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Ethyl 2-(5-cyclohexyl-3-methylsulfinyl-1-benzofuran-2-yl)acetate

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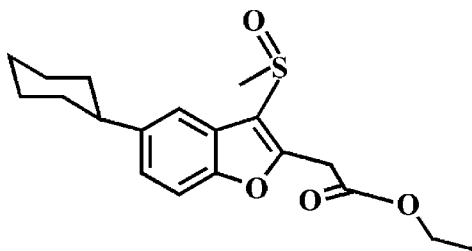
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 Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; disorder in main residue; R factor = 0.052; wR factor = 0.132; data-to-parameter ratio = 16.6.

 In the title compound, $\text{C}_{19}\text{H}_{24}\text{O}_4\text{S}$, the cyclohexyl ring adopts a chair conformation. In the crystal, molecules are linked by weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds. The O atom of the sulfinyl group is disordered over two orientations with site-occupancy factors of 0.875 (4) and 0.125 (4).

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 ethyl 2-(3-methylsulfinyl-1-benzofuran-2-yl) acetate derivatives, see: Choi *et al.* (2007a,b; 2009).


Experimental

Crystal data

 $\text{C}_{19}\text{H}_{24}\text{O}_4\text{S}$
 $M_r = 348.45$

 Monoclinic, $P2_1/c$
 $a = 16.1065$ (7) Å
 $b = 4.8485$ (2) Å
 $c = 22.3653$ (11) Å
 $\beta = 91.010$ (2)°
 $V = 1746.29$ (14) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.21$ mm⁻¹
 $T = 173$ K
 $0.28 \times 0.20 \times 0.18$ mm

Data collection

 Bruker SMART APEXII CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2009)
 $T_{\min} = 0.945$, $T_{\max} = 0.964$

 15388 measured reflections
 3804 independent reflections
 2917 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.049$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.132$
 $S = 1.02$
 3804 reflections
 229 parameters

 67 restraints
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.56$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.38$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C14}-\text{H14B}\cdots\text{O4A}^i$	0.99	2.60	3.479 (3)	149
$\text{C15}-\text{H15B}\cdots\text{O2}^{\text{ii}}$	0.99	2.56	3.437 (3)	147

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

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.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BV2187).

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supporting information

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Ethyl 2-(5-cyclohexyl-3-methylsulfinyl-1-benzofuran-2-yl)acetate

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S1. Comment

Recently, many compounds containing a benzofuran moiety have drawn much attention owing to their valuable 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 ethyl 2-(3-methylsulfinyl-1-benzofuran-2-yl)acetate analogues (Choi *et al.*, 2007*a,b*; 2009), we report herein on the crystal structure of the title compound

In the title molecule (Fig. 1), the benzofuran unit is essentially planar, with a mean deviation of 0.011 (2) Å from the least-squares plane defined by the nine constituent atoms. The cyclohexyl ring is in the chair form. The O atom of sulfinyl group is disordered over two positions with site-occupancy factors, from refinement of 0.875 (3) (part A) and 0.125 (4) (part B). The molecular packing (Fig. 2) is stabilized by weak intermolecular C—H...O hydrogen bonds; the first one between a cyclohexyl H atom and the O atom of the sulfinyl group (Table 1; C14—H14B...O4Aⁱ), and the second one between an H atom of the benzylic methylene group and the O atom of the carbonyl group (Table 1; C15—H15B...O2ⁱⁱ).

S2. Experimental

77% 3-chloroperoxybenzoic acid (224 mg, 1.0 mmol) was added in small portions to a stirred solution of ethyl 2-(5-cyclohexyl-3-methylsulfinyl-1-benzofuran-2-yl)acetate (298 mg, 0.9 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, 1:2 v/v) to afford the title compound as a colorless solid [yield 70%, m.p. 360–361 K; $R_f = 0.41$ (hexane–ethyl acetate, 1:2 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. The O atom of sulfinyl group is disordered over two positions with site-occupancy factors, from refinement of 0.875 (4) (part A) and 0.125 (4) (part B). The S—O distances were restrained to 0.001 Å using command SADI and DELU.

