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5-Chloro-2-(4-methylphenyl)-3-methylsulfinyl-1-benzofuran

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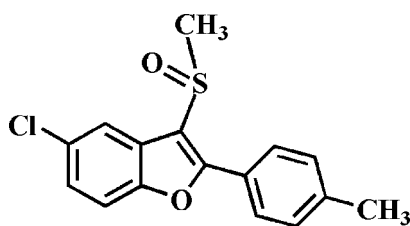
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 Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.038; wR factor = 0.103; data-to-parameter ratio = 18.7.

In the title compound, $\text{C}_{16}\text{H}_{13}\text{ClO}_2\text{S}$, the dihedral angle between the mean plane [r.m.s. deviation = 0.004 (2) Å] of the benzofuran ring system and the 4-methylphenyl ring is 29.25 (8)°. In the crystal, inversion dimers linked by pairs of weak $\text{C}-\text{H}\cdots\text{O}$ interactions generate $R_2^2(14)$ loops.

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 the crystal structures of related compounds, see: Choi *et al.* (2007, 2009).



Experimental

Crystal data

 $\text{C}_{16}\text{H}_{13}\text{ClO}_2\text{S}$
 $M_r = 304.77$
 Triclinic, $P\bar{1}$
 $a = 8.0694$ (8) Å
 $b = 8.0763$ (8) Å
 $c = 11.4208$ (11) Å

 $\alpha = 90.185$ (6)°
 $\beta = 96.280$ (6)°
 $\gamma = 111.701$ (6)°
 $V = 686.63$ (12) Å³
 $Z = 2$
 Mo $K\alpha$ radiation

 $\mu = 0.43$ mm⁻¹
 $T = 173$ K

 $0.35 \times 0.34 \times 0.10$ mm

Data collection

 Bruker SMART APEXII CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2009)
 $T_{\min} = 0.865$, $T_{\max} = 0.958$

 12808 measured reflections
 3423 independent reflections
 3021 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.103$
 $S = 1.05$
 3423 reflections

 183 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.32$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.28$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C14}-\text{H14}\cdots\text{O2}^i$	0.95	2.54	3.436 (2)	158

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

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 for Windows (Farrugia, 2012) 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: NK2211).

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supporting information

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5-Chloro-2-(4-methylphenyl)-3-methylsulfinyl-1-benzofuran

Hong Dae Choi, Pil Ja Seo and Uk Lee

S1. Comment

Many compounds having a benzofuran moiety have drawn much attention due 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 benzofuran derivatives occur in a wide range of natural products (Akgul & Anil, 2003; Soekamto *et al.*, 2003). As a part of our continuing study of 5-chloro-3-methylsulfinyl-1-benzofuran derivatives containing phenyl (Choi *et al.*, 2007) and 4-fluorophenyl (Choi *et al.*, 2009) substituents in 2-position, we report herein the crystal structure of the title compound.

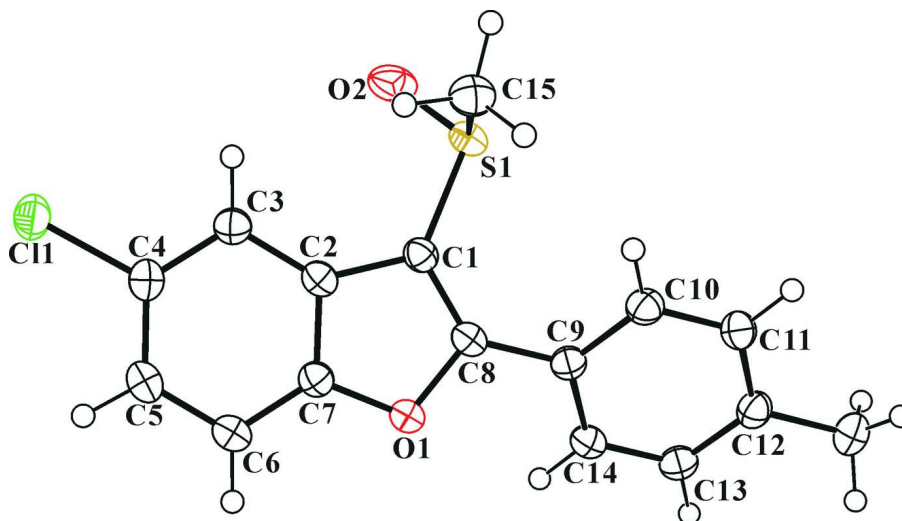
In the title molecule (Fig. 1), the benzofuran unit is essentially planar, with a mean deviation of 0.004 (2) Å from the least-squares plane defined by the nine constituent atoms. The dihedral angle between the mean plane of the benzofuran ring system and the 4-methylphenyl ring is 29.25 (8)°. In the crystal structure, molecules are connected by weak C—H···O hydrogen bonds (Table 1), resulting in a three-dimensional network.

