

3-(*tert*-Butoxycarbonyl)-2-(4-chlorophenyl)-1,3-thiazolidine-4-carboxylic acid. Corrigendum

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The author list in the paper by Ding [*Acta Cryst.* (2010), E66, o2633] is corrected.

In the paper by Ding (2010), the author list is incomplete. The full list of authors is given above.

References

Ding, S.-M. (2010). *Acta Cryst.* E66, o2633.

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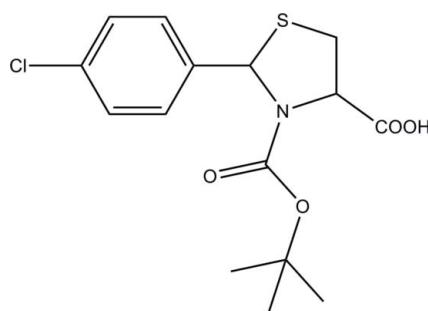
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Key indicators: single-crystal X-ray study; $T = 293\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.010\text{ \AA}$; R factor = 0.062; wR factor = 0.159; data-to-parameter ratio = 8.9.

In the title compound, $\text{C}_{15}\text{H}_{18}\text{ClNO}_4\text{S}$, the thiazolidine ring adopts a twisted conformation about the $\text{S}-\text{C}(\text{methylene})$ bond. The dihedral angle between the five- and six-membered rings is $77.2(3)^\circ$. In the crystal, the molecules are linked by $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds, generating $C(7)$ chains propagating in [100].

Related literature

For background to the biological properties of the title compound, see: Lu *et al.* (2010); Song *et al.* (2009). For reference bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{15}\text{H}_{18}\text{ClNO}_4\text{S}$
 $M_r = 343.81$

Monoclinic, $P2_1$
 $a = 6.4600(13)\text{ \AA}$

$b = 10.641(2)\text{ \AA}$
 $c = 12.411(3)\text{ \AA}$
 $\beta = 94.52(3)^\circ$
 $V = 850.5(3)\text{ \AA}^3$
 $Z = 2$

Mo $K\alpha$ radiation
 $\mu = 0.36\text{ mm}^{-1}$
 $T = 293\text{ K}$
 $0.30 \times 0.20 \times 0.10\text{ mm}$

Data collection

Enraf–Nonius CAD-4 diffractometer
Absorption correction: ψ scan (North *et al.*, 1968)
 $T_{\min} = 0.899$, $T_{\max} = 0.965$
1638 measured reflections

1638 independent reflections
1363 reflections with $I > 2\sigma(I)$
200 standard reflections every 3 reflections
intensity decay: 1%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.062$
 $wR(F^2) = 0.159$
 $S = 1.08$
1638 reflections
185 parameters
89 restraints

H-atom parameters constrained
 $\Delta\rho_{\max} = 0.43\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.37\text{ e \AA}^{-3}$
Absolute structure: Flack (1983)
Flack parameter: $-0.09(19)$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O}2-\text{H}2\text{A}\cdots\text{O}3^i$	0.82	1.83	2.638 (6)	167

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

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5637).

References

- Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–19.
- Enraf–Nonius (1989). *CAD-4 Software*. Enraf–Nonius, Delft, The Netherlands.
- Flack, H. D. (1983). *Acta Cryst. A* **39**, 876–881.
- Harms, K. & Wocadlo, S. (1995). *XCAD4*. University of Marburg, Germany.
- Lu, Y., Wang, Z., Li, C.-M., Chen, J.-J., Dalton, J. T., Li, W. & Miller, D. D. (2010). *Bioorg. Med. Chem.* **18**, 477–495.
- North, A. C. T., Phillips, D. C. & Mathews, F. S. (1968). *Acta Cryst. A* **24**, 351–359.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Song, Z.-C., Ma, G.-Y., Lv, P.-C., Li, H.-Q., Xiao, Z.-P. & Zhu, H.-L. (2009). *Eur. J. Med. Chem.* **44**, 3903–3908.

supporting information

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3-(*tert*-Butoxycarbonyl)-2-(4-chlorophenyl)-1,3-thiazolidine-4-carboxylic acid

