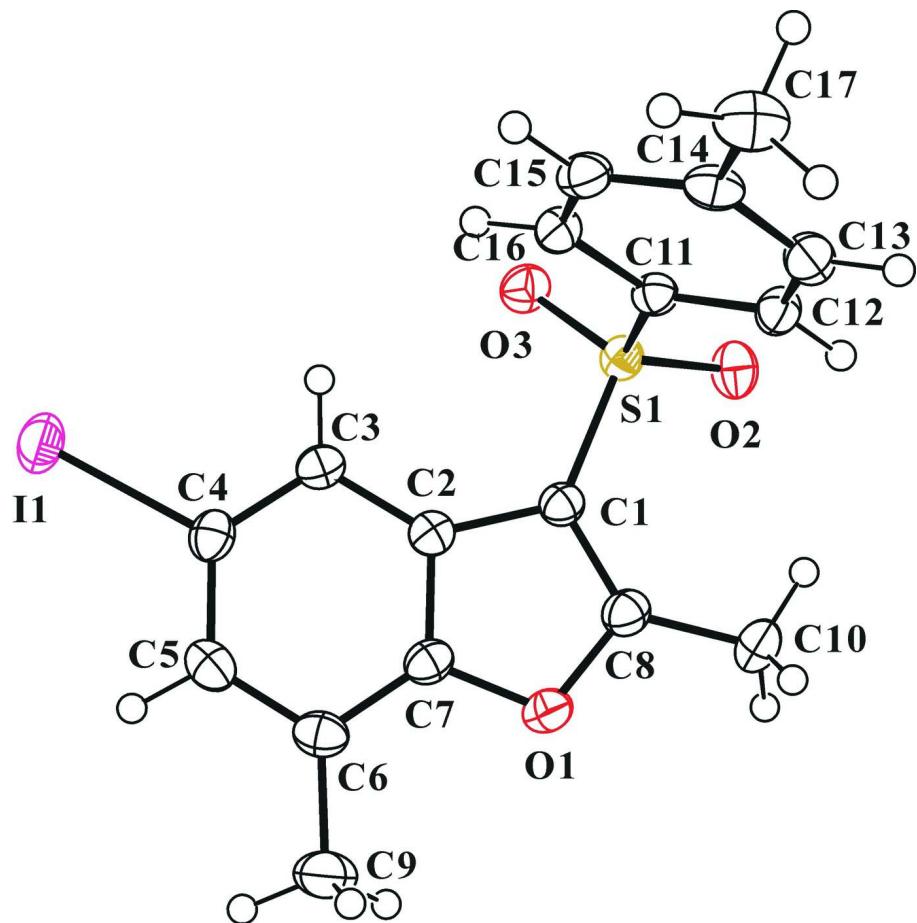
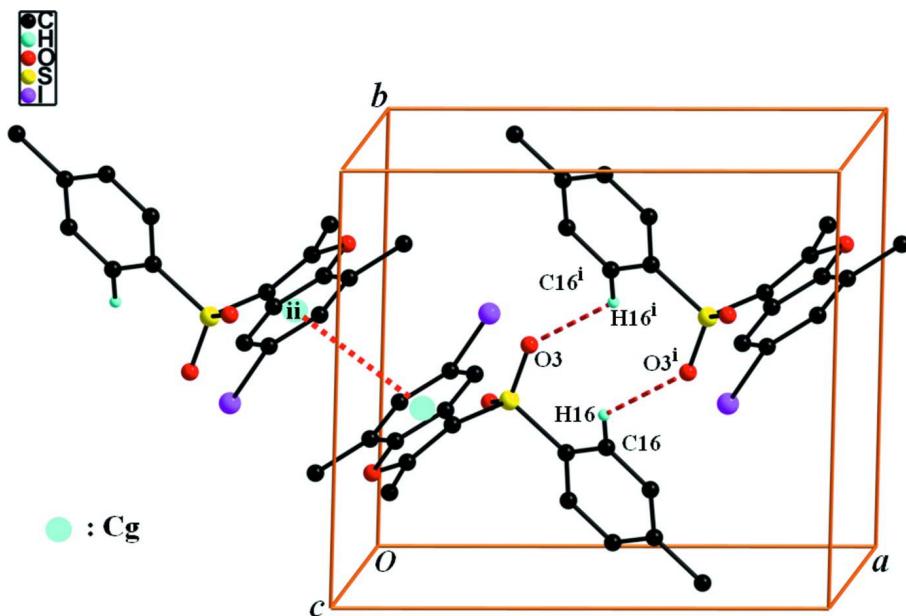


