

Propyl 2-(5-chloro-3-methylsulfinyl-1-benzofuran-2-yl)acetate

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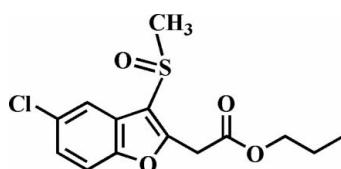
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Key indicators: single-crystal X-ray study; $T = 173\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.037; wR factor = 0.094; data-to-parameter ratio = 13.6.

In the title compound, $\text{C}_{14}\text{H}_{15}\text{ClO}_4\text{S}$, the O atom and the methyl group of the methylsulfinyl substituent lie on opposite sides of the plane of the benzofuran fragment. The crystal structure is stabilized by aromatic $\pi-\pi$ interactions between the benzene rings of neighbouring molecules [centroid-to-centroid distance = $3.635(3)\text{ \AA}$], and by C–H··· π interactions between a propyl methylene H atom and the furan ring of an adjacent molecule. In addition, the crystal structure exhibits weak intermolecular C–H···O hydrogen bonds.

Related literature

For the crystal structures of similar alkyl 2-(5-chloro-3-methylsulfinyl-1-benzofuran-2-yl)acetates, see: Choi *et al.* (2007, 2008).



Experimental

Crystal data



$M_r = 314.77$

Triclinic, $P\bar{1}$
 $a = 8.528(2)\text{ \AA}$
 $b = 9.585(3)\text{ \AA}$
 $c = 10.195(3)\text{ \AA}$
 $\alpha = 73.452(4)^\circ$
 $\beta = 81.773(5)^\circ$
 $\gamma = 65.747(4)^\circ$

$V = 728.0(4)\text{ \AA}^3$
 $Z = 2$
 $\text{Mo } K\alpha \text{ radiation}$
 $\mu = 0.42\text{ mm}^{-1}$
 $T = 173\text{ K}$
 $0.40 \times 0.40 \times 0.10\text{ mm}$

Data collection

Bruker SMART CCD
diffractometer
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1999)
 $R_{\text{int}} = 0.029$
 $T_{\text{min}} = 0.842$, $T_{\text{max}} = 0.961$

5062 measured reflections
2468 independent reflections
2304 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.029$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.094$
 $S = 1.07$
2468 reflections

182 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.28\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.32\text{ e \AA}^{-3}$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C12–H12A···Cg2 ⁱ	0.99	2.74	3.666(3)	155
C3–H3···O4 ⁱⁱ	0.95	2.44	3.353(2)	160
C5–H5···O3 ⁱⁱⁱ	0.95	2.50	3.373(2)	152
C9–H9A···O4 ^{iv}	0.99	2.35	3.321(2)	166
C9–H9B···O1 ^v	0.99	2.54	3.489(2)	161

Symmetry codes: (i) $x, y+1, z$; (ii) $-x, -y+1, -z+1$; (iii) $-x, -y+1, -z+2$; (iv) $-x+1, -y+1, -z+1$; (v) $-x+1, -y+1, -z+2$. Cg2 is the centroid of the C1/C2/C7/O1/C8 furan ring

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); 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*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2190).

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supporting information

Acta Cryst. (2009). E65, o521 [doi:10.1107/S1600536809004735]