Shu-Min Ding

S1. Comment

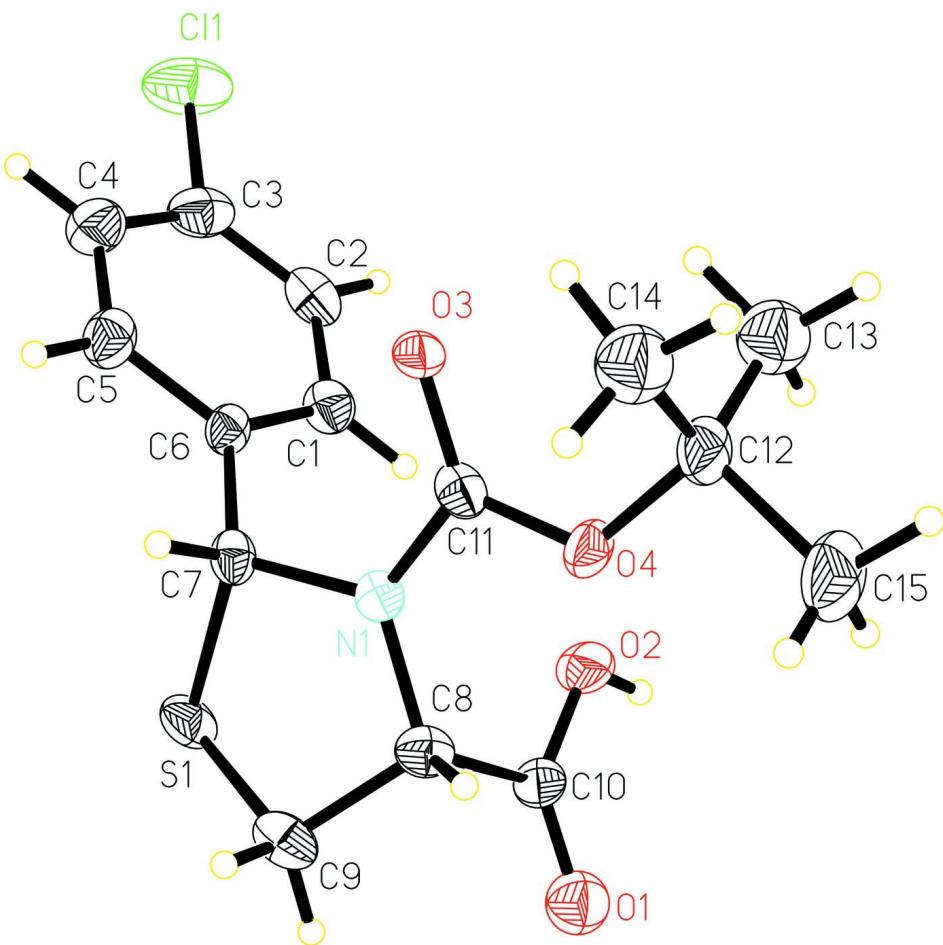
Recently, 3-*tert*-butoxycarbonyl-2-arylthiazolidine-4-carboxylic acid derivatives have been reported to possess antimicrobial and antitumor activities (Song *et al.*, 2009; Lu *et al.*, 2010). In this work, we report here the crystal structure of the title compound, (I). In (I), all bond lengths are within normal ranges (Allen *et al.*, 1987) (Fig. 1). There are intermolecular O—H···O hydrogen bonds in (I).

S2. Experimental

A mixture of *L*-cysteine (1.41 g, 10 mmol) and 4-chlorobenzaldehyde (1.4 g, 10 mmol) in methanol (100 ml) was stirred at room temperature for 10 h, and the separated solid was collected, washed with diethyl ether, and dried to obtain 2-(4-chlorophenyl)thiazolidine-4-carboxylic with yield of 90%. In ice water, 2-(4-chlorophenyl)thiazolidine-4-carboxylic (1 mmol) was dissolved in 1 N NaOH (1 ml) and 1,4-dioxane (10 ml); then di-*tert*-butyldicarbonate (1 mmol) was added slowly and stirred at room temperature for 6 h. The reaction mixture was concentrated in a vacuum and washed with ethyl acetate (10 ml). The aqueous phase was adjusted to pH 4 by adding 1 N HCl, then extracted with ethyl acetate, dried with magnesium sulfate, filtered, After keeping the filtrate in air for 5 d, colorless block-shaped crystals of (I) were formed.

S3. Refinement

All H atoms were positioned geometrically (C—H = 0.93 Å for the aromatic H atoms and C—H = 0.96 Å for the aliphatic H atoms) and were refined as riding, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$.

**Figure 1**

The structure of (I) showing 30% probability displacement ellipsoids.

