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2-Chloro-*N*-(2-methylbenzoyl)benzene-sulfonamide

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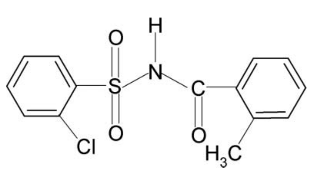
Key indicators: single-crystal X-ray study; T = 299 K; mean σ(C–C) = 0.004 Å; R factor = 0.041; wR factor = 0.110; data-to-parameter ratio = 15.7.

In the title compound, $C_{14}H_{12}ClNO_3S$, the N—H bond is antiperiplanar to the C=O bond. The dihedral angle between the two aromatic rings is 78.7 (1)°. The crystal structure features inversion-related dimers linked by pairs of N—H···O(S) hydrogen bonds.

Related literature

For background to our study of the effect of ring and side-chain substitutions on the crystal structures of *N*-aryl sulfonamides and for related structures, see: Gowda *et al.* (2009, 2010*a*,*b*).



Experimental

Crystal data

$C_{14}H_{12}ClNO_3S$
M_r = 309.76
Monoclinic, $P2_1/n$
a = 6.6086 (5) Å
b = 10.9621 (9) Å
c = 20.080 (2) Å
β = 95.586 (8)°
V = 1447.8 (2) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.41 mm^{-1}
T = 299 K
0.34 × 0.32 × 0.28 mm

Data collection

Oxford Diffraction Xcalibur diffractometer with a Sapphire CCD detector
Absorption correction: multi-scan (*CrysAlis RED*; Oxford Diffraction, 2009)
T_{min} = 0.872, T_{max} = 0.893
5395 measured reflections
2917 independent reflections
2592 reflections with $I > 2\sigma(I)$
R_{int} = 0.017

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.041
$wR(F^2)$ = 0.110
S = 1.07
2917 reflections
186 parameters
1 restraint
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.31 e Å^{-3}
$\Delta\rho_{min}$ = −0.33 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1N···O2^{i}	0.84 (2)	2.11 (2)	2.937 (2)	172 (2)

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

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2009); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2009); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); 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: BT5259).

References

Gowda, B. T., Foro, S., Suchetan, P. A. & Fuess, H. (2009). *Acta Cryst.* E**65**, o2516.
Gowda, B. T., Foro, S., Suchetan, P. A. & Fuess, H. (2010*a*). *Acta Cryst.* E**66**, o794.
Gowda, B. T., Foro, S., Suchetan, P. A. & Fuess, H. (2010*b*). *Acta Cryst.* E**66**, o326.
Oxford Diffraction (2009). *CrysAlis CCD* and *CrysAlis RED*. Oxford Diffraction Ltd, Abingdon, England.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.
Spek, A. L. (2009). *Acta Cryst.* D**65**, 148–155.

supporting information

Acta Cryst. (2010). E**66**, o1281 [https://doi.org/10.1107/S1600536810015990]

2-Chloro-*N*-(2-methylbenzoyl)benzenesulfonamide

P. A. Suchetan, B. Thimme Gowda, Sabine Foro and Hartmut Fuess

S1. Comment

In the present work, as a part of studying the effect of ring and the side chain substitutions on the crystal structures of *N*-aryl sulfonamides (Gowda *et al.*, 2009, 2010*a*,*b*), the structure of 2-chloro-*N*-(2-methylbenzoyl)benzenesulfonamide (I) has been determined. In the C—SO_2—NH—C(O) segment, the N—H bond is *anti* to the C=O bond (Fig.1), similar to those observed in *N*-(2-chlorobenzoyl)2-chlorobenzenesulfonamide (II) (Suchetan *et al.*, 2010), *N*-(benzoyl)benzene-sulfonamide (III) (Gowda *et al.*, 2009), and *N*-(benzoyl)2-chlorobenzenesulfonamide (IV) (Gowda *et al.*, 2010*a*).

