

Methyl 2-[2-(*tert*-butoxycarbonylamino)-1,3-benzothiazole-6-carboxamido]-acetate

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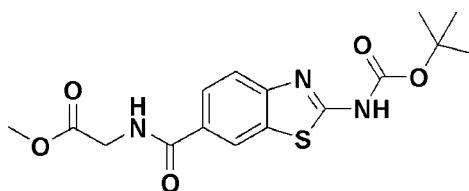
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Key indicators: single-crystal X-ray study; $T = 293\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$; R factor = 0.083; wR factor = 0.180; data-to-parameter ratio = 18.2.

In the title compound, $\text{C}_{16}\text{H}_{19}\text{N}_3\text{O}_5\text{S}$, the dihedral angle between the benzene ring and the carbonylamino group is $18.18(2)^\circ$. In the crystal, molecules form centrosymmetric dimers *via* pairs of $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds. The dimers are connected *via* $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds into a three-dimensional network..

Related literature

For benzothiazole derivatives with anti-tumor activity, see: Brantley *et al.* (2004); Ćaleta *et al.* (2009); Mortimer *et al.* (2006) and for benzothiazolines with anti-tuberculous properties, see: Palmer *et al.* (1971). For related benzothiazole structures, see: Lynch *et al.* (2002); Matković-Čalogović *et al.* (2003); Lei *et al.* (2010).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{19}\text{N}_3\text{O}_5\text{S}$

$M_r = 365.41$

Monoclinic, $P2_1/c$

$a = 16.861(3)\text{ \AA}$

$b = 11.317(2)\text{ \AA}$

$c = 9.6484(19)\text{ \AA}$

$\beta = 98.94(3)^\circ$

$V = 1818.7(6)\text{ \AA}^3$

$Z = 4$

Mo $K\alpha$ radiation

$\mu = 0.21\text{ mm}^{-1}$
 $T = 293\text{ K}$

$0.54 \times 0.33 \times 0.12\text{ mm}$

Data collection

Rigaku Saturn 724 CCD area-detector diffractometer
Absorption correction: numerical (*NUMABS*; Higashi, 2000)
 $T_{\min} = 0.921$, $T_{\max} = 0.975$

14817 measured reflections
4176 independent reflections
3718 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.054$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.083$
 $wR(F^2) = 0.180$
 $S = 1.26$
4176 reflections

230 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.34\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.23\text{ e \AA}^{-3}$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1 \cdots N2 ⁱ	0.86	2.16	3.005 (3)	167
N3—H3 \cdots O3 ⁱⁱ	0.86	2.11	2.802 (3)	137

Symmetry codes: (i) $-x + 1, -y + 1, -z$; (ii) $x, -y + \frac{3}{2}, z + \frac{1}{2}$.

Data collection: *CrystalClear* (Rigaku, 2007); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP* (McArdle, 1995); 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: FF2042).

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Methyl 2-[2-(*tert*-butoxycarbonylamino)-1,3-benzothiazole-6-carboxamido]acetate

D. Gao, X. Fang, H.-Y. Yu and J.-D. Wang

Comment

Benzothiazole is a heterocyclic compound, which contains a benzene ring and a thiazole ring, and is a component of a lot of natural products, biological pesticides, drugs, spices and so on. The benzothiazole derivatives have broad biological activities that make them play an important role in drug research and development. For example, they showed anti-tumor (Brantley *et al.*, 2004; Mortimer *et al.*, 2006; Caleta *et al.*, 2009) and anti-microbial activities (Palmer *et al.*, 1971). During our development of 2-aminobenzothiazole-based Urokinase-Type Plasminogen Activator (uPA) inhibitors, we synthesized the title compound (I) as an intermediate. The compound (I) has certain biological activity, and its IC₅₀ is 780 µM as uPA inhibitor itself.

