

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

2-(2-Nitrophenyl)-1,3-benzothiazole

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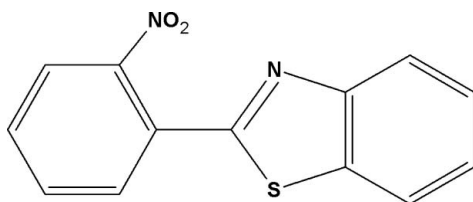
Received 25 June 2012; accepted 30 June 2012

Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.039; wR factor = 0.113; data-to-parameter ratio = 20.0.

In the title compound, $\text{C}_{13}\text{H}_8\text{N}_2\text{O}_2\text{S}$, the essentially planar benzothiazole system [maximum deviation = -0.012 (1) Å for the S atom] is oriented at a dihedral angle of 48.3 (1)° with respect to the benzene ring. The nitro group is substantially twisted from the plane of its attached benzene ring [dihedral angle = 52.0 (1)°]. The crystal packing features $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, which generate $C(6)$ helical chains propagating along [010]. Weak $\text{C}-\text{H}\cdots\pi$ interactions also occur in the crystal.

Related literature

For the pharmacological activity of benzothiazole derivatives, see: Repić *et al.* (2001); Schwartz *et al.* (1992). For related structures, see: Lakshmanan *et al.* (2011); Zhang *et al.* (2008).



Experimental

Crystal data

$\text{C}_{13}\text{H}_8\text{N}_2\text{O}_2\text{S}$
 $M_r = 256.27$
 Monoclinic, $P2_1/c$

$a = 7.6092$ (2) Å
 $b = 12.7854$ (3) Å
 $c = 11.9938$ (3) Å

$\beta = 90.556$ (2)°
 $V = 1166.78$ (5) Å³
 $Z = 4$
 Mo $K\alpha$ radiation

$\mu = 0.27$ mm⁻¹
 $T = 293$ K
 $0.24 \times 0.22 \times 0.16$ mm

Data collection

Bruker APEXII CCD diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.937$, $T_{\max} = 0.958$

14037 measured reflections
 3258 independent reflections
 2559 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.039$
 $wR(F^2) = 0.113$
 $S = 1.05$
 3258 reflections

163 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.24$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.32$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$Cg1$, $Cg2$ and $Cg3$ are the centroids of the S1/N1/C1/C2/C7 thiazole ring, the C2–C7 benzene ring and the C8–C13 benzene ring, respectively.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C11-H11\cdots O1^i$	0.93	2.51	3.236 (2)	135
$C9-H9\cdots Cg1^{ii}$	0.93	2.92	3.468 (2)	119
$C10-H10\cdots Cg2^{ii}$	0.93	2.90	3.536 (2)	127
$C3-H3\cdots Cg3^{iii}$	0.93	2.99	3.673 (2)	132

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

Data collection: APEX2 (Bruker, 2004); cell refinement: APEX2 and SAINT (Bruker, 2004); data reduction: SAINT and XPREP (Bruker, 2004); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 (Farrugia (1997)); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2009).

SM thank Dr Babu Vargheese, SAIF, IIT, Madras, India, for his help with the data collection.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6879).

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supplementary materials

Acta Cryst. (2012). E68, o2362 [doi:10.1107/S1600536812029844]

2-(2-Nitrophenyl)-1,3-benzothiazole

S. Vijayakumar, S. Murugavel, R. Selvakumar and M. Bakthadoss

Comment

The benzothiazole nucleus is associated with several pharmacological activities such as anti-tumor (Repić *et al.*, 2001) and antimicrobial (Schwartz *et al.*, 1992). As part of our studies in this area, the crystal structure of the title compound has been determined and the results are presented here.

Fig. 1. shows a displacement ellipsoid plot of (I), with the atom numbering scheme. The benzothiazole moiety (S1/N1/C1—C7) is essentially planar [maximum deviation = -0.012 (1) Å for the S atom] and lies at an angle 48.3 (1)° with respect to the benzene ring. The nitro group (N2/O1/O2) is twisted from the attached benzene ring, forming a dihedral angle of 52.0 (1)°. The geometric parameters of the title molecule agrees well with those reported for similar structures (Lakshmanan *et al.*, 2011, Zhang *et al.*, 2008).