Further, the conformation of the C=O bond in the C—SO_2—NH—C(O) segment of (I) is *syn* to the *ortho*-methyl group in the benzoyl ring, similar to that observed between the *ortho*-Cl and the C=O bond in (II).

The molecules are twisted at the *S* atom with the torsional angle of -64.0 (2)°, compared to those of 66.5 (2)° in (II), -66.9 (3)° in (III), and 66.7 (2)° in (IV).

The dihedral angles between the sulfonyl benzene ring and the —SO_2—NH—C—O segment is 88.4 (1)°, compared to the values of 86.9 (1)° in (II), 86.5(0.1) in (III) and 87.3 (1)° in (IV). Furthermore, the dihedral angle between the sulfonyl and the benzoyl benzene rings is 78.7 (1)°, compared to the values of 76.9 (1)° in (II), of 80.3(0.1) in (III), 69.8 (1)° (molecule 1) and 89.8 (1)° (molecule 2) in (III) and 73.3 (1)° in (IV).

The packing of molecules linked by N—H···O(S) hydrogen bonds (Table 1) is shown in Fig. 2.

S2. Experimental

The title compound was prepared by refluxing a mixture of 2-methylbenzoic acid, 2-chlorobenzenesulfonamide and phosphorous oxy chloride for 3 h on a water bath. The resultant mixture was cooled and poured into ice cold water. The solid obtained was filtered, washed thoroughly with water and then dissolved in sodium bicarbonate solution. The compound was later reprecipitated by acidifying the filtered solution with dilute HCl. It was filtered, dried and recrystallized.

Prism like colourless single crystals of the title compound used in X-ray diffraction studies were obtained by slow evaporation of its toluene solution at room temperature.

S3. Refinement

The H atom of the NH group was located in a difference map and its coordinates were refined with a distance restraint of N—H = 0.86 (2) %A. The other H atoms were positioned with idealized geometry using a riding model with C—H = 0.93–0.96 Å. All H atoms were refined with isotropic displacement parameters set to 1.2 times of the U_{eq} of the parent atom.

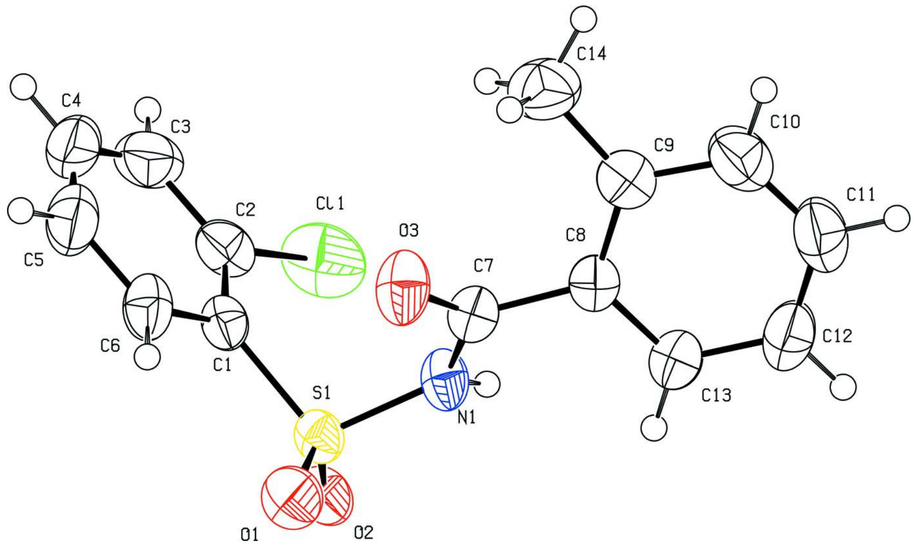

Figure 1

Molecular structure of the title compound, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.