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S1. Comment

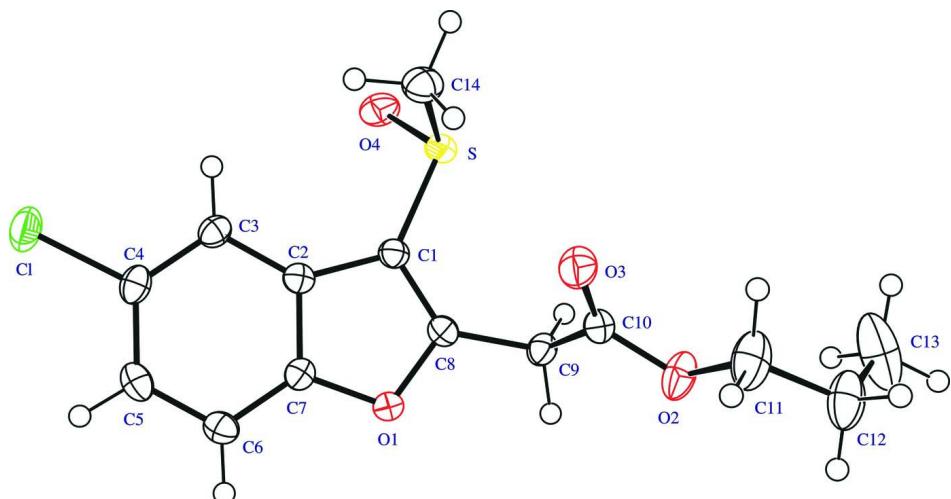
As a part of our ongoing research on the synthesis and structure of alkyl 2-(5-chloro-3-methylsulfinyl-1-benzofuran-2-yl)acetate analogues, we have described the crystal structures of ethyl 2-(5-chloro-3-methylsulfinyl-1-benzofuran-2-yl)acetate (Choi *et al.*, 2007) and methyl 2-(5-chloro-3-methylsulfinyl-1-benzofuran-2-yl)acetate (Choi *et al.*, 2008). Here we report the crystal structure of the title compound, propyl 2-(5-chloro-3-methylsulfinyl-1-benzofuran-2-yl)acetate (Fig. 1). Regardless of the lengths of the alkyl substituent arrangement of the molecules in the three crystal structures is very similar. The benzofuran unit is essentially planar, with a mean deviation of 0.013 (1) Å from the least-squares plane defined by the nine constituent atoms. The crystal packing (Fig. 2) is stabilized by aromatic π – π interactions between the benzene rings of neighbouring molecules. The Cg1···Cg1ⁱⁱ distance is 3.635 (3) Å (Cg1 is the centroid of the C2–C7 benzene ring; symmetry code as in Fig. 2). The molecular packing is further stabilized by C—H··· π interactions between the hydrogen of 12-methylene group and the furan ring of the benzofuran unit, with a C12—H12A···Cg2ⁱ separation of 2.74 Å (Table 1 and Fig. 2; Cg2 is the centroid of the C1/C2/C7/O1/C8 furan ring; symmetry code as in Fig. 2). Additionally, four different intermolecular C—H···O hydrogen bonds in the structure are observed (Table 1 & Fig. 3).

S2. Experimental

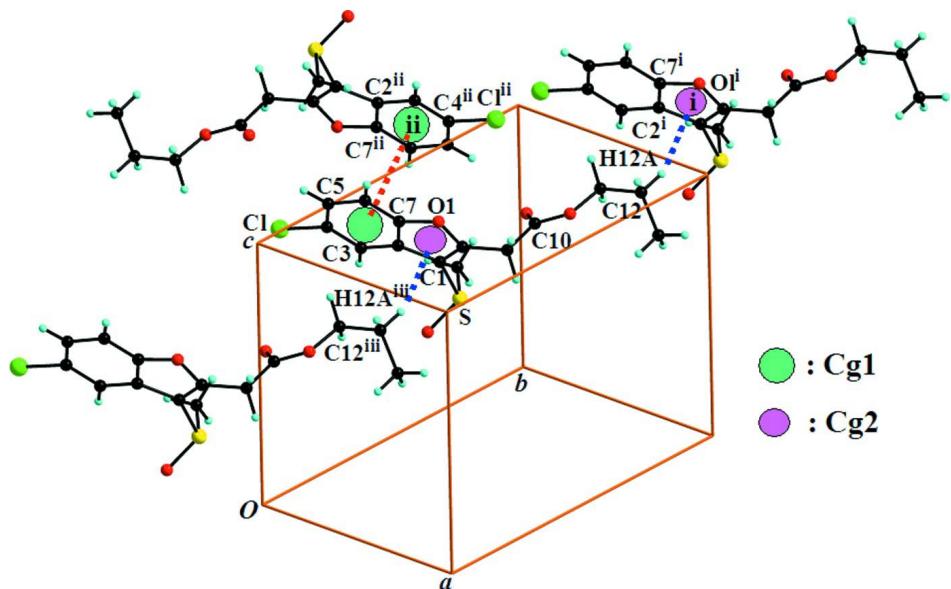
77% 3-Chloroperoxybenzoic acid (173 mg, 0.77 mmol) was added in small portions to a stirred solution of propyl 2-(5-chloro-3-methylsulfinyl-1-benzofuran-2-yl)acetate (209 mg, 0.7 mmol) in dichloromethane (20 ml) at 273 K. After being stirred for 3 h at room temperature, the mixture was washed with saturated sodium bicarbonate solution and the organic layer was separated, dried over magnesium sulfate, filtered and concentrated in vacuum. The residue was purified by column chromatography (hexane-ethyl acetate, 1:2 v/v) to afford the title compound as a colorless solid [yield 80%, m.p. 399–340 K; R_f = 0.51 (hexane-ethyl acetate, 1:2 v/v)]. Single crystals suitable for X-ray diffraction were prepared by evaporation of a solution of the title compound in benzene at room temperature. Spectroscopic analysis: ¹H NMR (CDCl₃, 400 MHz) δ 0.92 (t, J = 7.30 Hz, 3H), 1.65–1.74 (m, 2H), 3.05 (s, 3H), 4.03 (s, 2H), 4.13 (t, J = 6.92 Hz, 2H), 7.41 (d, J = 8.72 Hz, 1H), 7.50 (dd, J = 8.72 Hz and J = 1.83 Hz, 1H), 8.03 (d, J = 1.82 Hz, 1H); EI-MS 316 [M+2], 314 [M⁺].

