

1-(4-Bromophenylsulfinyl)-2-methyl-naphtho[2,1-*b*]furan

Hong Dae Choi,^a Pil Ja Seo^a and Uk Lee^{b*}

^aDepartment of Chemistry, Dongeui University, San 24 Kaya-dong Busanjin-gu, Busan 614-714, Republic of Korea, and ^bDepartment of Chemistry, Pukyong National University, 599-1 Daeyeon 3-dong, Nam-gu, Busan 608-737, Republic of Korea

Correspondence e-mail: uklee@pknu.ac.kr

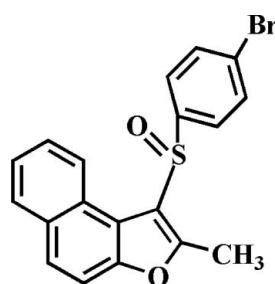
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Key indicators: single-crystal X-ray study; $T = 173\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.030; wR factor = 0.080; data-to-parameter ratio = 18.4.

In the title compound, $\text{C}_{19}\text{H}_{13}\text{BrO}_2\text{S}$, the 4-bromophenyl ring makes a dihedral angle of $83.75(4)^\circ$ with the mean plane of the naphthofuran fragment [r.m.s. deviation = $0.024(2)\text{ \AA}$]. 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 weak $\pi-\pi$ interactions between the central naphthofuran benzene rings of neighbouring molecules [centroid–centroid distance = $3.483(2)\text{ \AA}$, interplanar distance = $3.416(2)\text{ \AA}$ and slippage = $0.680(2)\text{ \AA}$].

Related literature

For background information and the crystal structures of related compounds, see: Choi *et al.* (2007, 2012).



Experimental

Crystal data

$\text{C}_{19}\text{H}_{13}\text{BrO}_2\text{S}$	$\gamma = 75.672(1)^\circ$
$M_r = 385.26$	$V = 778.74(2)\text{ \AA}^3$
Triclinic, $P\bar{1}$	$Z = 2$
$a = 8.7124(1)\text{ \AA}$	Mo $K\alpha$ radiation
$b = 9.4857(1)\text{ \AA}$	$\mu = 2.78\text{ mm}^{-1}$
$c = 10.2898(1)\text{ \AA}$	$T = 173\text{ K}$
$\alpha = 82.883(1)^\circ$	$0.33 \times 0.29 \times 0.27\text{ mm}$
$\beta = 71.126(1)^\circ$	

Data collection

Bruker SMART APEXII CCD diffractometer	14577 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2009)	3853 independent reflections
$T_{\min} = 0.462$, $T_{\max} = 0.517$	3461 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.031$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.030$	209 parameters
$wR(F^2) = 0.080$	H-atom parameters constrained
$S = 1.05$	$\Delta\rho_{\max} = 0.62\text{ e \AA}^{-3}$
3853 reflections	$\Delta\rho_{\min} = -0.60\text{ e \AA}^{-3}$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C19}-\text{H19}\cdots\text{O2}^i$	0.95	2.35	3.221 (2)	152

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

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: SJ5222).

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supporting information

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1-(4-Bromophenylsulfinyl)-2-methylnaphtho[2,1-*b*]furan

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S1. Comment

As a part of our ongoing study of 2-methylnaphtho[2,1-*b*]furan derivatives containing 1-phenylsulfinyl (Choi *et al.*, 2007) and 1-(4-methylphenylsulfinyl) (Choi *et al.*, 2012) substituents, we report herein the crystal structure of the title compound.

In the title molecule (Fig. 1), the naphthofuran unit is essentially planar, with a mean deviation of 0.024 (2) Å from the least-squares plane defined by the thirteen constituent atoms. The dihedral angle between the 4-bromophenyl ring and the mean plane of the naphthofuran fragment is 83.75 (4)°. 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 weak π — π interactions between the central naphthofuran benzene rings of neighbouring molecules, with a Cg···Cgⁱⁱ distance of 3.483 (2) Å and an interplanar distance of 3.416 (2) Å resulting in a slippage of 0.680 (2) Å (Fig. 2, Cg is the centroid of the C2/C3/C8/C9/C10/C11 benzene ring).