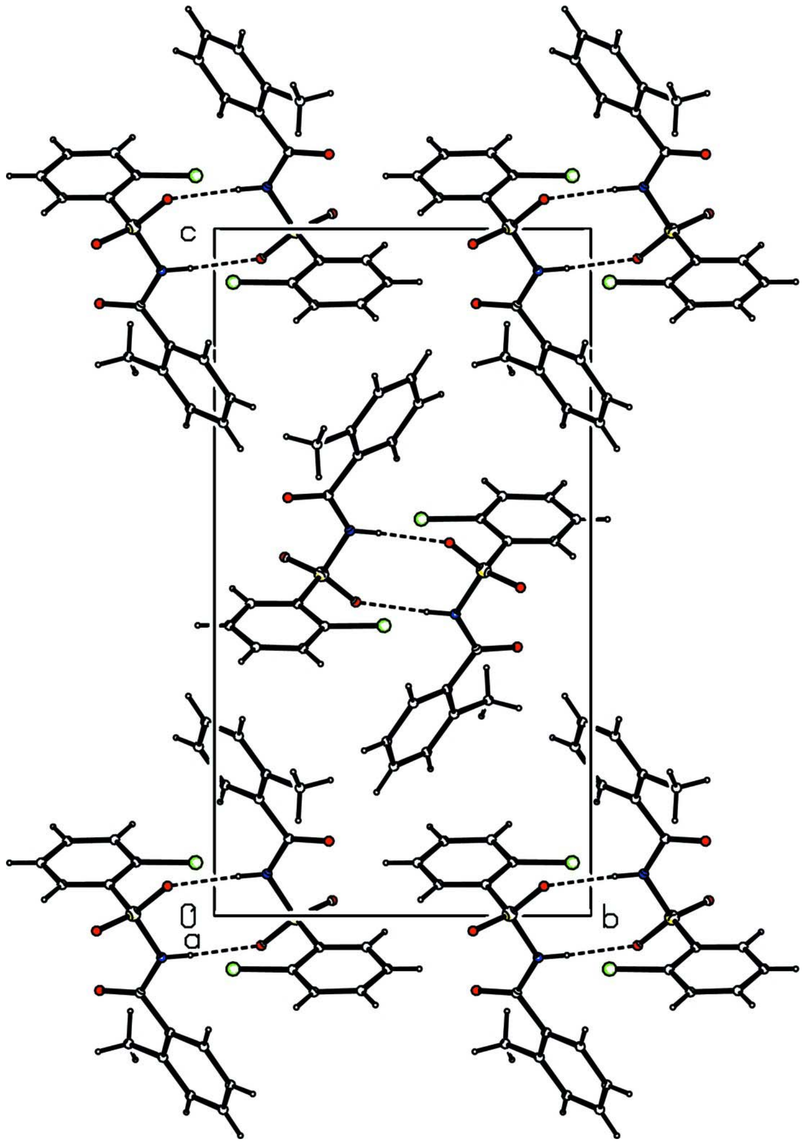

Figure 2

Molecular packing in the title compound. Hydrogen bonds are shown as dashed lines.

2-Chloro-*N*-(2-methylbenzoyl)benzenesulfonamide

Crystal data

$C_{14}H_{12}ClNO_3S$
M_r = 309.76
Monoclinic, $P2_1/n$
Hall symbol: -P 2yn
a = 6.6086 (5) Å
b = 10.9621 (9) Å
c = 20.080 (2) Å
β = 95.586 (8)°
V = 1447.8 (2) Å^3
Z = 4
F(000) = 640
D_x = 1.421 Mg m^{-3}
Mo $K\alpha$ radiation, λ = 0.71073 Å
Cell parameters from 3866 reflections
θ = 2.7–27.8°
μ = 0.41 mm^{-1}
T = 299 K
Prism, colourless
0.34 × 0.32 × 0.28 mm