S3. Refinement

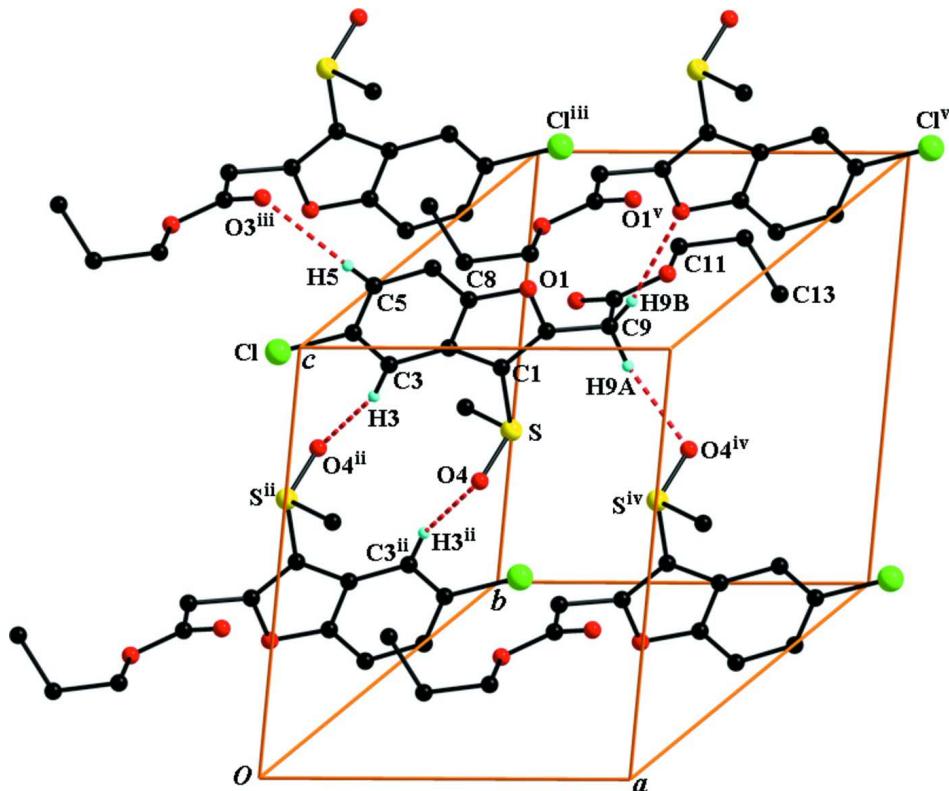
All H atoms were geometrically positioned and refined using a riding model, with C—H = 0.95 Å for the aryl, 0.99 Å for the methylene, and 0.98 Å for the methyl H atoms. U_{iso}(H) = 1.2U_{eq}(C) for the aryl and methylene H atoms, and 1.5U_{eq}(C) for the methyl H atoms.

**Figure 1**

The molecular structure of the title compound, showing displacement ellipsoids drawn at the 50% probability level.

**Figure 2**

π — π and C—H··· π interactions (dotted lines) in the title compound. Cg denotes the ring centroid. [Symmetry code: (i) x, y+1, z; (ii) -x, -y+1, -z+2; (iii) x, y-1, z.]

**Figure 3**

Intermolecular C—H···O hydrogen bonds (dotted lines) in the title compound. [Symmetry code: (ii) -x, -y+1, -z+1; (iii) -x, -y+1, -z+2; (iv) -x+1, -y+1, -z+1; (v) -x+1, -y+1, -z+2.]