S2. Experimental

77% 3-Chloroperoxybenzoic acid (202 mg, 0.9 mmol) was added in small portions to a stirred solution of 1-(4-bromophenylsulfanyl)-2-methylnaphtho [2,1-*b*]furan (295 mg, 0.8 mmol) in dichloromethane (40 mL) at 273 K. After being stirred at room temperature for 5 h, 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, 2:1 v/v) to afford the title compound as a colorless solid [yield 74%, m.p. 466–467 K; R_f = 0.59 (hexane–ethyl acetate, 2:1 v/v)]. 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 the 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 the methyl H atoms were optimized rotationally.

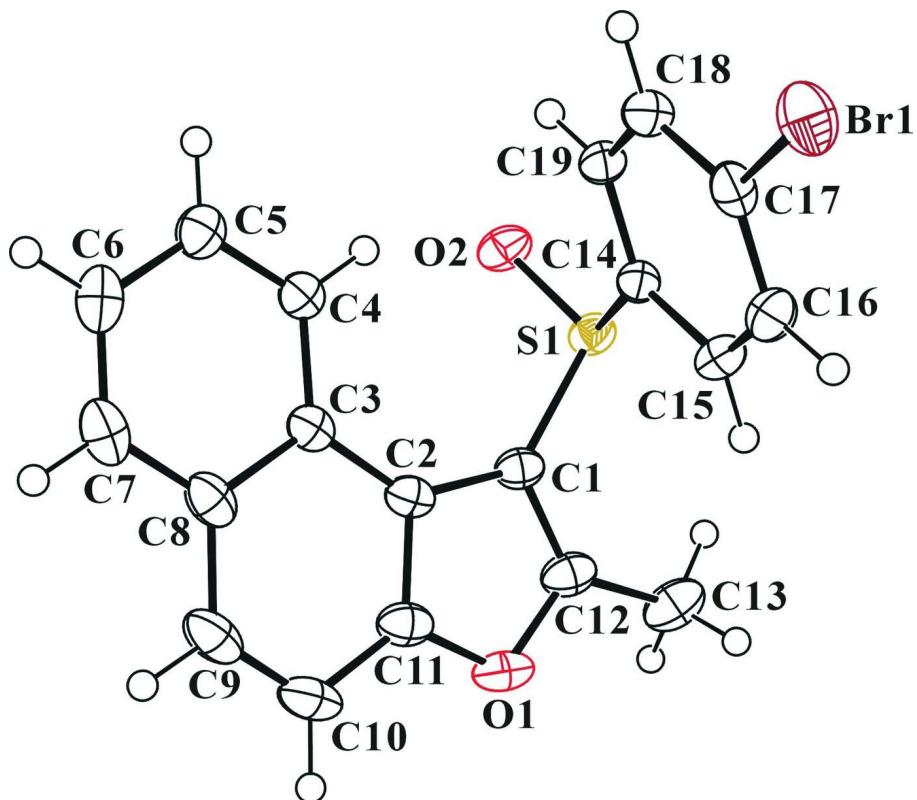
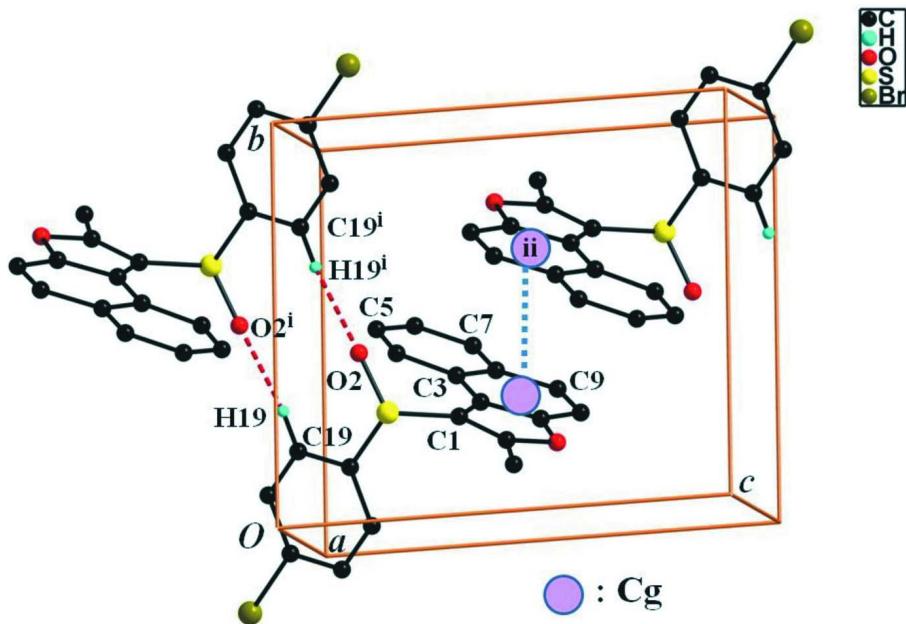