Data collection

Oxford Diffraction Xcalibur diffractometer with a Sapphire CCD detector
Radiation source: fine-focus sealed tube
Graphite monochromator
Rotation method data acquisition using ω and phi scans
Absorption correction: multi-scan (*CrysAlis RED*; Oxford Diffraction, 2009)
$T_{min} = 0.872$, $T_{max} = 0.893$
5395 measured reflections
2917 independent reflections
2592 reflections with $I > 2\sigma(I)$
$R_{int} = 0.017$
$\theta_{max} = 26.4°$, $\theta_{min} = 2.8°$
$h = -7 \rightarrow 8$
$k = -13 \rightarrow 7$
$l = -25 \rightarrow 25$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.041$
$wR(F^2) = 0.110$
$S = 1.07$
2917 reflections
186 parameters
1 restraint
Primary atom site location: structure-invariant direct methods
Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H atoms treated by a mixture of independent and constrained refinement
$w = 1/[\sigma^2(F_o^2) + (0.0502P)^2 + 0.7805P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max} = 0.31$ e Å^{-3}
$\Delta\rho_{min} = -0.33$ e Å^{-3}
Extinction correction: *SHELXL97* (Sheldrick, 2008), $Fc^* = kFc[1+0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.146 (5)

Special details

Experimental. CrysAlis RED (Oxford Diffraction, 2009) Empirical absorption correction using spherical harmonics, implemented in SCALE3 ABSPACK scaling algorithm.

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 $F^2 > \sigma(F^2)$ 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)

	x	y	z	U_{iso}*/U_{eq}
C1	0.2188 (3)	0.22154 (17)	0.45434 (9)	0.0389 (4)
C2	0.3529 (3)	0.2921 (2)	0.42162 (11)	0.0504 (5)
C3	0.5045 (4)	0.2357 (3)	0.38928 (13)	0.0717 (8)
H3	0.5940	0.2821	0.3667	0.086*
C4	0.5208 (4)	0.1097 (3)	0.39104 (14)	0.0771 (9)
H4	0.6237	0.0719	0.3702	0.092*
C5	0.3890 (4)	0.0403 (3)	0.42274 (13)	0.0658 (7)
H5	0.4005	−0.0443	0.4228	0.079*
C6	0.2392 (4)	0.0956 (2)	0.45452 (10)	0.0502 (5)
H6	0.1500	0.0480	0.4765	0.060*
C7	0.2598 (3)	0.30250 (17)	0.61222 (9)	0.0392 (4)
C8	0.3281 (3)	0.38164 (17)	0.67032 (9)	0.0389 (4)