Figure 1

Molecular structure of the title compound with displacement ellipsoids drawn at the 50% probability level. H atoms are presented as small spheres of arbitrary radius.

**Figure 2**

A view of the C—H···O and π ··· π interactions (dotted lines) in the crystal structure of the title compound. H atoms not participating in hydrogen-bonding were omitted for clarity. [Symmetry codes: (i) $-x + 1, -y + 1, -z + 1$; (ii) $-x, -y + 1, -z + 1$.]

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Crystal data

$C_{17}H_{15}IO_3S$
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 $c = 14.8911 (6)$ Å
 $\beta = 107.611 (1)$ °
 $V = 1629.10 (12)$ Å³
 $Z = 4$

$F(000) = 840$
 $D_x = 1.738$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 7311 reflections
 $\theta = 2.5\text{--}28.3$ °
 $\mu = 2.10$ mm⁻¹
 $T = 173$ K
Block, colourless
 $0.33 \times 0.27 \times 0.22$ 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)
 $T_{\min} = 0.634$, $T_{\max} = 0.746$

16012 measured reflections
4075 independent reflections
3619 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$
 $\theta_{\max} = 28.4$ °, $\theta_{\min} = 2.0$ °
 $h = -15 \rightarrow 15$
 $k = -11 \rightarrow 13$
 $l = -19 \rightarrow 19$

*Refinement*Refinement on F^2

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.027$$

$$wR(F^2) = 0.070$$

$$S = 1.05$$

4075 reflections

202 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

$$w = 1/[\sigma^2(F_o^2) + (0.0382P)^2 + 0.5412P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} = 0.001$$

$$\Delta\rho_{\max} = 0.49 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.91 \text{ e \AA}^{-3}$$

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 $F^2 > 2\text{sigma}(F^2)$ 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>	$U_{\text{iso}}^*/U_{\text{eq}}$
I1	0.279119 (14)	0.635214 (15)	0.727688 (10)	0.03911 (7)
S1	0.28800 (5)	0.38147 (4)	0.33837 (3)	0.02292 (10)
O1	0.02336 (13)	0.22668 (14)	0.42013 (11)	0.0294 (3)
O2	0.23919 (16)	0.36296 (14)	0.23798 (11)	0.0318 (3)
O3	0.32866 (14)	0.51329 (13)	0.37468 (10)	0.0294 (3)
C1	0.17945 (18)	0.33310 (19)	0.39126 (14)	0.0235 (4)
C2	0.17346 (18)	0.37898 (17)	0.48211 (14)	0.0230 (4)
C3	0.23659 (18)	0.47147 (19)	0.54969 (14)	0.0258 (4)
H3	0.3051	0.5191	0.5435	0.031*
C4	0.1941 (2)	0.4900 (2)	0.62619 (14)	0.0278 (4)
C5	0.0950 (2)	0.4200 (2)	0.63774 (15)	0.0303 (4)
H5	0.0701	0.4369	0.6920	0.036*
C6	0.03155 (19)	0.3262 (2)	0.57189 (15)	0.0290 (4)
C7	0.07526 (18)	0.31046 (19)	0.49528 (14)	0.0253 (4)
C8	0.08764 (19)	0.2427 (2)	0.35754 (15)	0.0277 (4)
C9	-0.0757 (2)	0.2491 (3)	0.58031 (19)	0.0416 (5)
H9A	-0.1400	0.2503	0.5197	0.062*
H9B	-0.1058	0.2906	0.6286	0.062*
H9C	-0.0518	0.1560	0.5981	0.062*
C10	0.0456 (2)	0.1610 (2)	0.27101 (18)	0.0397 (5)
H10A	0.0914	0.1860	0.2278	0.059*
H10B	-0.0412	0.1772	0.2406	0.059*
H10C	0.0587	0.0654	0.2872	0.059*
C11	0.40852 (17)	0.26776 (18)	0.38089 (13)	0.0222 (4)
C12	0.4107 (2)	0.15132 (19)	0.32984 (14)	0.0260 (4)