The molecule structure of the title compound (I) is shown in Fig. 1. The molecular skeleton is slightly distorted from a planar conformation with the angle between benzene and thiazole rings of 1.46 (1)°. And for the substituents, the dihedral angles between the thiazole ring and *tert*-butyl carbamate is 9.15 (6)°, the dihedral angles between benzene ring and carbonylamino group is 18.18 (2)°, and the dihedral angles between carbonylamino group and methyl acetate is 79.24 (3)°.

In the crystal, there are intermolecular hydrogen bonds of N1—H1···N2 and N3—H3···O3. Where, two molecules form a pair with inversion symmetry *via* N—H···N hydrogen bonds, and the pairs form a three dimensional network *via* N—H···O hydrogen bonds. No π—π interactions are found in this structure.

Experimental

In a 250 ml round bottom flask, the pale yellow solid of ethyl 2-(*tert*-butoxycarbonylamino) benzothiazole-6-carboxylate, N-Boc ester (3.22 g, 10 mmol) in a mixed solution of EtOH (100 ml) and 2 N aq NaOH (80 ml) were refluxed for 5 h. Then the solution was cooled with an ice bath and acidified with 1 N aq HCl, when pH<2, white floc generated and put it aside for 2 h. Then the mixture was filtered and the filter mass was washed to neutral by water and dried to afford white solid of 2-(*tert*-butoxycarbonylamino)benzothiazole-6-carboxylic acid, N-Boc acid (2.59 g, yield: 88%).

In a 100 ml round bottom flask, the mixture of N-Boc acid (705 mg, 2.4 mmol), 2-(1*H*-Benzotriazole-1-yl)-1,1,3,3-tetramethyluronium hexafluorophosphate (HBTU, 759 mg, 2 mmol), N, *N*-Diisopropylethylamine (DIEA, 310 mg, 2.4 mmol), and glycine methyl ester hydrochloride (251 mg, 2 mmol) in 20 ml dry DMF were stirred at room temperature for 20 h. Then the reaction solution was pured into 200 mL of 10% Na₂CO₃ solution and stirred for 1 h. The precipitate was filtered, washed with water, and dried to give the white solid of title compound (I) (591 mg, yield: 81%).

The solid was dissolved by DMF and filtered. The DMF was evaporated slowly at room temperature for 15–20 days, and colorless sheetlike crystals suitable for X-ray structure analysis were separated from the solution.

supplementary materials

Refinement

All hydrogen atoms were positioned geometrically and refined in a riding model approximation with U_{iso} (H) = 1.2 or 1.5 U_{eq} (C).

Figures

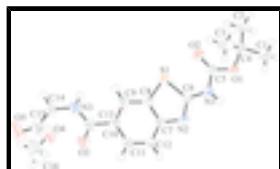


Fig. 1. A view of the molecular structure of (I), with displacement ellipsoids drawn at the 50% probability level for non-H atoms.

Methyl 2-[2-(tert-butoxycarbonylamino)-1,3-benzothiazole-6-carboxamido]acetate

Crystal data

C ₁₆ H ₁₉ N ₃ O ₅ S	$F(000) = 768.0$
$M_r = 365.41$	$D_x = 1.335 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Hall symbol: -P 2ybc	Cell parameters from 5508 reflections
$a = 16.861 (3) \text{ \AA}$	$\theta = 3.0\text{--}27.5^\circ$
$b = 11.317 (2) \text{ \AA}$	$\mu = 0.21 \text{ mm}^{-1}$
$c = 9.6484 (19) \text{ \AA}$	$T = 293 \text{ K}$
$\beta = 98.94 (3)^\circ$	Prism, colourless
$V = 1818.7 (6) \text{ \AA}^3$	$0.54 \times 0.33 \times 0.12 \text{ mm}$
$Z = 4$	