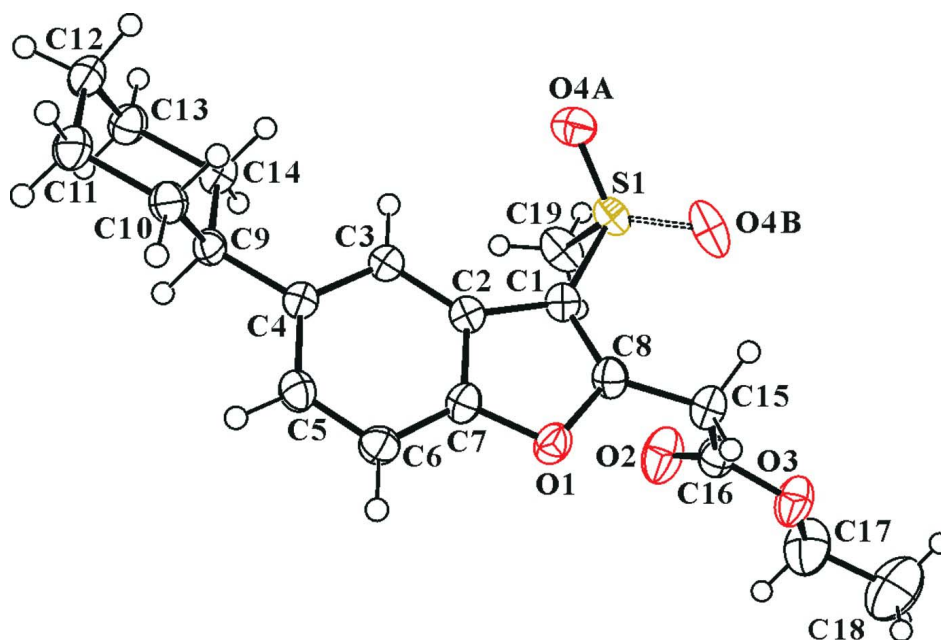


Figure 1

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. The O atom of sulfinyl group is disordered over two positions with site-occupancy factors, from refinement of 0.875 (3) (part A) and 0.125 (4) (part B).

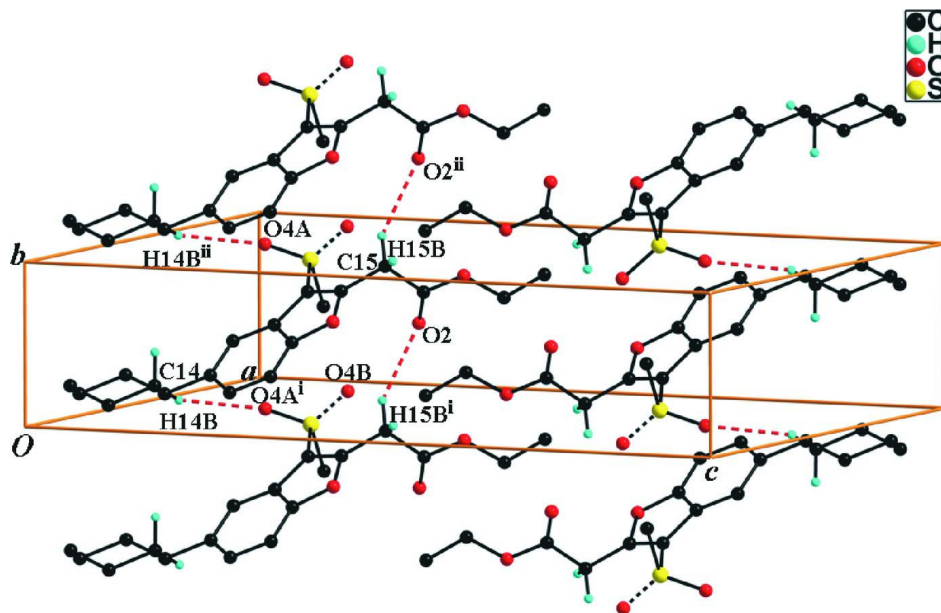


Figure 2

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

Ethyl 2-(5-cyclohexyl-3-methylsulfinyl-1-benzofuran-2-yl)acetate

Crystal data

$C_{19}H_{24}O_4S$	$F(000) = 744$
$M_r = 348.45$	$D_x = 1.325 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Hall symbol: $-P 2ybc$	Cell parameters from 3275 reflections
$a = 16.1065 (7) \text{ \AA}$	$\theta = 2.2\text{--}26.5^\circ$
$b = 4.8485 (2) \text{ \AA}$	$\mu = 0.21 \text{ mm}^{-1}$
$c = 22.3653 (11) \text{ \AA}$	$T = 173 \text{ K}$
$\beta = 91.010 (2)^\circ$	Block, colourless
$V = 1746.29 (14) \text{ \AA}^3$	$0.28 \times 0.20 \times 0.18 \text{ mm}$
$Z = 4$	