H12	0.3491	0.1343	0.2722	0.031*
C13	0.5049 (2)	0.0601 (2)	0.36479 (15)	0.0288 (4)
H13	0.5070	-0.0200	0.3305	0.035*
C14	0.59577 (19)	0.0837 (2)	0.44849 (15)	0.0277 (4)
C15	0.59026 (19)	0.2003 (2)	0.49941 (15)	0.0279 (4)
H15	0.6511	0.2167	0.5576	0.033*
C16	0.49679 (19)	0.2923 (2)	0.46593 (14)	0.0257 (4)
H16	0.4933	0.3712	0.5009	0.031*
C17	0.6981 (2)	-0.0151 (2)	0.48381 (19)	0.0383 (5)
H17A	0.6676	-0.1066	0.4671	0.057*
H17B	0.7315	-0.0075	0.5524	0.057*
H17C	0.7620	0.0044	0.4549	0.057*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
I1	0.03727 (11)	0.04423 (10)	0.03268 (10)	-0.00198 (6)	0.00588 (7)	-0.01371 (6)
S1	0.0247 (2)	0.0226 (2)	0.0205 (2)	-0.00060 (17)	0.00545 (19)	0.00125 (16)
O1	0.0244 (7)	0.0299 (7)	0.0336 (8)	-0.0057 (6)	0.0082 (6)	-0.0044 (6)
O2	0.0369 (9)	0.0364 (8)	0.0192 (7)	0.0038 (6)	0.0040 (7)	0.0024 (5)
O3	0.0339 (8)	0.0217 (6)	0.0328 (8)	-0.0032 (6)	0.0106 (7)	0.0010 (5)
C1	0.0213 (9)	0.0236 (8)	0.0248 (9)	0.0005 (7)	0.0058 (8)	0.0001 (7)
C2	0.0209 (9)	0.0219 (8)	0.0249 (9)	0.0031 (7)	0.0052 (8)	0.0016 (7)
C3	0.0222 (10)	0.0263 (9)	0.0278 (10)	-0.0010 (7)	0.0058 (8)	0.0002 (7)
C4	0.0268 (10)	0.0294 (9)	0.0238 (10)	0.0019 (8)	0.0027 (8)	-0.0011 (7)
C5	0.0296 (11)	0.0363 (10)	0.0268 (10)	0.0041 (9)	0.0113 (9)	0.0037 (8)
C6	0.0239 (10)	0.0313 (10)	0.0319 (11)	0.0017 (8)	0.0087 (9)	0.0058 (8)
C7	0.0220 (10)	0.0230 (9)	0.0290 (10)	0.0001 (7)	0.0047 (8)	0.0004 (7)
C8	0.0242 (10)	0.0264 (9)	0.0315 (11)	0.0000 (8)	0.0068 (8)	-0.0023 (8)
C9	0.0330 (13)	0.0508 (14)	0.0434 (13)	-0.0100 (10)	0.0153 (11)	0.0022 (11)
C10	0.0372 (13)	0.0400 (12)	0.0421 (13)	-0.0113 (10)	0.0124 (11)	-0.0185 (10)
C11	0.0215 (9)	0.0230 (8)	0.0221 (9)	-0.0015 (7)	0.0063 (7)	0.0015 (7)
C12	0.0273 (10)	0.0271 (9)	0.0226 (9)	-0.0037 (8)	0.0062 (8)	-0.0027 (7)
C13	0.0328 (11)	0.0248 (9)	0.0314 (10)	-0.0008 (8)	0.0137 (9)	-0.0004 (7)
C14	0.0249 (10)	0.0277 (9)	0.0332 (11)	-0.0017 (8)	0.0127 (9)	0.0070 (8)
C15	0.0222 (10)	0.0345 (10)	0.0250 (10)	-0.0055 (8)	0.0041 (8)	0.0020 (8)
C16	0.0262 (10)	0.0267 (9)	0.0246 (9)	-0.0042 (8)	0.0082 (8)	-0.0023 (7)
C17	0.0308 (12)	0.0344 (11)	0.0489 (14)	0.0051 (9)	0.0108 (11)	0.0111 (10)