C9	0.5185 (3)	0.3631 (2)	0.70605 (11)	0.0492 (5)
C10	0.5679 (4)	0.4362 (3)	0.76164 (13)	0.0661 (7)
H10	0.6937	0.4257	0.7860	0.079*
C11	0.4385 (4)	0.5231 (3)	0.78199 (13)	0.0680 (7)
H11	0.4759	0.5693	0.8201	0.082*
C12	0.2534 (4)	0.5422 (2)	0.74615 (12)	0.0608 (6)
H12	0.1660	0.6020	0.7594	0.073*
C13	0.1982 (3)	0.4717 (2)	0.69029 (10)	0.0472 (5)
H13	0.0731	0.4845	0.6658	0.057*
C14	0.6730 (4)	0.2723 (3)	0.68618 (15)	0.0696 (7)
H14A	0.6814	0.2770	0.6388	0.083*
H14B	0.6325	0.1915	0.6977	0.083*
H14C	0.8034	0.2905	0.7094	0.083*
N1	0.1386 (3)	0.36014 (14)	0.56089 (8)	0.0412 (4)
H1N	0.135 (3)	0.4361 (15)	0.5574 (11)	0.049*
O1	−0.0894 (2)	0.18713 (14)	0.52207 (8)	0.0559 (4)
O2	−0.0831 (2)	0.37563 (12)	0.45683 (8)	0.0497 (4)
O3	0.3005 (3)	0.19555 (13)	0.60869 (7)	0.0554 (4)
Cl1	0.34073 (12)	0.45035 (6)	0.42249 (4)	0.0807 (3)
S1	0.02324 (7)	0.28443 (4)	0.49721 (2)	0.03857 (18)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0448 (10)	0.0355 (10)	0.0351 (9)	0.0007 (8)	−0.0027 (7)	−0.0024 (7)
C2	0.0472 (11)	0.0577 (13)	0.0447 (11)	−0.0085 (10)	−0.0030 (9)	0.0012 (9)
C3	0.0462 (12)	0.117 (3)	0.0523 (13)	−0.0120 (14)	0.0068 (10)	−0.0090 (15)
C4	0.0600 (15)	0.108 (2)	0.0607 (15)	0.0241 (16)	−0.0054 (12)	−0.0369 (16)
C5	0.0772 (17)	0.0629 (16)	0.0551 (14)	0.0211 (13)	−0.0052 (12)	−0.0178 (12)
C6	0.0670 (13)	0.0384 (11)	0.0440 (11)	0.0063 (10)	0.0001 (9)	−0.0051 (9)
C7	0.0477 (10)	0.0337 (9)	0.0366 (9)	0.0017 (8)	0.0067 (8)	0.0037 (7)
C8	0.0473 (10)	0.0349 (9)	0.0351 (9)	−0.0035 (8)	0.0076 (8)	0.0031 (7)
C9	0.0507 (11)	0.0501 (12)	0.0465 (11)	−0.0030 (9)	0.0038 (9)	0.0045 (9)
C10	0.0616 (14)	0.0768 (18)	0.0574 (14)	−0.0082 (13)	−0.0074 (11)	−0.0048 (13)
C11	0.0824 (18)	0.0715 (17)	0.0497 (13)	−0.0146 (14)	0.0038 (12)	−0.0161 (12)
C12	0.0743 (15)	0.0569 (14)	0.0534 (13)	0.0010 (12)	0.0185 (11)	−0.0140 (11)
C13	0.0527 (11)	0.0465 (11)	0.0432 (10)	0.0024 (9)	0.0085 (9)	−0.0018 (9)
C14	0.0565 (14)	0.0680 (17)	0.0833 (18)	0.0118 (12)	0.0019 (13)	0.0033 (14)
N1	0.0570 (10)	0.0242 (7)	0.0411 (8)	0.0016 (7)	−0.0015 (7)	−0.0002 (6)
O1	0.0591 (9)	0.0400 (8)	0.0701 (10)	−0.0135 (7)	0.0146 (8)	−0.0011 (7)
O2	0.0509 (8)	0.0349 (7)	0.0597 (9)	0.0040 (6)	−0.0124 (7)	−0.0026 (6)
O3	0.0846 (11)	0.0320 (7)	0.0483 (8)	0.0118 (7)	−0.0006 (7)	0.0017 (6)
Cl1	0.0810 (5)	0.0573 (4)	0.1049 (6)	−0.0258 (3)	0.0148 (4)	0.0216 (4)
S1	0.0428 (3)	0.0266 (3)	0.0456 (3)	−0.00243 (18)	0.00085 (19)	−0.00064 (18)

Geometric parameters (Å, º)