Data collection

Bruker SMART APEXII CCD diffractometer	15388 measured reflections
Radiation source: rotating anode	3804 independent reflections
Graphite multilayer monochromator	2917 reflections with $I > 2\sigma(I)$
Detector resolution: $10.0 \text{ pixels mm}^{-1}$	$R_{\text{int}} = 0.049$
φ and ω scans	$\theta_{\text{max}} = 27.0^\circ$, $\theta_{\text{min}} = 1.3^\circ$
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2009)	$h = -20 \rightarrow 19$
$T_{\text{min}} = 0.945$, $T_{\text{max}} = 0.964$	$k = -6 \rightarrow 6$
	$l = -28 \rightarrow 28$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.052$	H-atom parameters constrained
$wR(F^2) = 0.132$	$w = 1/[\sigma^2(F_o^2) + (0.0537P)^2 + 1.3587P]$
$S = 1.02$	where $P = (F_o^2 + 2F_c^2)/3$
3804 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
229 parameters	$\Delta\rho_{\text{max}} = 0.56 \text{ e \AA}^{-3}$
67 restraints	$\Delta\rho_{\text{min}} = -0.38 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
S1	0.36614 (3)	0.95799 (12)	0.29116 (3)	0.03096 (17)	
O1	0.18470 (9)	0.6531 (3)	0.38601 (6)	0.0325 (4)	
O2	0.38323 (12)	0.5969 (4)	0.44371 (8)	0.0532 (5)	
O3	0.36351 (11)	0.9044 (4)	0.51602 (7)	0.0443 (5)	