The crystal packing features C—H...O hydrogen bonds. Atom C11 at x, y, z donates one proton to atom O1 at $1 - x, -1/2 + y, 3/2 - z$, forming C(6) zigzag chains along the b axis (Fig. 2). The crystal packing also features three weak C—H... π interactions, the first one between a benzene H9 atom and the thiazole ring (S1/N1/C1/C2/C7) of an adjacent molecule, with a C9—H9...Cg1ⁱⁱ separation of 2.92 Å, the second one between a benzene H10 atom and the benzene ring (C2—C7) of a neighbouring molecule, with a C10—H10...Cg2ⁱⁱ separation of 2.90 Å and the third one between a benzene H3 atom and the benzene ring (C8—C13) of a neighbouring molecule, with a C3—H3...Cg3ⁱⁱⁱ separation of 2.99 Å (Table 1 and Fig. 3; Cg1, Cg2 and Cg3 are the centroids of the (S1/N1/C1/C2/C7) thiazole ring, (C2—C7) benzene ring and (C8—C13) benzene ring, respectively. symmetry code as in Fig. 3).

Experimental

A mixture of 2-nitrobenzaldehyde (1 g, 6.6 mmol), 2-aminobenzenethiol (0.827 g, 6.6 mmol) and bakers' yeast (2.05 g) were stirred at room temperature for 24 h in dichloro methane(DCM). After completion of the reaction, the bakers' yeast was filtered through a bed of Celite, and the filtrate was concentrated under reduced pressure. On cooling, the solid product (1.60 g, 94%) obtained was separated and crystallized from ethylacetate to afford the title compound as yellow blocks.

Refinement

All the H atoms were positioned geometrically, with C—H = 0.93–0.96 Å and constrained to ride on their parent atom, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Computing details

Data collection: *APEX2* (Bruker, 2004); cell refinement: *APEX2* and *SAINT* (Bruker, 2004); data reduction: *SAINT* and *XPREP* (Bruker, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia (1997); software used to prepare material for publication: *SHELXL97* (Sheldrick, 2008) and *PLATON* (Spek, 2009).

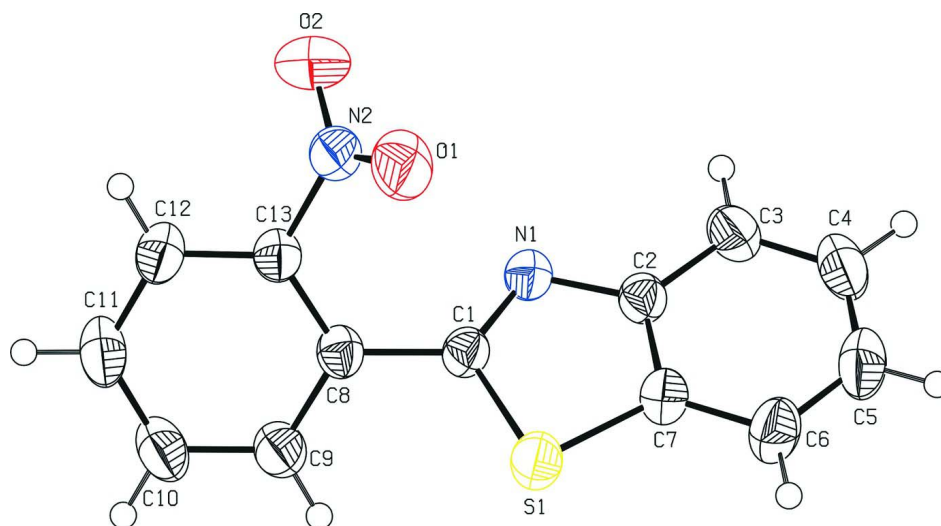


Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level.