Data collection

Rigaku Saturn 724 CCD area-detector diffractometer	4176 independent reflections
Radiation source: fine-focus sealed tube graphite	3718 reflections with $I > 2\sigma(I)$
Detector resolution: 28.5714 pixels mm^{-1}	$R_{\text{int}} = 0.054$
dtp010.ref scans	$\theta_{\text{max}} = 27.5^\circ, \theta_{\text{min}} = 3.0^\circ$
Absorption correction: numerical (<i>NUMABS</i> ; Higashi, 2000)	$h = -20 \rightarrow 21$
$T_{\text{min}} = 0.921, T_{\text{max}} = 0.975$	$k = -14 \rightarrow 14$
14817 measured reflections	$l = -12 \rightarrow 12$

Refinement

Refinement on F^2	Primary atom site location: structure-invariant direct methods
Least-squares matrix: full	Secondary atom site location: difference Fourier map

$R[F^2 > 2\sigma(F^2)] = 0.083$	Hydrogen site location: inferred from neighbouring sites
$wR(F^2) = 0.180$	H-atom parameters constrained
$S = 1.26$	$w = 1/[\sigma^2(F_o^2) + (0.0486P)^2 + 1.240P]$ where $P = (F_o^2 + 2F_c^2)/3$
4176 reflections	$(\Delta/\sigma)_{\max} < 0.001$
230 parameters	$\Delta\rho_{\max} = 0.34 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\min} = -0.23 \text{ e } \text{\AA}^{-3}$

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	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.36614 (4)	0.72248 (7)	0.16296 (8)	0.0467 (2)
O3	0.05637 (13)	0.7065 (2)	-0.3160 (2)	0.0637 (7)
O4	-0.06380 (16)	0.6314 (3)	-0.0995 (3)	0.0789 (8)
O2	0.49650 (13)	0.7178 (2)	0.3633 (2)	0.0574 (6)
O1	0.60458 (12)	0.59926 (19)	0.3591 (2)	0.0479 (5)
O5	-0.16027 (18)	0.7431 (4)	-0.2127 (5)	0.1271 (15)
N1	0.50500 (13)	0.5954 (2)	0.1795 (2)	0.0415 (6)
H1	0.5363	0.5490	0.1426	0.050*
N2	0.40231 (13)	0.5568 (2)	-0.0059 (2)	0.0377 (5)
N3	0.05095 (15)	0.7905 (3)	-0.1077 (3)	0.0524 (7)
H3	0.0756	0.8095	-0.0259	0.063*
C9	0.21789 (16)	0.7285 (3)	-0.0217 (3)	0.0431 (7)
H9	0.1985	0.7890	0.0290	0.052*
C3	0.5917 (3)	0.5968 (4)	0.6084 (4)	0.0761 (12)
H3A	0.5441	0.6448	0.5945	0.114*
H3B	0.6199	0.6089	0.7017	0.114*
H3C	0.5770	0.5151	0.5962	0.114*
C15	-0.0922 (2)	0.7299 (4)	-0.1594 (4)	0.0696 (11)
C16	-0.1190 (3)	0.5336 (5)	-0.1009 (7)	0.133 (2)
H16A	-0.1640	0.5574	-0.0578	0.199*
H16B	-0.0922	0.4684	-0.0498	0.199*
H16C	-0.1373	0.5100	-0.1960	0.199*
C1	0.7200 (2)	0.5557 (4)	0.5209 (4)	0.0855 (14)
H1A	0.7053	0.4740	0.5087	0.128*