Propyl 2-(5-chloro-3-methylsulfinyl-1-benzofuran-2-yl)acetate

Crystal data

$C_{14}H_{15}ClO_4S$
 $M_r = 314.77$
Triclinic, $P\bar{1}$
Hall symbol: -P 1
 $a = 8.528 (2)$ Å
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 $\beta = 81.773 (5)^\circ$
 $\gamma = 65.747 (4)^\circ$
 $V = 728.0 (4)$ Å³

$Z = 2$
 $F(000) = 328$
 $D_x = 1.436 \text{ Mg m}^{-3}$
 $Mo K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 4784 reflections
 $\theta = 2.4\text{--}28.3^\circ$
 $\mu = 0.42 \text{ mm}^{-1}$
 $T = 173$ K
Block, colorless
 $0.40 \times 0.40 \times 0.10$ mm

Data collection

Bruker SMART CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 10.0 pixels mm⁻¹
 φ and ω scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1999)
 $T_{\min} = 0.842$, $T_{\max} = 0.961$

5062 measured reflections
2468 independent reflections
2304 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.029$
 $\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.1^\circ$
 $h = -10 \rightarrow 10$
 $k = -11 \rightarrow 11$
 $l = -12 \rightarrow 12$

*Refinement*Refinement on F^2

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.037$$

$$wR(F^2) = 0.094$$

$$S = 1.07$$

2468 reflections

182 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0381P)^2 + 0.4505P]$$
$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.28 \text{ e \AA}^{-3}$$

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

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
S	0.23799 (6)	0.59458 (6)	0.53751 (4)	0.02410 (15)
Cl	-0.18859 (7)	0.22051 (7)	0.88822 (6)	0.03715 (17)
O1	0.33263 (16)	0.45103 (15)	0.93160 (12)	0.0223 (3)
O2	0.46225 (18)	0.85866 (17)	0.78142 (17)	0.0349 (4)
O3	0.21787 (18)	0.86811 (17)	0.71372 (15)	0.0338 (4)
O4	0.24182 (19)	0.46691 (18)	0.47748 (14)	0.0321 (4)
C1	0.2395 (2)	0.5204 (2)	0.71715 (18)	0.0206 (4)
C2	0.1483 (2)	0.4276 (2)	0.80462 (18)	0.0203 (4)
C3	0.0205 (2)	0.3780 (2)	0.78783 (19)	0.0228 (4)
H3	-0.0289	0.4059	0.7016	0.027*
C4	-0.0300 (2)	0.2866 (2)	0.9031 (2)	0.0249 (4)
C5	0.0395 (3)	0.2434 (2)	1.0326 (2)	0.0266 (4)
H5	0.0019	0.1786	1.1082	0.032*
C6	0.1625 (2)	0.2954 (2)	1.04984 (19)	0.0252 (4)
H6	0.2100	0.2693	1.1366	0.030*
C7	0.2134 (2)	0.3870 (2)	0.93509 (19)	0.0209 (4)
C8	0.3462 (2)	0.5306 (2)	0.79717 (18)	0.0211 (4)
C9	0.4628 (2)	0.6175 (2)	0.76992 (19)	0.0229 (4)
H9A	0.5337	0.5986	0.6860	0.028*
H9B	0.5416	0.5757	0.8470	0.028*
C10	0.3636 (2)	0.7945 (2)	0.75211 (19)	0.0244 (4)
C11	0.3854 (3)	1.0300 (3)	0.7669 (3)	0.0439 (6)
H11A	0.3053	1.0552	0.8453	0.053*
H11B	0.3204	1.0848	0.6815	0.053*
C12	0.5296 (3)	1.0819 (3)	0.7631 (3)	0.0451 (6)