Figure 1

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 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 non-participating in hydrogen-bonding were omitted for clarity. [Symmetry codes: (i) $-x + 1, -y + 1, -z$; (ii) $-x + 1, -y + 1, -z + 1$.]

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Crystal data

$C_{19}H_{13}BrO_2S$
 $M_r = 385.26$
Triclinic, $P\bar{1}$
Hall symbol: -P 1
 $a = 8.7124 (1)$ Å
 $b = 9.4857 (1)$ Å
 $c = 10.2898 (1)$ Å
 $\alpha = 82.883 (1)^\circ$
 $\beta = 71.126 (1)^\circ$
 $\gamma = 75.672 (1)^\circ$
 $V = 778.74 (2)$ Å³

$Z = 2$
 $F(000) = 388$
 $D_x = 1.643$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 7454 reflections
 $\theta = 2.5\text{--}28.3^\circ$
 $\mu = 2.78$ mm⁻¹
 $T = 173$ K
Block, colourless
 $0.33 \times 0.29 \times 0.27$ 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)
 $T_{\min} = 0.462$, $T_{\max} = 0.517$

14577 measured reflections
3853 independent reflections
3461 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.031$
 $\theta_{\max} = 28.3^\circ$, $\theta_{\min} = 2.1^\circ$
 $h = -11 \rightarrow 11$
 $k = -12 \rightarrow 12$
 $l = -13 \rightarrow 13$

*Refinement*Refinement on F^2

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.030$$

$$wR(F^2) = 0.080$$

$$S = 1.05$$

3853 reflections

209 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

$$w = 1/[\sigma^2(F_o^2) + (0.0424P)^2 + 0.3115P]$$

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

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

$$\Delta\rho_{\max} = 0.62 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.60 \text{ e \AA}^{-3}$$

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	1.00152 (2)	-0.12635 (2)	-0.16983 (2)	0.03951 (8)
S1	0.35059 (5)	0.29100 (5)	0.20858 (4)	0.02531 (10)
O1	0.39380 (18)	0.19036 (15)	0.57662 (13)	0.0339 (3)
O2	0.34216 (16)	0.44277 (15)	0.14815 (14)	0.0308 (3)
C1	0.4076 (2)	0.27335 (19)	0.36023 (17)	0.0248 (3)
C2	0.5387 (2)	0.31553 (19)	0.39413 (17)	0.0241 (3)
C3	0.6643 (2)	0.39490 (19)	0.32814 (17)	0.0236 (3)
C4	0.6876 (2)	0.46220 (19)	0.19438 (17)	0.0243 (3)
H4	0.6154	0.4567	0.1439	0.029*
C5	0.8127 (2)	0.5349 (2)	0.1373 (2)	0.0295 (4)
H5	0.8265	0.5795	0.0475	0.035*
C6	0.9214 (2)	0.5446 (2)	0.2102 (2)	0.0357 (4)
H6	1.0098	0.5933	0.1689	0.043*
C7	0.8993 (3)	0.4836 (2)	0.3402 (2)	0.0357 (4)
H7	0.9721	0.4918	0.3892	0.043*
C8	0.7706 (2)	0.4086 (2)	0.40403 (19)	0.0295 (4)
C9	0.7452 (3)	0.3495 (2)	0.5420 (2)	0.0357 (4)
H9	0.8165	0.3611	0.5910	0.043*
C10	0.6220 (3)	0.2772 (2)	0.60542 (19)	0.0353 (4)
H10	0.6045	0.2392	0.6976	0.042*
C11	0.5225 (2)	0.2618 (2)	0.52789 (18)	0.0293 (4)
C12	0.3248 (2)	0.2003 (2)	0.47280 (19)	0.0301 (4)
C13	0.1834 (3)	0.1278 (3)	0.5032 (2)	0.0419 (5)
H13A	0.1309	0.1541	0.4299	0.063*
H13B	0.1017	0.1594	0.5912	0.063*