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Crystal data



$M_r = 343.81$

Monoclinic, $P2_1$

Hall symbol: P 2yb

$a = 6.4600 (13)$ Å

$b = 10.641 (2)$ Å

$c = 12.411 (3)$ Å

$\beta = 94.52 (3)^\circ$

$V = 850.5 (3)$ Å³

$Z = 2$

$F(000) = 360$

$D_x = 1.343 \text{ Mg m}^{-3}$

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

Cell parameters from 25 reflections

$\theta = 9\text{--}12^\circ$

$\mu = 0.36 \text{ mm}^{-1}$

$T = 293$ K

Block, colorless

$0.30 \times 0.20 \times 0.10$ mm

Data collection

Enraf–Nonius CAD-4
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\omega/2\theta$ scan

Absorption correction: ψ scan
(North *et al.*, 1968)

$T_{\min} = 0.899$, $T_{\max} = 0.965$

1638 measured reflections

1638 independent reflections

1363 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.000$
 $\theta_{\text{max}} = 25.3^\circ$, $\theta_{\text{min}} = 1.7^\circ$
 $h = -7 \rightarrow 7$
 $k = 0 \rightarrow 12$

$l = 0 \rightarrow 14$
200 standard reflections every 3 reflections
intensity decay: 1%

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.062$
 $wR(F^2) = 0.159$
 $S = 1.08$
1638 reflections
185 parameters
89 restraints
Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0649P)^2 + 1.2912P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.43 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.37 \text{ e } \text{\AA}^{-3}$
Absolute structure: Flack (1983)
Absolute structure parameter: -0.09 (19)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s 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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.3575 (11)	0.4607 (7)	0.7980 (6)	0.0457 (16)
H1	0.4776	0.4984	0.7768	0.055*
C2	0.3229 (12)	0.3336 (8)	0.7816 (6)	0.0519 (17)
H2	0.4183	0.2863	0.7465	0.062*
C3	0.1505 (12)	0.2766 (8)	0.8162 (7)	0.0555 (18)
C4	0.0021 (13)	0.3440 (8)	0.8654 (6)	0.0585 (18)
H4	-0.1163	0.3054	0.8879	0.070*
C5	0.0368 (11)	0.4720 (7)	0.8800 (6)	0.0492 (16)
H5	-0.0604	0.5195	0.9136	0.059*
C6	0.2109 (9)	0.5308 (6)	0.8464 (5)	0.0349 (13)
C7	0.2361 (9)	0.6677 (6)	0.8692 (4)	0.0358 (13)
H7	0.1036	0.7023	0.8887	0.043*
C8	0.4770 (9)	0.8351 (7)	0.8161 (5)	0.0409 (14)
H8	0.4364	0.9165	0.7842	0.049*
O1	0.8408 (8)	0.8639 (5)	0.8109 (4)	0.060
C9	0.4827 (11)	0.8471 (7)	0.9375 (5)	0.0494 (16)
H9A	0.3760	0.9043	0.9583	0.059*
H9B	0.6170	0.8774	0.9671	0.059*
C10	0.6910 (9)	0.8031 (8)	0.7835 (5)	0.0459 (17)