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H1B	0.7497	0.5673	0.6132	0.128*
H1C	0.7526	0.5781	0.4521	0.128*
C6	0.43024 (16)	0.6160 (2)	0.1075 (3)	0.0372 (6)
C7	0.32479 (16)	0.5921 (2)	-0.0557 (3)	0.0363 (6)
C10	0.17080 (16)	0.6823 (3)	-0.1401 (3)	0.0409 (6)
C8	0.29461 (16)	0.6830 (2)	0.0201 (3)	0.0384 (6)
C12	0.27640 (17)	0.5443 (3)	-0.1727 (3)	0.0427 (7)
H12	0.2951	0.4828	-0.2228	0.051*
C11	0.20071 (17)	0.5895 (3)	-0.2130 (3)	0.0440 (7)
H11	0.1684	0.5575	-0.2909	0.053*
C13	0.08877 (18)	0.7271 (3)	-0.1946 (3)	0.0460 (7)
C5	0.53326 (17)	0.6445 (3)	0.3083 (3)	0.0419 (6)
C4	0.64493 (19)	0.6309 (3)	0.5033 (3)	0.0484 (7)
C14	-0.03095 (19)	0.8269 (4)	-0.1511 (4)	0.0637 (10)
H14A	-0.0350	0.8638	-0.2427	0.076*
H14B	-0.0441	0.8864	-0.0860	0.076*
C2	0.6663 (2)	0.7601 (3)	0.5084 (4)	0.0668 (10)
H2A	0.6974	0.7777	0.4356	0.100*
H2B	0.6972	0.7785	0.5980	0.100*
H2C	0.6181	0.8065	0.4947	0.100*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0417 (4)	0.0507 (5)	0.0453 (4)	0.0087 (3)	-0.0005 (3)	-0.0150 (3)
O3	0.0521 (13)	0.096 (2)	0.0399 (12)	0.0100 (13)	-0.0022 (10)	-0.0002 (12)
O4	0.0595 (16)	0.095 (2)	0.0791 (18)	-0.0064 (15)	0.0023 (14)	0.0225 (16)
O2	0.0514 (13)	0.0652 (15)	0.0529 (13)	0.0140 (11)	-0.0002 (10)	-0.0218 (11)
O1	0.0480 (11)	0.0523 (13)	0.0397 (11)	0.0121 (9)	-0.0045 (9)	-0.0131 (9)
O5	0.0533 (18)	0.141 (3)	0.172 (4)	0.0003 (19)	-0.029 (2)	0.035 (3)
N1	0.0374 (12)	0.0463 (14)	0.0399 (12)	0.0078 (10)	0.0028 (10)	-0.0081 (10)
N2	0.0387 (12)	0.0367 (12)	0.0376 (12)	0.0003 (9)	0.0053 (10)	-0.0051 (10)
N3	0.0409 (14)	0.0684 (19)	0.0456 (14)	0.0096 (12)	-0.0003 (11)	-0.0011 (13)
C9	0.0388 (15)	0.0472 (17)	0.0439 (15)	0.0045 (12)	0.0080 (12)	-0.0043 (13)
C3	0.103 (3)	0.079 (3)	0.0425 (19)	-0.007 (2)	0.003 (2)	0.0012 (19)
C15	0.0423 (19)	0.099 (3)	0.065 (2)	0.0063 (19)	0.0001 (16)	0.007 (2)
C16	0.097 (4)	0.135 (5)	0.160 (6)	-0.038 (4)	0.000 (4)	0.054 (5)
C1	0.085 (3)	0.092 (3)	0.067 (2)	0.040 (2)	-0.029 (2)	-0.028 (2)
C6	0.0377 (14)	0.0371 (14)	0.0367 (14)	0.0016 (11)	0.0049 (11)	-0.0013 (11)
C7	0.0362 (14)	0.0352 (14)	0.0377 (14)	-0.0004 (11)	0.0064 (11)	-0.0003 (11)
C10	0.0381 (14)	0.0473 (17)	0.0374 (14)	-0.0010 (12)	0.0064 (11)	0.0013 (12)
C8	0.0372 (14)	0.0384 (15)	0.0392 (14)	0.0007 (11)	0.0050 (11)	-0.0016 (12)
C12	0.0445 (16)	0.0406 (16)	0.0428 (15)	0.0016 (12)	0.0059 (12)	-0.0043 (13)
C11	0.0441 (16)	0.0492 (17)	0.0371 (14)	-0.0039 (13)	0.0013 (12)	-0.0023 (13)
C13	0.0409 (16)	0.0563 (19)	0.0400 (15)	0.0027 (13)	0.0037 (12)	0.0069 (14)
C5	0.0397 (15)	0.0447 (16)	0.0407 (15)	0.0016 (12)	0.0048 (12)	-0.0061 (13)
C4	0.0539 (18)	0.0501 (18)	0.0372 (15)	0.0083 (14)	-0.0057 (13)	-0.0104 (13)
C14	0.0473 (18)	0.074 (3)	0.067 (2)	0.0189 (17)	0.0000 (16)	0.0019 (19)