C1—C6	1.387 (3)	C9—C10	1.387 (3)
C1—C2	1.389 (3)	C9—C14	1.507 (3)
C1—S1	1.761 (2)	C10—C11	1.369 (4)
C2—C3	1.391 (4)	C10—H10	0.9300
C2—Cl1	1.737 (3)	C11—C12	1.373 (4)
C3—C4	1.386 (4)	C11—H11	0.9300
C3—H3	0.9300	C12—C13	1.382 (3)
C4—C5	1.361 (4)	C12—H12	0.9300
C4—H4	0.9300	C13—H13	0.9300
C5—C6	1.370 (3)	C14—H14A	0.9600
C5—H5	0.9300	C14—H14B	0.9600
C6—H6	0.9300	C14—H14C	0.9600
C7—O3	1.206 (2)	N1—S1	1.6478 (17)
C7—N1	1.394 (2)	N1—H1N	0.836 (16)
C7—C8	1.488 (3)	O1—S1	1.4188 (15)
C8—C13	1.393 (3)	O2—S1	1.4283 (14)
C8—C9	1.401 (3)		
C6—C1—C2	119.3 (2)	C11—C10—C9	122.5 (2)
C6—C1—S1	117.65 (17)	C11—C10—H10	118.7
C2—C1—S1	123.06 (16)	C9—C10—H10	118.7
C1—C2—C3	119.7 (2)	C10—C11—C12	120.0 (2)
C1—C2—Cl1	121.28 (18)	C10—C11—H11	120.0
C3—C2—Cl1	119.0 (2)	C12—C11—H11	120.0
C4—C3—C2	119.2 (3)	C11—C12—C13	119.5 (2)
C4—C3—H3	120.4	C11—C12—H12	120.3
C2—C3—H3	120.4	C13—C12—H12	120.3
C5—C4—C3	121.2 (2)	C12—C13—C8	120.6 (2)
C5—C4—H4	119.4	C12—C13—H13	119.7
C3—C4—H4	119.4	C8—C13—H13	119.7
C4—C5—C6	119.7 (3)	C9—C14—H14A	109.5
C4—C5—H5	120.2	C9—C14—H14B	109.5
C6—C5—H5	120.2	H14A—C14—H14B	109.5
C5—C6—C1	120.9 (2)	C9—C14—H14C	109.5
C5—C6—H6	119.5	H14A—C14—H14C	109.5
C1—C6—H6	119.5	H14B—C14—H14C	109.5
O3—C7—N1	120.83 (18)	C7—N1—S1	122.33 (13)
O3—C7—C8	124.07 (18)	C7—N1—H1N	121.6 (16)
N1—C7—C8	115.09 (16)	S1—N1—H1N	115.4 (16)
C13—C8—C9	120.23 (19)	O1—S1—O2	118.67 (10)
C13—C8—C7	119.36 (18)	O1—S1—N1	108.94 (9)
C9—C8—C7	120.38 (18)	O2—S1—N1	104.66 (8)
C10—C9—C8	117.2 (2)	O1—S1—C1	108.22 (9)
C10—C9—C14	118.8 (2)	O2—S1—C1	109.90 (9)
C8—C9—C14	123.9 (2)	N1—S1—C1	105.69 (9)

C6—C1—C2—C3	0.4 (3)	C8—C9—C10—C11	0.2 (4)
S1—C1—C2—C3	179.48 (17)	C14—C9—C10—C11	178.2 (3)
C6—C1—C2—Cl1	−177.48 (16)	C9—C10—C11—C12	−1.3 (4)
S1—C1—C2—Cl1	1.6 (2)	C10—C11—C12—C13	1.0 (4)
C1—C2—C3—C4	−0.8 (3)	C11—C12—C13—C8	0.2 (4)
Cl1—C2—C3—C4	177.1 (2)	C9—C8—C13—C12	−1.2 (3)
C2—C3—C4—C5	1.2 (4)	C7—C8—C13—C12	176.6 (2)
C3—C4—C5—C6	−1.1 (4)	O3—C7—N1—S1	7.1 (3)
C4—C5—C6—C1	0.6 (4)	C8—C7—N1—S1	−171.69 (14)
C2—C1—C6—C5	−0.2 (3)	C7—N1—S1—O1	52.08 (18)
S1—C1—C6—C5	−179.40 (17)	C7—N1—S1—O2	179.96 (16)
O3—C7—C8—C13	−143.2 (2)	C7—N1—S1—C1	−64.00 (18)
N1—C7—C8—C13	35.5 (3)	C6—C1—S1—O1	−3.36 (19)
O3—C7—C8—C9	34.6 (3)	C2—C1—S1—O1	177.51 (16)
N1—C7—C8—C9	−146.66 (19)	C6—C1—S1—O2	−134.38 (16)
C13—C8—C9—C10	1.0 (3)	C2—C1—S1—O2	46.49 (18)
C7—C8—C9—C10	−176.8 (2)	C6—C1—S1—N1	113.21 (16)
C13—C8—C9—C14	−176.9 (2)	C2—C1—S1—N1	−65.91 (18)
C7—C8—C9—C14	5.4 (3)		

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N1—H1*N*···O2^{i}	0.84 (2)	2.11 (2)	2.937 (2)	172 (2)

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