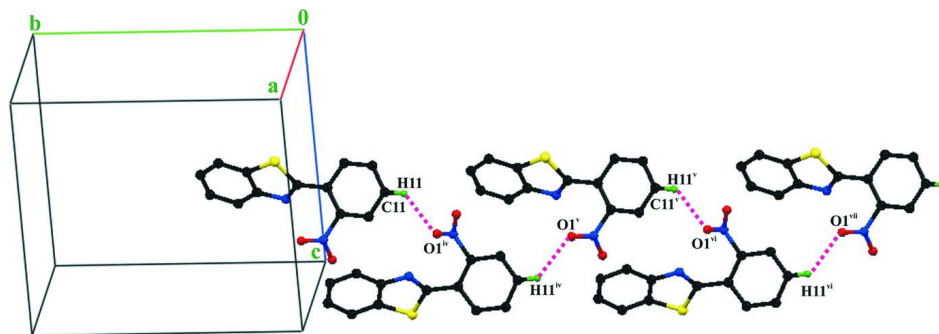
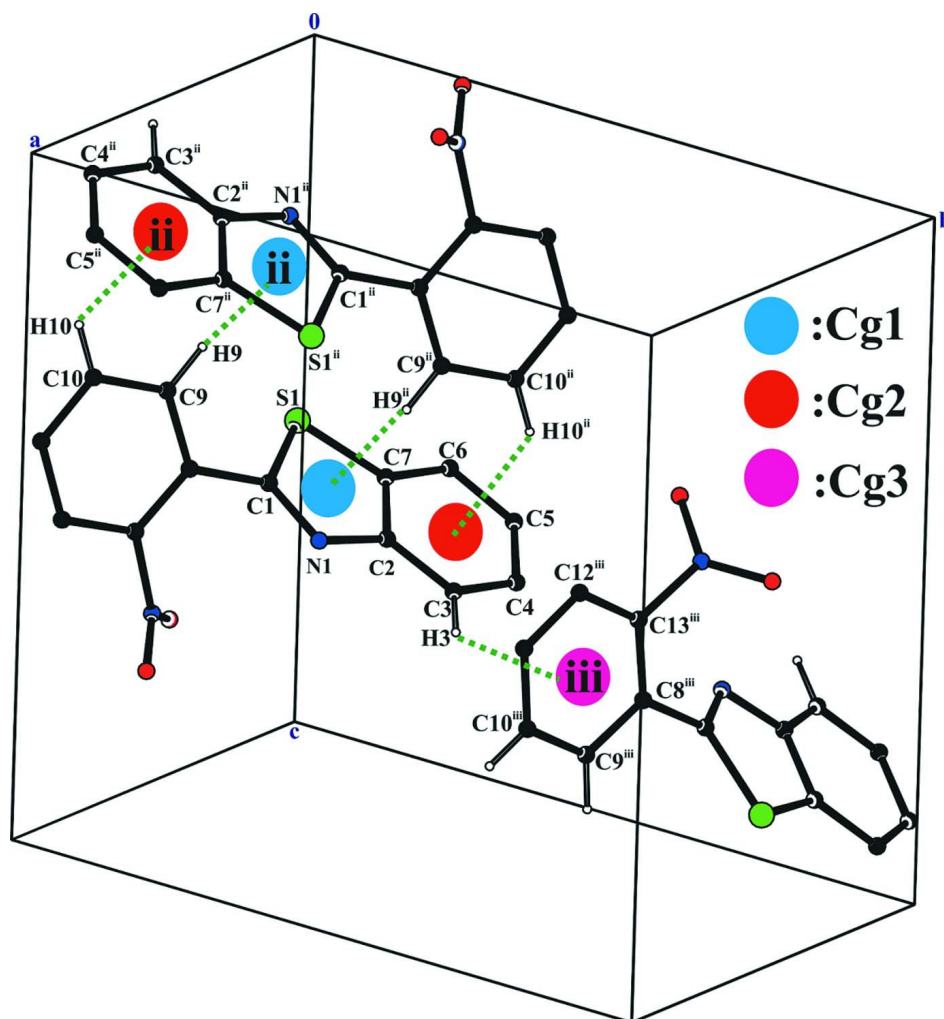


Figure 2

Part of the crystal structure of (I) showing intermolecular C—H...O hydrogen bonds (dotted lines), forming C(6) zigzag chains along the *b* axis. For clarity H atoms involved in the hydrogen bonds are shown. [Symmetry codes:(iv) $1 - x, -1/2 + y, 3/2 - z$; (v) $x, -1 + y, z$; (vi) $1 - x, -3/2 + y, 3/2 - z$; (vii) $x, -2 + y, z$].


Figure 3

A view of the C—H \cdots π interactions (dotted lines) in the crystal structure of the title compound. Cg1, Cg2 and Cg3 denotes centroid of the S1/N1/C1/C2/C7 thiazole ring, C2–C7 benzene ring and C8–C13 benzene ring, respectively. [Symmetry codes: (ii)- x , $-y$, $1 - z$; (iii)- x , $1/2 + y$, $3/2 - z$].