O4A	0.35467 (11)	1.0448 (4)	0.22886 (8)	0.0392 (6)	0.875 (4)
O4B	0.3933 (9)	1.160 (2)	0.3361 (5)	0.057 (4)	0.125 (4)
C1	0.27598 (12)	0.7963 (4)	0.31809 (9)	0.0266 (5)	
C2	0.22021 (12)	0.6069 (4)	0.28840 (9)	0.0253 (4)	
C3	0.21169 (12)	0.5003 (4)	0.23082 (9)	0.0262 (5)	
H3	0.2479	0.5575	0.2001	0.031*	
C4	0.14931 (12)	0.3085 (4)	0.21906 (9)	0.0257 (5)	
C5	0.09692 (13)	0.2284 (5)	0.26516 (10)	0.0316 (5)	
H5	0.0548	0.0962	0.2567	0.038*	
C6	0.10392 (13)	0.3335 (5)	0.32258 (10)	0.0327 (5)	
H6	0.0679	0.2773	0.3535	0.039*	
C7	0.16600 (13)	0.5242 (5)	0.33243 (9)	0.0283 (5)	
C8	0.25217 (13)	0.8178 (5)	0.37491 (10)	0.0297 (5)	
C9	0.13521 (13)	0.1884 (4)	0.15724 (9)	0.0269 (5)	
H9	0.1225	-0.0119	0.1624	0.032*	
C10	0.05931 (13)	0.3205 (5)	0.12640 (9)	0.0308 (5)	
H10A	0.0100	0.2924	0.1515	0.037*	
H10B	0.0686	0.5216	0.1227	0.037*	
C11	0.04212 (14)	0.1998 (6)	0.06457 (10)	0.0379 (6)	
H11A	0.0255	0.0042	0.0686	0.045*	
H11B	-0.0046	0.3007	0.0453	0.045*	
C12	0.11783 (14)	0.2186 (5)	0.02536 (10)	0.0362 (5)	
H12A	0.1304	0.4148	0.0173	0.043*	
H12B	0.1058	0.1270	-0.0134	0.043*	
C13	0.19283 (15)	0.0828 (5)	0.05528 (10)	0.0358 (5)	
H13A	0.2420	0.1071	0.0299	0.043*	
H13B	0.1825	-0.1175	0.0595	0.043*	
C14	0.21039 (13)	0.2080 (5)	0.11680 (9)	0.0317 (5)	
H14A	0.2261	0.4041	0.1121	0.038*	
H14B	0.2579	0.1106	0.1359	0.038*	
C15	0.28665 (14)	0.9685 (5)	0.42765 (10)	0.0338 (5)	
H15A	0.2408	1.0187	0.4544	0.041*	
H15B	0.3131	1.1413	0.4141	0.041*	
C16	0.34984 (14)	0.7987 (5)	0.46210 (10)	0.0347 (5)	
C17	0.42421 (19)	0.7598 (7)	0.55380 (13)	0.0591 (8)	
H17A	0.4054	0.5687	0.5612	0.071*	
H17B	0.4784	0.7519	0.5337	0.071*	
C18	0.4327 (2)	0.9083 (9)	0.61010 (14)	0.0775 (11)	
H18A	0.4544	1.0937	0.6026	0.116*	
H18B	0.4711	0.8088	0.6369	0.116*	
H18C	0.3783	0.9225	0.6288	0.116*	
C19	0.42883 (14)	0.6559 (5)	0.28894 (12)	0.0403 (6)	
H19A	0.4039	0.5228	0.2610	0.060*	
H19B	0.4325	0.5739	0.3290	0.060*	
H19C	0.4846	0.7046	0.2757	0.060*	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0292 (3)	0.0237 (3)	0.0402 (3)	-0.0018 (2)	0.0030 (2)	0.0002 (3)
O1	0.0286 (8)	0.0414 (9)	0.0276 (8)	-0.0026 (7)	0.0032 (6)	-0.0066 (7)
O2	0.0576 (12)	0.0484 (11)	0.0531 (11)	0.0198 (10)	-0.0138 (9)	-0.0149 (10)
O3	0.0490 (10)	0.0465 (11)	0.0368 (9)	0.0081 (9)	-0.0149 (8)	-0.0089 (8)
O4A	0.0363 (10)	0.0412 (12)	0.0402 (11)	-0.0020 (9)	0.0010 (8)	0.0072 (9)
O4B	0.058 (9)	0.030 (7)	0.083 (9)	-0.013 (6)	-0.001 (8)	-0.008 (7)
C1	0.0227 (10)	0.0267 (11)	0.0302 (10)	0.0026 (9)	-0.0007 (8)	-0.0023 (9)
C2	0.0211 (9)	0.0254 (11)	0.0294 (10)	0.0025 (9)	0.0000 (8)	-0.0003 (9)
C3	0.0242 (10)	0.0272 (11)	0.0271 (10)	0.0014 (9)	-0.0001 (8)	-0.0002 (9)
C4	0.0251 (10)	0.0253 (11)	0.0266 (10)	0.0027 (9)	-0.0009 (8)	-0.0011 (9)
C5	0.0279 (11)	0.0319 (12)	0.0350 (11)	-0.0035 (10)	-0.0028 (9)	-0.0014 (10)
C6	0.0267 (10)	0.0405 (13)	0.0312 (11)	-0.0049 (10)	0.0050 (9)	-0.0006 (11)
C7	0.0251 (10)	0.0318 (12)	0.0279 (10)	0.0028 (9)	-0.0016 (8)	-0.0043 (10)
C8	0.0245 (10)	0.0308 (12)	0.0336 (11)	-0.0002 (9)	0.0002 (9)	-0.0048 (10)
C9	0.0313 (11)	0.0222 (11)	0.0270 (10)	-0.0006 (9)	-0.0023 (8)	-0.0014 (9)
C10	0.0248 (10)	0.0342 (12)	0.0335 (11)	-0.0021 (10)	-0.0017 (9)	-0.0003 (10)
C11	0.0323 (12)	0.0469 (15)	0.0341 (12)	-0.0067 (11)	-0.0080 (9)	0.0010 (11)
C12	0.0384 (12)	0.0431 (14)	0.0270 (11)	-0.0051 (11)	-0.0041 (9)	-0.0024 (11)
C13	0.0416 (13)	0.0358 (13)	0.0301 (11)	0.0010 (11)	0.0030 (10)	-0.0049 (11)
C14	0.0297 (11)	0.0350 (12)	0.0305 (11)	0.0053 (10)	-0.0016 (9)	-0.0043 (10)
C15	0.0338 (11)	0.0365 (13)	0.0309 (11)	0.0016 (10)	-0.0017 (9)	-0.0073 (10)
C16	0.0322 (12)	0.0371 (13)	0.0346 (12)	-0.0027 (11)	-0.0018 (9)	-0.0044 (11)
C17	0.0637 (18)	0.0570 (18)	0.0557 (17)	0.0124 (16)	-0.0278 (15)	-0.0027 (15)
C18	0.077 (2)	0.102 (3)	0.0525 (18)	0.025 (2)	-0.0274 (16)	-0.0068 (19)
C19	0.0274 (11)	0.0317 (13)	0.0620 (16)	0.0040 (10)	0.0077 (11)	0.0001 (12)