S2. Experimental

3-Chloroperoxybenzoic acid (77%, 269 mg, 1.2 mmol) was added in small portions to a stirred solution of 5-chloro-2-(4-methylphenyl)-3-methylsulfonyl-1-benzofuran (317 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 70%, m.p. 459–460 K; $R_f = 0.49$ (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 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. $U_{iso}(H) = 1.2U_{eq}(C)$ for aryl and $1.5U_{eq}(C)$ for methyl H atoms. The positions of methyl hydrogens were optimized rotationally.

**Figure 1**

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

5-Chloro-2-(4-methylphenyl)-3-methylsulfinyl-1-benzofuran

Crystal data

$C_{16}H_{13}ClO_2S$

$M_r = 304.77$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 8.0694$ (8) Å

$b = 8.0763$ (8) Å

$c = 11.4208$ (11) Å

$\alpha = 90.185$ (6)°

$\beta = 96.280$ (6)°

$\gamma = 111.701$ (6)°

$V = 686.63$ (12) Å³

$Z = 2$

$F(000) = 316$

$D_x = 1.474$ Mg m⁻³

Melting point = 459–460 K

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

Cell parameters from 4424 reflections

$\theta = 2.7$ – 28.1 °

$\mu = 0.43$ mm⁻¹

$T = 173$ K

Block, colourless

$0.35 \times 0.34 \times 0.10$ 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)

$T_{\min} = 0.865$, $T_{\max} = 0.958$

12808 measured reflections

3423 independent reflections

3021 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.048$

$\theta_{\max} = 28.4$ °, $\theta_{\min} = 1.8$ °

$h = -10 \rightarrow 10$

$k = -10 \rightarrow 10$

$l = -15 \rightarrow 15$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.038$

$wR(F^2) = 0.103$

$S = 1.05$

3423 reflections

183 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: difference Fourier map
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0432P)^2 + 0.3137P]$
 where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.32 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.28 \text{ e } \text{Å}^{-3}$

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 (Å^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	0.21613 (6)	-0.06293 (6)	0.00434 (4)	0.03430 (13)
S1	0.79728 (5)	0.32171 (5)	0.40498 (4)	0.02612 (12)
O1	0.30621 (15)	0.15287 (15)	0.50057 (10)	0.0257 (2)
O2	0.82257 (18)	0.19866 (17)	0.31756 (13)	0.0401 (3)
C1	0.5655 (2)	0.2423 (2)	0.41897 (14)	0.0239 (3)
C2	0.4219 (2)	0.1456 (2)	0.32835 (14)	0.0236 (3)
C3	0.4092 (2)	0.0990 (2)	0.20888 (14)	0.0259 (3)
H3	0.5122	0.1328	0.1679	0.031*
C4	0.2396 (2)	0.0014 (2)	0.15325 (14)	0.0269 (3)
C5	0.0852 (2)	-0.0505 (2)	0.21040 (15)	0.0291 (3)
H5	-0.0284	-0.1179	0.1680	0.035*
C6	0.0970 (2)	-0.0040 (2)	0.32891 (15)	0.0282 (3)
H6	-0.0061	-0.0370	0.3698	0.034*
C7	0.2666 (2)	0.0929 (2)	0.38407 (14)	0.0241 (3)
C8	0.4897 (2)	0.2427 (2)	0.52001 (14)	0.0238 (3)
C9	0.5607 (2)	0.3179 (2)	0.63960 (14)	0.0238 (3)
C10	0.7141 (2)	0.4725 (2)	0.66158 (15)	0.0280 (3)
H10	0.7738	0.5314	0.5976	0.034*
C11	0.7804 (2)	0.5414 (2)	0.77602 (15)	0.0296 (4)
H11	0.8862	0.6458	0.7898	0.036*
C12	0.6930 (2)	0.4586 (2)	0.87084 (15)	0.0282 (3)
C13	0.5390 (2)	0.3061 (2)	0.84809 (15)	0.0281 (3)
H13	0.4780	0.2489	0.9121	0.034*
C14	0.4720 (2)	0.2353 (2)	0.73466 (14)	0.0256 (3)
H14	0.3661	0.1309	0.7213	0.031*
C15	0.7646 (3)	0.5323 (3)	0.99514 (16)	0.0390 (4)
H15A	0.6653	0.5005	1.0436	0.059*
H15B	0.8228	0.6624	0.9952	0.059*
H15C	0.8522	0.4818	1.0276	0.059*
C16	0.8112 (3)	0.5192 (2)	0.32875 (16)	0.0318 (4)
H16A	0.9311	0.5749	0.3036	0.048*