H12A	0.4800	1.1959	0.7603	0.054*
H12B	0.5932	1.0244	0.8489	0.054*
C13	0.6534 (5)	1.0561 (4)	0.6453 (4)	0.0753 (10)
H13A	0.5929	1.1158	0.5596	0.090*
H13B	0.7052	0.9432	0.6478	0.090*
H13C	0.7439	1.0921	0.6509	0.090*
C14	0.0216 (3)	0.7411 (2)	0.5235 (2)	0.0318 (5)
H14A	0.0002	0.8006	0.4277	0.048*
H14B	0.0062	0.8137	0.5798	0.048*
H14C	-0.0595	0.6891	0.5554	0.048*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S	0.0256 (3)	0.0286 (3)	0.0184 (2)	-0.0127 (2)	-0.00144 (18)	-0.00292 (19)
Cl	0.0348 (3)	0.0380 (3)	0.0473 (3)	-0.0243 (2)	-0.0025 (2)	-0.0075 (2)
O1	0.0237 (7)	0.0252 (7)	0.0198 (6)	-0.0117 (6)	-0.0034 (5)	-0.0035 (5)
O2	0.0254 (8)	0.0228 (7)	0.0590 (10)	-0.0088 (6)	-0.0060 (7)	-0.0128 (7)
O3	0.0249 (8)	0.0292 (8)	0.0437 (9)	-0.0098 (6)	-0.0075 (6)	-0.0019 (7)
O4	0.0341 (8)	0.0393 (8)	0.0260 (7)	-0.0132 (7)	-0.0016 (6)	-0.0143 (6)
C1	0.0228 (9)	0.0198 (9)	0.0190 (9)	-0.0080 (7)	-0.0015 (7)	-0.0047 (7)
C2	0.0211 (9)	0.0173 (9)	0.0207 (9)	-0.0054 (7)	-0.0012 (7)	-0.0053 (7)
C3	0.0234 (10)	0.0217 (9)	0.0254 (10)	-0.0086 (8)	-0.0026 (7)	-0.0085 (8)
C4	0.0214 (10)	0.0210 (10)	0.0346 (11)	-0.0088 (8)	0.0007 (8)	-0.0105 (8)
C5	0.0268 (10)	0.0209 (10)	0.0274 (10)	-0.0088 (8)	0.0031 (8)	-0.0019 (8)
C6	0.0252 (10)	0.0244 (10)	0.0225 (9)	-0.0074 (8)	-0.0020 (8)	-0.0033 (8)
C7	0.0196 (9)	0.0192 (9)	0.0240 (9)	-0.0069 (7)	-0.0013 (7)	-0.0062 (7)
C8	0.0220 (9)	0.0196 (9)	0.0204 (9)	-0.0074 (8)	-0.0004 (7)	-0.0039 (7)
C9	0.0202 (9)	0.0244 (10)	0.0261 (9)	-0.0103 (8)	-0.0015 (7)	-0.0064 (8)
C10	0.0250 (10)	0.0269 (10)	0.0233 (9)	-0.0139 (9)	0.0003 (8)	-0.0039 (8)
C11	0.0322 (12)	0.0236 (11)	0.0760 (18)	-0.0075 (10)	-0.0016 (11)	-0.0177 (11)
C12	0.0357 (13)	0.0242 (11)	0.0760 (18)	-0.0114 (10)	-0.0014 (12)	-0.0139 (11)
C13	0.080 (2)	0.0462 (17)	0.093 (2)	-0.0344 (16)	0.0380 (19)	-0.0128 (16)
C14	0.0314 (11)	0.0270 (11)	0.0325 (11)	-0.0069 (9)	-0.0078 (9)	-0.0043 (9)

Geometric parameters (\AA , ^\circ)

S—O4	1.5038 (16)	C6—C7	1.382 (3)
S—C1	1.7656 (18)	C6—H6	0.9500
S—C14	1.797 (2)	C8—C9	1.493 (3)
Cl—C4	1.752 (2)	C9—C10	1.522 (3)
O1—C7	1.379 (2)	C9—H9A	0.9900
O1—C8	1.380 (2)	C9—H9B	0.9900
O2—C10	1.333 (2)	C11—C12	1.496 (3)
O2—C11	1.468 (3)	C11—H11A	0.9900
O3—C10	1.207 (2)	C11—H11B	0.9900
C1—C8	1.354 (3)	C12—C13	1.487 (4)
C1—C2	1.452 (3)	C12—H12A	0.9900