Geometric parameters (\AA , $^\circ$)

S1—O4B	1.463 (2)	C10—H10A	0.9900
S1—O4A	1.4641 (19)	C10—H10B	0.9900
S1—C1	1.765 (2)	C11—C12	1.517 (3)
S1—C19	1.780 (2)	C11—H11A	0.9900
O1—C8	1.375 (3)	C11—H11B	0.9900
O1—C7	1.380 (2)	C12—C13	1.520 (3)
O2—C16	1.193 (3)	C12—H12A	0.9900
O3—C16	1.325 (3)	C12—H12B	0.9900
O3—C17	1.460 (3)	C13—C14	1.526 (3)
C1—C8	1.338 (3)	C13—H13A	0.9900
C1—C2	1.439 (3)	C13—H13B	0.9900
C2—C7	1.387 (3)	C14—H14A	0.9900
C2—C3	1.392 (3)	C14—H14B	0.9900
C3—C4	1.391 (3)	C15—C16	1.510 (3)
C3—H3	0.9500	C15—H15A	0.9900
C4—C5	1.399 (3)	C15—H15B	0.9900
C4—C9	1.514 (3)	C17—C18	1.455 (4)

C5—C6	1.385 (3)	C17—H17A	0.9900
C5—H5	0.9500	C17—H17B	0.9900
C6—C7	1.377 (3)	C18—H18A	0.9800
C6—H6	0.9500	C18—H18B	0.9800
C8—C15	1.487 (3)	C18—H18C	0.9800
C9—C14	1.527 (3)	C19—H19A	0.9800
C9—C10	1.533 (3)	C19—H19B	0.9800
C9—H9	1.0000	C19—H19C	0.9800
C10—C11	1.523 (3)		
O4B—S1—O4A	119.5 (6)	C10—C11—H11B	109.3
O4B—S1—C1	107.5 (6)	H11A—C11—H11B	108.0
O4A—S1—C1	111.19 (10)	C11—C12—C13	111.10 (19)
O4B—S1—C19	114.0 (6)	C11—C12—H12A	109.4
O4A—S1—C19	105.79 (12)	C13—C12—H12A	109.4
C1—S1—C19	96.58 (11)	C11—C12—H12B	109.4
C8—O1—C7	105.52 (16)	C13—C12—H12B	109.4
C16—O3—C17	116.1 (2)	H12A—C12—H12B	108.0
C8—C1—C2	107.52 (19)	C12—C13—C14	110.96 (19)
C8—C1—S1	122.73 (17)	C12—C13—H13A	109.4
C2—C1—S1	129.54 (16)	C14—C13—H13A	109.4
C7—C2—C3	119.77 (19)	C12—C13—H13B	109.4
C7—C2—C1	104.63 (18)	C14—C13—H13B	109.4
C3—C2—C1	135.6 (2)	H13A—C13—H13B	108.0
C4—C3—C2	118.8 (2)	C13—C14—C9	111.91 (18)
C4—C3—H3	120.6	C13—C14—H14A	109.2
C2—C3—H3	120.6	C9—C14—H14A	109.2
C3—C4—C5	119.3 (2)	C13—C14—H14B	109.2
C3—C4—C9	121.74 (19)	C9—C14—H14B	109.2
C5—C4—C9	118.92 (19)	H14A—C14—H14B	107.9
C6—C5—C4	122.8 (2)	C8—C15—C16	112.1 (2)
C6—C5—H5	118.6	C8—C15—H15A	109.2
C4—C5—H5	118.6	C16—C15—H15A	109.2
C7—C6—C5	116.3 (2)	C8—C15—H15B	109.2
C7—C6—H6	121.9	C16—C15—H15B	109.2
C5—C6—H6	121.9	H15A—C15—H15B	107.9
C6—C7—O1	126.1 (2)	O2—C16—O3	124.2 (2)
C6—C7—C2	123.0 (2)	O2—C16—C15	125.1 (2)
O1—C7—C2	110.79 (18)	O3—C16—C15	110.7 (2)
C1—C8—O1	111.53 (18)	C18—C17—O3	108.3 (2)
C1—C8—C15	133.0 (2)	C18—C17—H17A	110.0
O1—C8—C15	115.31 (19)	O3—C17—H17A	110.0
C4—C9—C14	114.16 (17)	C18—C17—H17B	110.0
C4—C9—C10	110.85 (17)	O3—C17—H17B	110.0
C14—C9—C10	110.00 (18)	H17A—C17—H17B	108.4
C4—C9—H9	107.2	C17—C18—H18A	109.5
C14—C9—H9	107.2	C17—C18—H18B	109.5
C10—C9—H9	107.2	H18A—C18—H18B	109.5