Geometric parameters (\AA , ^\circ)

I1—C4	2.106 (2)	C9—H9B	0.9800
S1—O3	1.4408 (14)	C9—H9C	0.9800
S1—O2	1.4410 (16)	C10—H10A	0.9800
S1—C1	1.738 (2)	C10—H10B	0.9800
S1—C11	1.7556 (19)	C10—H10C	0.9800
O1—C8	1.365 (2)	C11—C16	1.386 (3)
O1—C7	1.378 (2)	C11—C12	1.389 (3)

C1—C8	1.364 (3)	C12—C13	1.390 (3)
C1—C2	1.449 (3)	C12—H12	0.9500
C2—C7	1.387 (3)	C13—C14	1.385 (3)
C2—C3	1.395 (3)	C13—H13	0.9500
C3—C4	1.382 (3)	C14—C15	1.396 (3)
C3—H3	0.9500	C14—C17	1.504 (3)
C4—C5	1.394 (3)	C15—C16	1.387 (3)
C5—C6	1.391 (3)	C15—H15	0.9500
C5—H5	0.9500	C16—H16	0.9500
C6—C7	1.389 (3)	C17—H17A	0.9800
C6—C9	1.494 (3)	C17—H17B	0.9800
C8—C10	1.476 (3)	C17—H17C	0.9800
C9—H9A	0.9800		
O3—S1—O2	119.09 (9)	C6—C9—H9C	109.5
O3—S1—C1	106.23 (9)	H9A—C9—H9C	109.5
O2—S1—C1	108.99 (10)	H9B—C9—H9C	109.5
O3—S1—C11	108.50 (9)	C8—C10—H10A	109.5
O2—S1—C11	108.09 (9)	C8—C10—H10B	109.5
C1—S1—C11	105.09 (9)	H10A—C10—H10B	109.5
C8—O1—C7	106.93 (15)	C8—C10—H10C	109.5
C8—C1—C2	107.44 (17)	H10A—C10—H10C	109.5
C8—C1—S1	127.02 (16)	H10B—C10—H10C	109.5
C2—C1—S1	125.50 (15)	C16—C11—C12	121.08 (18)
C7—C2—C3	119.54 (18)	C16—C11—S1	120.01 (15)
C7—C2—C1	104.45 (17)	C12—C11—S1	118.86 (15)
C3—C2—C1	135.97 (18)	C11—C12—C13	118.69 (19)
C4—C3—C2	116.36 (18)	C11—C12—H12	120.7
C4—C3—H3	121.8	C13—C12—H12	120.7
C2—C3—H3	121.8	C14—C13—C12	121.37 (19)
C3—C4—C5	123.01 (19)	C14—C13—H13	119.3
C3—C4—I1	118.57 (15)	C12—C13—H13	119.3
C5—C4—I1	118.38 (15)	C13—C14—C15	118.84 (19)
C6—C5—C4	121.65 (19)	C13—C14—C17	120.4 (2)
C6—C5—H5	119.2	C15—C14—C17	120.8 (2)
C4—C5—H5	119.2	C16—C15—C14	120.7 (2)
C7—C6—C5	114.20 (19)	C16—C15—H15	119.7
C7—C6—C9	121.9 (2)	C14—C15—H15	119.7
C5—C6—C9	123.9 (2)	C11—C16—C15	119.33 (19)
O1—C7—C2	110.81 (17)	C11—C16—H16	120.3
O1—C7—C6	123.92 (18)	C15—C16—H16	120.3
C2—C7—C6	125.24 (19)	C14—C17—H17A	109.5
C1—C8—O1	110.37 (18)	C14—C17—H17B	109.5
C1—C8—C10	134.2 (2)	H17A—C17—H17B	109.5
O1—C8—C10	115.38 (18)	C14—C17—H17C	109.5
C6—C9—H9A	109.5	H17A—C17—H17C	109.5
C6—C9—H9B	109.5	H17B—C17—H17C	109.5
H9A—C9—H9B	109.5		