C2—C3	1.403 (3)	C12—H12B	0.9900
C2—C7	1.403 (3)	C13—H13A	0.9800
C3—C4	1.383 (3)	C13—H13B	0.9800
C3—H3	0.9500	C13—H13C	0.9800
C4—C5	1.407 (3)	C14—H14A	0.9800
C5—C6	1.383 (3)	C14—H14B	0.9800
C5—H5	0.9500	C14—H14C	0.9800
O4—S—C1	106.94 (9)	C10—C9—H9A	109.2
O4—S—C14	106.77 (10)	C8—C9—H9B	109.2
C1—S—C14	98.32 (9)	C10—C9—H9B	109.2
C7—O1—C8	106.32 (14)	H9A—C9—H9B	107.9
C10—O2—C11	116.86 (16)	O3—C10—O2	124.34 (19)
C8—C1—C2	107.36 (16)	O3—C10—C9	125.35 (19)
C8—C1—S	122.84 (15)	O2—C10—C9	110.29 (16)
C2—C1—S	129.51 (15)	O2—C11—C12	107.24 (18)
C3—C2—C7	119.33 (17)	O2—C11—H11A	110.3
C3—C2—C1	136.18 (17)	C12—C11—H11A	110.3
C7—C2—C1	104.48 (17)	O2—C11—H11B	110.3
C4—C3—C2	116.72 (17)	C12—C11—H11B	110.3
C4—C3—H3	121.6	H11A—C11—H11B	108.5
C2—C3—H3	121.6	C13—C12—C11	114.7 (3)
C3—C4—C5	123.34 (19)	C13—C12—H12A	108.6
C3—C4—Cl	118.78 (15)	C11—C12—H12A	108.6
C5—C4—Cl	117.87 (15)	C13—C12—H12B	108.6
C6—C5—C4	119.91 (18)	C11—C12—H12B	108.6
C6—C5—H5	120.0	H12A—C12—H12B	107.6
C4—C5—H5	120.0	C12—C13—H13A	109.5
C7—C6—C5	117.01 (18)	C12—C13—H13B	109.5
C7—C6—H6	121.5	H13A—C13—H13B	109.5
C5—C6—H6	121.5	C12—C13—H13C	109.5
O1—C7—C6	125.69 (17)	H13A—C13—H13C	109.5
O1—C7—C2	110.66 (16)	H13B—C13—H13C	109.5
C6—C7—C2	123.65 (18)	S—C14—H14A	109.5
C1—C8—O1	111.16 (17)	S—C14—H14B	109.5
C1—C8—C9	132.90 (17)	H14A—C14—H14B	109.5
O1—C8—C9	115.80 (16)	S—C14—H14C	109.5
C8—C9—C10	112.11 (15)	H14A—C14—H14C	109.5
C8—C9—H9A	109.2	H14B—C14—H14C	109.5
O4—S—C1—C8	132.24 (16)	C3—C2—C7—O1	-177.81 (15)
C14—S—C1—C8	-117.31 (17)	C1—C2—C7—O1	1.2 (2)
O4—S—C1—C2	-40.91 (19)	C3—C2—C7—C6	2.1 (3)
C14—S—C1—C2	69.54 (19)	C1—C2—C7—C6	-178.94 (17)
C8—C1—C2—C3	178.0 (2)	C2—C1—C8—O1	0.0 (2)
S—C1—C2—C3	-8.0 (3)	S—C1—C8—O1	-174.43 (12)
C8—C1—C2—C7	-0.7 (2)	C2—C1—C8—C9	-175.32 (19)
S—C1—C2—C7	173.25 (14)	S—C1—C8—C9	10.2 (3)

C7—C2—C3—C4	−1.9 (3)	C7—O1—C8—C1	0.7 (2)
C1—C2—C3—C4	179.6 (2)	C7—O1—C8—C9	176.90 (15)
C2—C3—C4—C5	0.3 (3)	C1—C8—C9—C10	72.8 (3)
C2—C3—C4—C1	−179.76 (13)	O1—C8—C9—C10	−102.43 (18)
C3—C4—C5—C6	1.2 (3)	C11—O2—C10—O3	0.8 (3)
C1—C4—C5—C6	−178.73 (15)	C11—O2—C10—C9	179.06 (18)
C4—C5—C6—C7	−1.1 (3)	C8—C9—C10—O3	−26.0 (3)
C8—O1—C7—C6	178.95 (18)	C8—C9—C10—O2	155.70 (16)
C8—O1—C7—C2	−1.16 (19)	C10—O2—C11—C12	−163.3 (2)
C5—C6—C7—O1	179.33 (17)	O2—C11—C12—C13	62.5 (3)
C5—C6—C7—C2	−0.6 (3)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C12—H12A···Cg2 ⁱ	0.99	2.74	3.666 (3)	155
C3—H3···O4 ⁱⁱ	0.95	2.44	3.353 (2)	160
C5—H5···O3 ⁱⁱⁱ	0.95	2.50	3.373 (2)	152
C9—H9A···O4 ^{iv}	0.99	2.35	3.321 (2)	166
C9—H9B···O1 ^v	0.99	2.54	3.489 (2)	161

Symmetry codes: (i) $x, y+1, z$; (ii) $-x, -y+1, -z+1$; (iii) $-x, -y+1, -z+2$; (iv) $-x+1, -y+1, -z+1$; (v) $-x+1, -y+1, -z+2$.