C2	0.065 (2)	0.059 (2)	0.071 (2)	-0.0095 (17)	-0.0060 (18)	-0.0116 (19)
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Geometric parameters (\AA , $^\circ$)

S1—C8	1.742 (3)	C3—H3C	0.9600
S1—C6	1.757 (3)	C15—C14	1.501 (6)
O3—C13	1.235 (4)	C16—H16A	0.9600
O4—C15	1.312 (5)	C16—H16B	0.9600
O4—C16	1.444 (5)	C16—H16C	0.9600
O2—C5	1.207 (3)	C1—C4	1.513 (4)
O1—C5	1.329 (3)	C1—H1A	0.9600
O1—C4	1.494 (3)	C1—H1B	0.9600
O5—C15	1.191 (4)	C1—H1C	0.9600
N1—C6	1.362 (3)	C7—C12	1.395 (4)
N1—C5	1.377 (3)	C7—C8	1.403 (4)
N1—H1	0.8600	C10—C11	1.401 (4)
N2—C6	1.307 (3)	C10—C13	1.490 (4)
N2—C7	1.380 (3)	C12—C11	1.373 (4)
N3—C13	1.338 (4)	C12—H12	0.9300
N3—C14	1.440 (4)	C11—H11	0.9300
N3—H3	0.8600	C4—C2	1.505 (5)
C9—C10	1.389 (4)	C14—H14A	0.9700
C9—C8	1.393 (4)	C14—H14B	0.9700
C9—H9	0.9300	C2—H2A	0.9600
C3—C4	1.506 (5)	C2—H2B	0.9600
C3—H3A	0.9600	C2—H2C	0.9600
C3—H3B	0.9600		
C8—S1—C6	88.06 (13)	N2—C7—C8	115.6 (2)
C15—O4—C16	117.2 (3)	C12—C7—C8	119.5 (2)
C5—O1—C4	120.4 (2)	C9—C10—C11	119.4 (3)
C6—N1—C5	123.7 (2)	C9—C10—C13	122.9 (3)
C6—N1—H1	118.2	C11—C10—C13	117.7 (3)
C5—N1—H1	118.2	C9—C8—C7	121.1 (3)
C6—N2—C7	110.0 (2)	C9—C8—S1	129.2 (2)
C13—N3—C14	120.0 (3)	C7—C8—S1	109.7 (2)
C13—N3—H3	120.0	C11—C12—C7	119.0 (3)
C14—N3—H3	120.0	C11—C12—H12	120.5
C10—C9—C8	119.1 (3)	C7—C12—H12	120.5
C10—C9—H9	120.5	C12—C11—C10	121.9 (3)
C8—C9—H9	120.5	C12—C11—H11	119.1
C4—C3—H3A	109.5	C10—C11—H11	119.1
C4—C3—H3B	109.5	O3—C13—N3	120.8 (3)
H3A—C3—H3B	109.5	O3—C13—C10	121.3 (3)
C4—C3—H3C	109.5	N3—C13—C10	117.9 (3)
H3A—C3—H3C	109.5	O2—C5—O1	126.9 (3)
H3B—C3—H3C	109.5	O2—C5—N1	123.0 (3)
O5—C15—O4	123.9 (4)	O1—C5—N1	110.1 (2)
O5—C15—C14	122.7 (4)	O1—C4—C2	109.6 (3)
O4—C15—C14	113.4 (3)	O1—C4—C3	109.4 (3)