H16B	0.7896	0.6031	0.3815	0.048*
H16C	0.7206	0.4877	0.2595	0.048*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0365 (3)	0.0364 (2)	0.0259 (2)	0.01030 (19)	-0.00156 (17)	-0.00172 (16)
S1	0.0199 (2)	0.0253 (2)	0.0326 (2)	0.00772 (16)	0.00336 (15)	0.00359 (15)
O1	0.0207 (6)	0.0284 (6)	0.0265 (6)	0.0073 (5)	0.0031 (4)	-0.0001 (4)
O2	0.0316 (7)	0.0302 (6)	0.0614 (9)	0.0115 (6)	0.0178 (6)	-0.0034 (6)
C1	0.0214 (8)	0.0241 (7)	0.0262 (7)	0.0090 (6)	0.0017 (6)	0.0007 (6)
C2	0.0218 (8)	0.0215 (7)	0.0281 (8)	0.0091 (6)	0.0014 (6)	0.0021 (6)
C3	0.0257 (8)	0.0260 (7)	0.0269 (8)	0.0107 (6)	0.0035 (6)	0.0021 (6)
C4	0.0303 (9)	0.0243 (7)	0.0255 (7)	0.0106 (7)	-0.0008 (6)	0.0001 (6)
C5	0.0252 (8)	0.0272 (8)	0.0315 (8)	0.0076 (7)	-0.0026 (7)	0.0006 (6)
C6	0.0226 (8)	0.0287 (8)	0.0329 (8)	0.0089 (7)	0.0031 (6)	0.0024 (6)
C7	0.0238 (8)	0.0237 (7)	0.0256 (7)	0.0098 (6)	0.0021 (6)	0.0015 (6)
C8	0.0201 (8)	0.0218 (7)	0.0290 (8)	0.0073 (6)	0.0021 (6)	0.0021 (6)
C9	0.0241 (8)	0.0240 (7)	0.0258 (7)	0.0117 (6)	0.0034 (6)	0.0014 (6)
C10	0.0301 (9)	0.0248 (7)	0.0282 (8)	0.0081 (7)	0.0072 (7)	0.0031 (6)
C11	0.0297 (9)	0.0234 (7)	0.0329 (8)	0.0065 (7)	0.0046 (7)	-0.0028 (6)
C12	0.0311 (9)	0.0278 (8)	0.0275 (8)	0.0132 (7)	0.0030 (7)	-0.0024 (6)
C13	0.0298 (9)	0.0296 (8)	0.0270 (8)	0.0123 (7)	0.0075 (7)	0.0048 (6)
C14	0.0228 (8)	0.0236 (7)	0.0301 (8)	0.0081 (6)	0.0040 (6)	0.0027 (6)
C15	0.0417 (11)	0.0422 (10)	0.0295 (9)	0.0116 (9)	0.0035 (8)	-0.0084 (7)
C16	0.0342 (10)	0.0272 (8)	0.0355 (9)	0.0118 (7)	0.0087 (7)	0.0073 (7)

Geometric parameters (Å, °)

C11—C4	1.7462 (16)	C9—C10	1.393 (2)
S1—O2	1.4879 (13)	C9—C14	1.399 (2)
S1—C1	1.7648 (17)	C10—C11	1.388 (2)
S1—C16	1.7925 (17)	C10—H10	0.9500
O1—C7	1.3768 (18)	C11—C12	1.393 (2)
O1—C8	1.3790 (19)	C11—H11	0.9500
C1—C8	1.364 (2)	C12—C13	1.387 (2)
C1—C2	1.445 (2)	C12—C15	1.505 (2)
C2—C7	1.393 (2)	C13—C14	1.382 (2)
C2—C3	1.397 (2)	C13—H13	0.9500
C3—C4	1.381 (2)	C14—H14	0.9500
C3—H3	0.9500	C15—H15A	0.9800
C4—C5	1.396 (2)	C15—H15B	0.9800
C5—C6	1.387 (2)	C15—H15C	0.9800
C5—H5	0.9500	C16—H16A	0.9800
C6—C7	1.378 (2)	C16—H16B	0.9800
C6—H6	0.9500	C16—H16C	0.9800
C8—C9	1.459 (2)		