O3—S1—C1—C8	155.49 (18)	C9—C6—C7—O1	1.6 (3)
O2—S1—C1—C8	26.0 (2)	C5—C6—C7—C2	0.3 (3)
C11—S1—C1—C8	-89.6 (2)	C9—C6—C7—C2	179.4 (2)
O3—S1—C1—C2	-26.97 (19)	C2—C1—C8—O1	-0.2 (2)
O2—S1—C1—C2	-156.44 (16)	S1—C1—C8—O1	177.67 (14)
C11—S1—C1—C2	87.91 (18)	C2—C1—C8—C10	-179.7 (2)
C8—C1—C2—C7	-0.2 (2)	S1—C1—C8—C10	-1.8 (4)
S1—C1—C2—C7	-178.18 (15)	C7—O1—C8—C1	0.6 (2)
C8—C1—C2—C3	-177.5 (2)	C7—O1—C8—C10	-179.82 (19)
S1—C1—C2—C3	4.5 (3)	O3—S1—C11—C16	27.66 (18)
C7—C2—C3—C4	-1.0 (3)	O2—S1—C11—C16	158.10 (15)
C1—C2—C3—C4	176.0 (2)	C1—S1—C11—C16	-85.63 (17)
C2—C3—C4—C5	0.9 (3)	O3—S1—C11—C12	-154.88 (15)
C2—C3—C4—I1	-176.91 (14)	O2—S1—C11—C12	-24.44 (18)
C3—C4—C5—C6	-0.2 (3)	C1—S1—C11—C12	91.83 (16)
I1—C4—C5—C6	177.66 (16)	C16—C11—C12—C13	-1.2 (3)
C4—C5—C6—C7	-0.4 (3)	S1—C11—C12—C13	-178.63 (15)
C4—C5—C6—C9	-179.5 (2)	C11—C12—C13—C14	-0.3 (3)
C8—O1—C7—C2	-0.8 (2)	C12—C13—C14—C15	1.5 (3)
C8—O1—C7—C6	177.30 (19)	C12—C13—C14—C17	-178.7 (2)
C3—C2—C7—O1	178.48 (17)	C13—C14—C15—C16	-1.3 (3)
C1—C2—C7—O1	0.6 (2)	C17—C14—C15—C16	178.98 (19)
C3—C2—C7—C6	0.4 (3)	C12—C11—C16—C15	1.5 (3)
C1—C2—C7—C6	-177.42 (19)	S1—C11—C16—C15	178.85 (15)
C5—C6—C7—O1	-177.48 (18)	C14—C15—C16—C11	-0.2 (3)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C16—H16···O3 ⁱ	0.95	2.58	3.246 (2)	127

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