C11—C10—C9	112.27 (19)	C17—C18—H18C	109.5
C11—C10—H10A	109.2	H18A—C18—H18C	109.5
C9—C10—H10A	109.2	H18B—C18—H18C	109.5
C11—C10—H10B	109.2	S1—C19—H19A	109.5
C9—C10—H10B	109.2	S1—C19—H19B	109.5
H10A—C10—H10B	107.9	H19A—C19—H19B	109.5
C12—C11—C10	111.51 (18)	S1—C19—H19C	109.5
C12—C11—H11A	109.3	H19A—C19—H19C	109.5
C10—C11—H11A	109.3	H19B—C19—H19C	109.5
C12—C11—H11B	109.3		
O4B—S1—C1—C8	-13.8 (7)	C2—C1—C8—O1	0.8 (2)
O4A—S1—C1—C8	-146.31 (19)	S1—C1—C8—O1	-174.34 (14)
C19—S1—C1—C8	103.9 (2)	C2—C1—C8—C15	176.7 (2)
O4B—S1—C1—C2	172.3 (7)	S1—C1—C8—C15	1.6 (4)
O4A—S1—C1—C2	39.7 (2)	C7—O1—C8—C1	-0.3 (2)
C19—S1—C1—C2	-70.1 (2)	C7—O1—C8—C15	-176.97 (19)
C8—C1—C2—C7	-0.9 (2)	C3—C4—C9—C14	21.3 (3)
S1—C1—C2—C7	173.75 (17)	C5—C4—C9—C14	-160.1 (2)
C8—C1—C2—C3	179.9 (2)	C3—C4—C9—C10	-103.6 (2)
S1—C1—C2—C3	-5.4 (4)	C5—C4—C9—C10	75.0 (2)
C7—C2—C3—C4	-1.0 (3)	C4—C9—C10—C11	-178.72 (18)
C1—C2—C3—C4	178.1 (2)	C14—C9—C10—C11	54.1 (2)
C2—C3—C4—C5	0.1 (3)	C9—C10—C11—C12	-54.8 (3)
C2—C3—C4—C9	178.68 (19)	C10—C11—C12—C13	55.2 (3)
C3—C4—C5—C6	0.5 (3)	C11—C12—C13—C14	-55.9 (3)
C9—C4—C5—C6	-178.2 (2)	C12—C13—C14—C9	56.5 (3)
C4—C5—C6—C7	0.0 (3)	C4—C9—C14—C13	179.72 (18)
C5—C6—C7—O1	-179.3 (2)	C10—C9—C14—C13	-54.9 (2)
C5—C6—C7—C2	-1.0 (3)	C1—C8—C15—C16	-88.6 (3)
C8—O1—C7—C6	178.2 (2)	O1—C8—C15—C16	87.2 (2)
C8—O1—C7—C2	-0.3 (2)	C17—O3—C16—O2	0.5 (4)
C3—C2—C7—C6	1.5 (3)	C17—O3—C16—C15	-179.5 (2)
C1—C2—C7—C6	-177.8 (2)	C8—C15—C16—O2	16.1 (3)
C3—C2—C7—O1	-179.89 (18)	C8—C15—C16—O3	-163.9 (2)
C1—C2—C7—O1	0.8 (2)	C16—O3—C17—C18	178.5 (3)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C14—H14B \cdots O4A ⁱ	0.99	2.60	3.479 (3)	149
C15—H15B \cdots O2 ⁱⁱ	0.99	2.56	3.437 (3)	147

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