C11	0.2163 (9)	0.7473 (7)	0.6838 (5)	0.0373 (14)
C12	0.2095 (11)	0.8500 (8)	0.5098 (5)	0.0520 (18)
C13	0.2564 (14)	0.7386 (9)	0.4443 (8)	0.070
H13A	0.3999	0.7161	0.4586	0.106*
H13B	0.2293	0.7579	0.3690	0.106*
H13C	0.1705	0.6695	0.4629	0.106*
C14	-0.0195 (13)	0.8803 (9)	0.4998 (8)	0.069
H14A	-0.0975	0.8056	0.5123	0.104*
H14B	-0.0583	0.9117	0.4286	0.104*
H14C	-0.0483	0.9429	0.5523	0.104*
C15	0.3316 (15)	0.9649 (10)	0.4750 (7)	0.078 (3)
H15A	0.2943	1.0371	0.5156	0.116*
H15B	0.2992	0.9802	0.3993	0.116*
H15C	0.4777	0.9492	0.4883	0.116*
Cl1	0.1156 (5)	0.1156 (2)	0.7961 (2)	0.0908 (9)
N1	0.3167 (7)	0.7456 (5)	0.7848 (4)	0.0336 (11)
O2	0.6980 (6)	0.7001 (6)	0.7256 (4)	0.0534 (12)
H2A	0.8185	0.6858	0.7131	0.080*
O3	0.0721 (6)	0.6749 (5)	0.6573 (3)	0.0398 (10)
O4	0.2916 (7)	0.8324 (5)	0.6216 (3)	0.0448 (11)
S1	0.4357 (3)	0.68989 (18)	0.98363 (12)	0.0481 (5)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.047 (4)	0.049 (3)	0.041 (4)	0.001 (3)	0.003 (3)	0.001 (3)
C2	0.059 (4)	0.055 (4)	0.041 (4)	0.010 (3)	0.001 (3)	-0.006 (3)
C3	0.064 (4)	0.047 (4)	0.053 (4)	-0.006 (3)	-0.014 (3)	0.007 (3)
C4	0.060 (4)	0.057 (4)	0.058 (4)	-0.014 (3)	0.004 (3)	0.004 (4)
C5	0.046 (3)	0.056 (4)	0.045 (4)	-0.004 (3)	0.003 (3)	0.011 (3)
C6	0.039 (3)	0.035 (3)	0.031 (3)	0.000 (2)	0.002 (2)	0.008 (3)
C7	0.035 (3)	0.043 (4)	0.030 (3)	0.005 (3)	0.003 (2)	0.006 (3)
C8	0.042 (3)	0.038 (3)	0.042 (3)	0.001 (3)	-0.004 (3)	0.001 (3)
O1	0.060	0.060	0.060	0.000	0.005	0.000
C9	0.052 (4)	0.052 (4)	0.043 (3)	0.008 (3)	-0.006 (3)	-0.009 (3)
C10	0.026 (3)	0.074 (5)	0.037 (3)	-0.015 (3)	-0.006 (2)	-0.002 (3)
C11	0.028 (3)	0.053 (4)	0.031 (3)	0.009 (3)	-0.002 (2)	-0.002 (3)
C12	0.050 (4)	0.070 (5)	0.035 (3)	-0.008 (4)	-0.003 (3)	0.019 (4)
C13	0.070	0.070	0.070	0.000	0.006	0.000
C14	0.070	0.070	0.070	0.000	0.006	0.000
C15	0.086 (6)	0.085 (7)	0.061 (5)	-0.007 (5)	0.006 (4)	0.036 (5)
Cl1	0.115 (2)	0.0475 (12)	0.1032 (19)	-0.0122 (13)	-0.0338 (16)	0.0021 (13)
N1	0.030 (2)	0.038 (3)	0.033 (2)	0.004 (2)	0.0010 (19)	0.001 (2)
O2	0.032 (2)	0.066 (3)	0.064 (3)	-0.010 (2)	0.012 (2)	-0.018 (3)
O3	0.029 (2)	0.058 (3)	0.0317 (19)	-0.007 (2)	-0.0026 (15)	-0.005 (2)
O4	0.044 (2)	0.055 (3)	0.035 (2)	-0.015 (2)	0.0001 (18)	0.016 (2)
S1	0.0578 (10)	0.0564 (10)	0.0285 (7)	-0.0005 (9)	-0.0071 (6)	-0.0025 (9)

Geometric parameters (\AA , $\text{^{\circ}}$)

C1—C6	1.379 (9)	C9—H9A	0.9700
C1—C2	1.384 (11)	C9—H9B	0.9700
C1—H1	0.9300	C10—O2	1.313 (9)
C2—C3	1.367 (11)	C11—O3	1.234 (8)
C2—H2	0.9300	C11—O4	1.308 (8)
C3—C4	1.378 (12)	C11—N1	1.365 (8)
C3—Cl1	1.744 (8)	C12—O4	1.457 (8)
C4—C5	1.390 (11)	C12—C13	1.483 (12)
C4—H4	0.9300	C12—C14	1.510 (11)
C5—C6	1.380 (9)	C12—C15	1.535 (11)
C5—H5	0.9300	C13—H13A	0.9600
C6—C7	1.490 (9)	C13—H13B	0.9600
C7—N1	1.464 (8)	C13—H13C	0.9600
C7—S1	1.857 (6)	C14—H14A	0.9600
C7—H7	0.9800	C14—H14B	0.9600
C8—N1	1.437 (8)	C14—H14C	0.9600
C8—C10	1.510 (9)	C15—H15A	0.9600
C8—C9	1.510 (9)	C15—H15B	0.9600
C8—H8	0.9800	C15—H15C	0.9600
O1—C10	1.191 (8)	O2—H2A	0.8200
C9—S1	1.801 (8)		
C6—C1—C2	119.0 (7)	O1—C10—O2	123.2 (6)
C6—C1—H1	120.5	O1—C10—C8	122.8 (7)
C2—C1—H1	120.5	O2—C10—C8	114.0 (5)
C3—C2—C1	120.9 (7)	O3—C11—O4	125.6 (5)
C3—C2—H2	119.6	O3—C11—N1	122.1 (6)
C1—C2—H2	119.6	O4—C11—N1	112.3 (5)
C2—C3—C4	121.4 (8)	O4—C12—C13	110.1 (7)
C2—C3—Cl1	119.4 (7)	O4—C12—C14	112.7 (6)
C4—C3—Cl1	119.2 (7)	C13—C12—C14	111.5 (7)
C3—C4—C5	117.2 (8)	O4—C12—C15	102.4 (6)
C3—C4—H4	121.4	C13—C12—C15	110.5 (6)
C5—C4—H4	121.4	C14—C12—C15	109.2 (7)
C6—C5—C4	122.2 (8)	C12—C13—H13A	109.5
C6—C5—H5	118.9	C12—C13—H13B	109.5
C4—C5—H5	118.9	H13A—C13—H13B	109.5
C1—C6—C5	119.3 (6)	C12—C13—H13C	109.5
C1—C6—C7	122.9 (6)	H13A—C13—H13C	109.5
C5—C6—C7	117.8 (6)	H13B—C13—H13C	109.5
N1—C7—C6	117.2 (5)	C12—C14—H14A	109.5
N1—C7—S1	102.2 (4)	C12—C14—H14B	109.5
C6—C7—S1	109.2 (4)	H14A—C14—H14B	109.5
N1—C7—H7	109.3	C12—C14—H14C	109.5
C6—C7—H7	109.3	H14A—C14—H14C	109.5
S1—C7—H7	109.3	H14B—C14—H14C	109.5