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O4—C16—H16A	109.5	C2—C4—C3	113.1 (3)
O4—C16—H16B	109.5	O1—C4—C1	102.8 (2)
H16A—C16—H16B	109.5	C2—C4—C1	110.5 (3)
O4—C16—H16C	109.5	C3—C4—C1	110.9 (3)
H16A—C16—H16C	109.5	N3—C14—C15	115.3 (3)
H16B—C16—H16C	109.5	N3—C14—H14A	108.4
C4—C1—H1A	109.5	C15—C14—H14A	108.4
C4—C1—H1B	109.5	N3—C14—H14B	108.4
H1A—C1—H1B	109.5	C15—C14—H14B	108.4
C4—C1—H1C	109.5	H14A—C14—H14B	107.5
H1A—C1—H1C	109.5	C4—C2—H2A	109.5
H1B—C1—H1C	109.5	C4—C2—H2B	109.5
N2—C6—N1	121.6 (2)	H2A—C2—H2B	109.5
N2—C6—S1	116.6 (2)	C4—C2—H2C	109.5
N1—C6—S1	121.8 (2)	H2A—C2—H2C	109.5
N2—C7—C12	124.9 (2)	H2B—C2—H2C	109.5
C16—O4—C15—O5	0.7 (7)	C8—C7—C12—C11	1.5 (4)
C16—O4—C15—C14	179.2 (4)	C7—C12—C11—C10	0.1 (4)
C7—N2—C6—N1	178.2 (2)	C9—C10—C11—C12	-1.6 (5)
C7—N2—C6—S1	-1.2 (3)	C13—C10—C11—C12	178.3 (3)
C5—N1—C6—N2	-171.5 (3)	C14—N3—C13—O3	5.3 (5)
C5—N1—C6—S1	7.8 (4)	C14—N3—C13—C10	-174.5 (3)
C8—S1—C6—N2	0.2 (2)	C9—C10—C13—O3	161.8 (3)
C8—S1—C6—N1	-179.1 (3)	C11—C10—C13—O3	-18.2 (5)
C6—N2—C7—C12	-178.0 (3)	C9—C10—C13—N3	-18.4 (5)
C6—N2—C7—C8	1.8 (3)	C11—C10—C13—N3	161.7 (3)
C8—C9—C10—C11	1.4 (4)	C4—O1—C5—O2	5.0 (5)
C8—C9—C10—C13	-178.6 (3)	C4—O1—C5—N1	-174.9 (2)
C10—C9—C8—C7	0.3 (4)	C6—N1—C5—O2	-6.2 (5)
C10—C9—C8—S1	-179.6 (2)	C6—N1—C5—O1	173.7 (3)
N2—C7—C8—C9	178.4 (3)	C5—O1—C4—C2	-65.7 (4)
C12—C7—C8—C9	-1.8 (4)	C5—O1—C4—C3	58.8 (4)
N2—C7—C8—S1	-1.7 (3)	C5—O1—C4—C1	176.7 (3)
C12—C7—C8—S1	178.2 (2)	C13—N3—C14—C15	71.6 (4)
C6—S1—C8—C9	-179.3 (3)	O5—C15—C14—N3	-168.8 (4)
C6—S1—C8—C7	0.8 (2)	O4—C15—C14—N3	12.7 (5)
N2—C7—C12—C11	-178.6 (3)		

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

$D—\text{H}\cdots A$	$D—\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D—\text{H}\cdots A$
N1—H1 ⁱ —N2 ⁱ	0.86	2.16	3.005 (3)	167.
N3—H3 ⁱⁱ —O3 ⁱⁱ	0.86	2.11	2.802 (3)	137.

Symmetry codes: (i) $-x+1, -y+1, -z$; (ii) $x, -y+3/2, z+1/2$.

Fig. 1