O2—S1—C1	106.64 (7)	C10—C9—C8	121.45 (14)
O2—S1—C16	106.07 (8)	C14—C9—C8	119.67 (15)
C1—S1—C16	97.73 (8)	C11—C10—C9	120.65 (15)
C7—O1—C8	106.62 (12)	C11—C10—H10	119.7
C8—C1—C2	107.24 (14)	C9—C10—H10	119.7
C8—C1—S1	126.49 (13)	C10—C11—C12	120.54 (16)
C2—C1—S1	125.89 (12)	C10—C11—H11	119.7
C7—C2—C3	119.27 (15)	C12—C11—H11	119.7
C7—C2—C1	105.00 (14)	C13—C12—C11	118.46 (15)
C3—C2—C1	135.73 (15)	C13—C12—C15	120.74 (16)
C4—C3—C2	116.75 (15)	C11—C12—C15	120.80 (16)
C4—C3—H3	121.6	C14—C13—C12	121.65 (16)
C2—C3—H3	121.6	C14—C13—H13	119.2
C3—C4—C5	123.29 (15)	C12—C13—H13	119.2
C3—C4—C11	118.71 (13)	C13—C14—C9	119.81 (15)
C5—C4—C11	118.00 (13)	C13—C14—H14	120.1
C6—C5—C4	120.20 (15)	C9—C14—H14	120.1
C6—C5—H5	119.9	C12—C15—H15A	109.5
C4—C5—H5	119.9	C12—C15—H15B	109.5
C7—C6—C5	116.26 (15)	H15A—C15—H15B	109.5
C7—C6—H6	121.9	C12—C15—H15C	109.5
C5—C6—H6	121.9	H15A—C15—H15C	109.5
O1—C7—C6	125.14 (14)	H15B—C15—H15C	109.5
O1—C7—C2	110.63 (14)	S1—C16—H16A	109.5
C6—C7—C2	124.23 (15)	S1—C16—H16B	109.5
C1—C8—O1	110.51 (14)	H16A—C16—H16B	109.5
C1—C8—C9	133.98 (15)	S1—C16—H16C	109.5
O1—C8—C9	115.51 (13)	H16A—C16—H16C	109.5
C10—C9—C14	118.87 (15)	H16B—C16—H16C	109.5
O2—S1—C1—C8	142.96 (15)	C1—C2—C7—C6	-179.96 (15)
C16—S1—C1—C8	-107.64 (15)	C2—C1—C8—O1	-0.17 (17)
O2—S1—C1—C2	-29.09 (16)	S1—C1—C8—O1	-173.43 (11)
C16—S1—C1—C2	80.31 (15)	C2—C1—C8—C9	-179.34 (16)
C8—C1—C2—C7	-0.40 (17)	S1—C1—C8—C9	7.4 (3)
S1—C1—C2—C7	172.92 (12)	C7—O1—C8—C1	0.68 (17)
C8—C1—C2—C3	179.75 (17)	C7—O1—C8—C9	-179.99 (12)
S1—C1—C2—C3	-6.9 (3)	C1—C8—C9—C10	29.6 (3)
C7—C2—C3—C4	-0.1 (2)	O1—C8—C9—C10	-149.54 (15)
C1—C2—C3—C4	179.69 (16)	C1—C8—C9—C14	-151.36 (18)
C2—C3—C4—C5	0.1 (2)	O1—C8—C9—C14	29.5 (2)
C2—C3—C4—C11	-179.18 (11)	C14—C9—C10—C11	1.5 (2)
C3—C4—C5—C6	0.2 (3)	C8—C9—C10—C11	-179.49 (15)
C11—C4—C5—C6	179.45 (12)	C9—C10—C11—C12	-1.0 (3)
C4—C5—C6—C7	-0.4 (2)	C10—C11—C12—C13	0.2 (2)
C8—O1—C7—C6	179.86 (15)	C10—C11—C12—C15	179.82 (16)
C8—O1—C7—C2	-0.94 (16)	C11—C12—C13—C14	0.3 (2)
C5—C6—C7—O1	179.43 (14)	C15—C12—C13—C14	-179.38 (16)

C5—C6—C7—C2	0.3 (2)	C12—C13—C14—C9	0.2 (2)
C3—C2—C7—O1	-179.29 (13)	C10—C9—C14—C13	-1.0 (2)
C1—C2—C7—O1	0.83 (17)	C8—C9—C14—C13	179.91 (14)
C3—C2—C7—C6	-0.1 (2)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C14—H14 \cdots O2 ⁱ	0.95	2.54	3.436 (2)	158

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