H13C	0.2243	0.0219	0.5087	0.063*
C14	0.5374 (2)	0.17880 (19)	0.10434 (17)	0.0234 (3)
C15	0.5817 (3)	0.0335 (2)	0.14588 (19)	0.0313 (4)
H15	0.5173	-0.0031	0.2308	0.038*
C16	0.7191 (3)	-0.0578 (2)	0.0640 (2)	0.0333 (4)
H16	0.7497	-0.1574	0.0916	0.040*
C17	0.8116 (2)	-0.0017 (2)	-0.05890 (19)	0.0283 (4)
C18	0.7683 (2)	0.1426 (2)	-0.10104 (18)	0.0286 (4)
H18	0.8335	0.1792	-0.1855	0.034*
C19	0.6289 (2)	0.2341 (2)	-0.01936 (17)	0.0259 (3)
H19	0.5970	0.3332	-0.0480	0.031*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.03626 (13)	0.03675 (14)	0.04460 (13)	-0.00190 (9)	-0.01114 (9)	-0.01442 (9)
S1	0.0229 (2)	0.0272 (2)	0.0268 (2)	-0.00701 (17)	-0.01025 (16)	0.00538 (16)
O1	0.0371 (7)	0.0337 (8)	0.0231 (6)	-0.0025 (6)	-0.0056 (5)	0.0069 (5)
O2	0.0278 (6)	0.0283 (7)	0.0344 (7)	-0.0035 (5)	-0.0129 (5)	0.0092 (5)
C1	0.0234 (8)	0.0242 (9)	0.0231 (8)	-0.0015 (7)	-0.0060 (6)	0.0016 (6)
C2	0.0258 (8)	0.0217 (8)	0.0212 (7)	0.0024 (6)	-0.0078 (6)	-0.0021 (6)
C3	0.0239 (8)	0.0210 (8)	0.0239 (8)	0.0013 (6)	-0.0079 (6)	-0.0047 (6)
C4	0.0247 (8)	0.0214 (8)	0.0261 (8)	-0.0015 (7)	-0.0089 (6)	-0.0025 (6)
C5	0.0287 (9)	0.0254 (9)	0.0324 (9)	-0.0046 (7)	-0.0069 (7)	-0.0034 (7)
C6	0.0304 (9)	0.0314 (11)	0.0481 (11)	-0.0095 (8)	-0.0112 (8)	-0.0084 (8)
C7	0.0330 (10)	0.0342 (11)	0.0458 (11)	-0.0035 (8)	-0.0194 (9)	-0.0117 (8)
C8	0.0307 (9)	0.0272 (10)	0.0318 (9)	0.0023 (7)	-0.0147 (7)	-0.0092 (7)
C9	0.0418 (11)	0.0343 (11)	0.0346 (10)	0.0050 (8)	-0.0240 (9)	-0.0080 (8)
C10	0.0452 (11)	0.0334 (11)	0.0232 (8)	0.0054 (9)	-0.0152 (8)	-0.0022 (7)
C11	0.0313 (9)	0.0271 (9)	0.0240 (8)	0.0020 (7)	-0.0077 (7)	-0.0001 (7)
C12	0.0284 (9)	0.0273 (10)	0.0272 (8)	-0.0004 (7)	-0.0040 (7)	0.0026 (7)
C13	0.0353 (11)	0.0409 (12)	0.0419 (11)	-0.0124 (9)	-0.0024 (9)	0.0092 (9)
C14	0.0276 (8)	0.0228 (9)	0.0229 (8)	-0.0076 (7)	-0.0114 (6)	0.0020 (6)
C15	0.0411 (10)	0.0259 (10)	0.0266 (8)	-0.0120 (8)	-0.0092 (8)	0.0058 (7)
C16	0.0460 (11)	0.0202 (9)	0.0339 (9)	-0.0064 (8)	-0.0143 (8)	0.0017 (7)
C17	0.0314 (9)	0.0274 (9)	0.0300 (9)	-0.0063 (7)	-0.0132 (7)	-0.0061 (7)
C18	0.0326 (9)	0.0314 (10)	0.0236 (8)	-0.0103 (8)	-0.0096 (7)	0.0018 (7)
C19	0.0325 (9)	0.0229 (9)	0.0245 (8)	-0.0073 (7)	-0.0130 (7)	0.0044 (6)