N1—C8—C10	115.7 (6)	C12—C15—H15A	109.5
N1—C8—C9	106.7 (5)	C12—C15—H15B	109.5
C10—C8—C9	109.6 (5)	H15A—C15—H15B	109.5
N1—C8—H8	108.2	C12—C15—H15C	109.5
C10—C8—H8	108.2	H15A—C15—H15C	109.5
C9—C8—H8	108.2	H15B—C15—H15C	109.5
C8—C9—S1	104.3 (5)	C11—N1—C8	121.3 (5)
C8—C9—H9A	110.9	C11—N1—C7	119.6 (5)
S1—C9—H9A	110.9	C8—N1—C7	118.1 (5)
C8—C9—H9B	110.9	C10—O2—H2A	109.5
S1—C9—H9B	110.9	C11—O4—C12	121.8 (5)
H9A—C9—H9B	108.9	C9—S1—C7	90.0 (3)
C6—C1—C2—C3	-2.5 (11)	O3—C11—N1—C8	-176.7 (6)
C1—C2—C3—C4	2.1 (12)	O4—C11—N1—C8	3.6 (8)
C1—C2—C3—Cl1	-178.8 (6)	O3—C11—N1—C7	-8.1 (9)
C2—C3—C4—C5	-1.1 (11)	O4—C11—N1—C7	172.2 (5)
Cl1—C3—C4—C5	179.8 (6)	C10—C8—N1—C11	-83.3 (7)
C3—C4—C5—C6	0.5 (11)	C9—C8—N1—C11	154.5 (5)
C2—C1—C6—C5	1.9 (10)	C10—C8—N1—C7	107.9 (6)
C2—C1—C6—C7	178.9 (6)	C9—C8—N1—C7	-14.2 (7)
C4—C5—C6—C1	-0.9 (10)	C6—C7—N1—C11	56.7 (7)
C4—C5—C6—C7	-178.1 (6)	S1—C7—N1—C11	176.0 (4)
C1—C6—C7—N1	42.7 (8)	C6—C7—N1—C8	-134.4 (6)
C5—C6—C7—N1	-140.2 (6)	S1—C7—N1—C8	-15.1 (6)
C1—C6—C7—S1	-72.7 (7)	O3—C11—O4—C12	-0.4 (10)
C5—C6—C7—S1	104.3 (6)	N1—C11—O4—C12	179.3 (6)
N1—C8—C9—S1	37.6 (6)	C13—C12—O4—C11	-67.2 (8)
C10—C8—C9—S1	-88.4 (6)	C14—C12—O4—C11	58.1 (10)
N1—C8—C10—O1	-174.4 (6)	C15—C12—O4—C11	175.3 (6)
C9—C8—C10—O1	-53.8 (9)	C8—C9—S1—C7	-40.6 (5)
N1—C8—C10—O2	3.4 (8)	N1—C7—S1—C9	31.7 (4)
C9—C8—C10—O2	124.0 (6)	C6—C7—S1—C9	156.5 (5)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O2—H2A···O3 ⁱ	0.82	1.83	2.638 (6)	167

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