Geometric parameters (\AA , ^\circ)

Br1—C17	1.8932 (19)	C8—C9	1.428 (3)
S1—O2	1.4899 (13)	C9—C10	1.360 (3)
S1—C1	1.7621 (17)	C9—H9	0.9500
S1—C14	1.7966 (19)	C10—C11	1.395 (3)
O1—C12	1.369 (2)	C10—H10	0.9500
O1—C11	1.378 (2)	C12—C13	1.486 (3)
C1—C12	1.360 (2)	C13—H13A	0.9800

C1—C2	1.451 (2)	C13—H13B	0.9800
C2—C11	1.382 (2)	C13—H13C	0.9800
C2—C3	1.424 (2)	C14—C19	1.384 (2)
C3—C4	1.417 (2)	C14—C15	1.389 (3)
C3—C8	1.427 (2)	C15—C16	1.381 (3)
C4—C5	1.366 (2)	C15—H15	0.9500
C4—H4	0.9500	C16—C17	1.385 (3)
C5—C6	1.411 (3)	C16—H16	0.9500
C5—H5	0.9500	C17—C18	1.380 (3)
C6—C7	1.363 (3)	C18—C19	1.390 (3)
C6—H6	0.9500	C18—H18	0.9500
C7—C8	1.413 (3)	C19—H19	0.9500
C7—H7	0.9500		
O2—S1—C1	110.36 (8)	C9—C10—H10	121.8
O2—S1—C14	107.32 (8)	C11—C10—H10	121.8
C1—S1—C14	97.62 (8)	O1—C11—C2	111.20 (16)
C12—O1—C11	106.47 (13)	O1—C11—C10	123.59 (16)
C12—C1—C2	107.38 (15)	C2—C11—C10	125.21 (18)
C12—C1—S1	119.35 (14)	C1—C12—O1	110.59 (16)
C2—C1—S1	133.18 (13)	C1—C12—C13	133.92 (18)
C11—C2—C3	118.75 (16)	O1—C12—C13	115.46 (16)
C11—C2—C1	104.35 (15)	C12—C13—H13A	109.5
C3—C2—C1	136.87 (15)	C12—C13—H13B	109.5
C4—C3—C2	124.19 (15)	H13A—C13—H13B	109.5
C4—C3—C8	118.74 (16)	C12—C13—H13C	109.5
C2—C3—C8	117.06 (15)	H13A—C13—H13C	109.5
C5—C4—C3	120.72 (16)	H13B—C13—H13C	109.5
C5—C4—H4	119.6	C19—C14—C15	120.80 (17)
C3—C4—H4	119.6	C19—C14—S1	120.37 (14)
C4—C5—C6	120.68 (17)	C15—C14—S1	118.69 (13)
C4—C5—H5	119.7	C16—C15—C14	120.14 (17)
C6—C5—H5	119.7	C16—C15—H15	119.9
C7—C6—C5	119.72 (18)	C14—C15—H15	119.9
C7—C6—H6	120.1	C15—C16—C17	118.89 (18)
C5—C6—H6	120.1	C15—C16—H16	120.6
C6—C7—C8	121.57 (17)	C17—C16—H16	120.6
C6—C7—H7	119.2	C18—C17—C16	121.34 (18)
C8—C7—H7	119.2	C18—C17—Br1	119.83 (14)
C7—C8—C3	118.50 (17)	C16—C17—Br1	118.84 (15)
C7—C8—C9	121.04 (17)	C17—C18—C19	119.77 (16)
C3—C8—C9	120.45 (18)	C17—C18—H18	120.1
C10—C9—C8	122.03 (18)	C19—C18—H18	120.1
C10—C9—H9	119.0	C14—C19—C18	119.05 (16)
C8—C9—H9	119.0	C14—C19—H19	120.5
C9—C10—C11	116.45 (17)	C18—C19—H19	120.5
O2—S1—C1—C12	135.73 (15)	C12—O1—C11—C10	178.95 (18)

C14—S1—C1—C12	-112.55 (16)	C3—C2—C11—O1	178.96 (15)
O2—S1—C1—C2	-48.3 (2)	C1—C2—C11—O1	0.4 (2)
C14—S1—C1—C2	63.40 (19)	C3—C2—C11—C10	-0.8 (3)
C12—C1—C2—C11	0.2 (2)	C1—C2—C11—C10	-179.41 (18)
S1—C1—C2—C11	-176.07 (15)	C9—C10—C11—O1	179.36 (18)
C12—C1—C2—C3	-177.9 (2)	C9—C10—C11—C2	-0.9 (3)
S1—C1—C2—C3	5.8 (3)	C2—C1—C12—O1	-0.8 (2)
C11—C2—C3—C4	-176.38 (17)	S1—C1—C12—O1	176.13 (13)
C1—C2—C3—C4	1.6 (3)	C2—C1—C12—C13	-178.5 (2)
C11—C2—C3—C8	2.4 (3)	S1—C1—C12—C13	-1.6 (3)
C1—C2—C3—C8	-179.6 (2)	C11—O1—C12—C1	1.0 (2)
C2—C3—C4—C5	-179.11 (17)	C11—O1—C12—C13	179.19 (18)
C8—C3—C4—C5	2.1 (3)	O2—S1—C14—C19	-9.52 (16)
C3—C4—C5—C6	0.0 (3)	C1—S1—C14—C19	-123.69 (14)
C4—C5—C6—C7	-1.5 (3)	O2—S1—C14—C15	174.66 (14)
C5—C6—C7—C8	0.9 (3)	C1—S1—C14—C15	60.49 (15)
C6—C7—C8—C3	1.2 (3)	C19—C14—C15—C16	0.4 (3)
C6—C7—C8—C9	-177.87 (19)	S1—C14—C15—C16	176.20 (15)
C4—C3—C8—C7	-2.7 (3)	C14—C15—C16—C17	0.4 (3)
C2—C3—C8—C7	178.44 (17)	C15—C16—C17—C18	-0.5 (3)
C4—C3—C8—C9	176.43 (17)	C15—C16—C17—Br1	179.21 (14)
C2—C3—C8—C9	-2.5 (3)	C16—C17—C18—C19	-0.2 (3)
C7—C8—C9—C10	179.9 (2)	Br1—C17—C18—C19	-179.89 (13)
C3—C8—C9—C10	0.8 (3)	C15—C14—C19—C18	-1.1 (3)
C8—C9—C10—C11	0.9 (3)	S1—C14—C19—C18	-176.81 (13)
C12—O1—C11—C2	-0.9 (2)	C17—C18—C19—C14	1.0 (3)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C19—H19···O2 ⁱ	0.95	2.35	3.221 (2)	152

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