2-(2-Nitrophenyl)-1,3-benzothiazole

Crystal data

$C_{13}H_8N_2O_2S$

$M_r = 256.27$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2_1/c$

$a = 7.6092$ (2) Å

$b = 12.7854$ (3) Å

$c = 11.9938$ (3) Å

$\beta = 90.556$ (2)°

$V = 1166.78$ (5) Å³

$Z = 4$

$F(000) = 528$

$D_x = 1.459$ Mg m⁻³

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

Cell parameters from 3261 reflections

$\theta = 2.3$ – 29.5 °

$\mu = 0.27$ mm⁻¹

$T = 293$ K

Block, yellow

$0.24 \times 0.22 \times 0.16$ mm

Data collection

Bruker APEXII CCD diffractometer	14037 measured reflections 3258 independent reflections
Radiation source: fine-focus sealed tube	2559 reflections with $I > 2\sigma(I)$
Graphite monochromator	$R_{\text{int}} = 0.027$
Detector resolution: 10.0 pixels mm^{-1}	$\theta_{\text{max}} = 29.5^\circ$, $\theta_{\text{min}} = 2.3^\circ$
ω scans	$h = -6 \rightarrow 10$
Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 1996)	$k = -17 \rightarrow 17$
$T_{\text{min}} = 0.937$, $T_{\text{max}} = 0.958$	$l = -16 \rightarrow 16$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.039$	H-atom parameters constrained
$wR(F^2) = 0.113$	$w = 1/[\sigma^2(F_o^2) + (0.0565P)^2 + 0.2357P]$
$S = 1.05$	where $P = (F_o^2 + 2F_c^2)/3$
3258 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
163 parameters	$\Delta\rho_{\text{max}} = 0.24 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.32 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.23422 (17)	0.05259 (10)	0.59828 (11)	0.0353 (3)
C2	0.16130 (18)	0.21138 (10)	0.64971 (11)	0.0363 (3)
C3	0.0966 (2)	0.29175 (12)	0.71628 (13)	0.0480 (3)
H3	0.0530	0.2776	0.7869	0.058*
C4	0.0984 (2)	0.39208 (12)	0.67587 (16)	0.0548 (4)
H4	0.0561	0.4462	0.7199	0.066*
C5	0.1620 (2)	0.41411 (12)	0.57096 (16)	0.0560 (4)
H5	0.1621	0.4830	0.5460	0.067*
C6	0.2249 (2)	0.33699 (12)	0.50273 (15)	0.0531 (4)
H6	0.2667	0.3523	0.4320	0.064*
C7	0.22399 (19)	0.23476 (11)	0.54327 (12)	0.0404 (3)
C8	0.26409 (17)	-0.06135 (10)	0.60273 (11)	0.0362 (3)
C9	0.2087 (2)	-0.12556 (11)	0.51565 (13)	0.0465 (3)
H9	0.1533	-0.0963	0.4536	0.056*
C10	0.2349 (2)	-0.23238 (12)	0.52027 (15)	0.0542 (4)
H10	0.1979	-0.2743	0.4612	0.065*

C11	0.3152 (2)	-0.27701 (11)	0.61136 (16)	0.0546 (4)
H11	0.3325	-0.3490	0.6137	0.065*
C12	0.3705 (2)	-0.21558 (11)	0.69957 (14)	0.0470 (3)
H12	0.4235	-0.2455	0.7621	0.056*
C13	0.34560 (18)	-0.10908 (10)	0.69301 (11)	0.0376 (3)
N1	0.16909 (16)	0.10658 (8)	0.67886 (9)	0.0385 (3)
N2	0.42054 (19)	-0.04475 (10)	0.78230 (11)	0.0482 (3)
O1	0.51720 (18)	0.02623 (10)	0.75526 (12)	0.0675 (4)
O2	0.3857 (2)	-0.06687 (13)	0.87769 (10)	0.0812 (4)
S1	0.29145 (6)	0.12103 (3)	0.47927 (3)	0.04852 (13)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0351 (6)	0.0335 (6)	0.0372 (6)	0.0036 (5)	-0.0008 (5)	0.0019 (5)
C2	0.0360 (7)	0.0340 (6)	0.0389 (6)	0.0019 (5)	-0.0033 (5)	-0.0014 (5)
C3	0.0543 (9)	0.0430 (7)	0.0469 (8)	0.0053 (6)	0.0005 (7)	-0.0092 (6)
C4	0.0577 (10)	0.0379 (7)	0.0687 (10)	0.0062 (7)	-0.0064 (8)	-0.0153 (7)
C5	0.0586 (10)	0.0316 (7)	0.0776 (11)	-0.0010 (7)	-0.0063 (8)	0.0036 (7)
C6	0.0607 (10)	0.0383 (8)	0.0605 (9)	-0.0017 (7)	0.0062 (8)	0.0109 (7)
C7	0.0416 (7)	0.0338 (6)	0.0459 (7)	0.0012 (5)	0.0020 (6)	0.0019 (5)
C8	0.0339 (6)	0.0326 (6)	0.0422 (7)	0.0034 (5)	-0.0006 (5)	-0.0012 (5)
C9	0.0464 (8)	0.0423 (7)	0.0506 (8)	0.0074 (6)	-0.0117 (7)	-0.0064 (6)
C10	0.0537 (9)	0.0404 (8)	0.0681 (11)	0.0038 (7)	-0.0123 (8)	-0.0167 (7)
C11	0.0561 (9)	0.0309 (7)	0.0766 (11)	0.0033 (6)	-0.0052 (8)	-0.0038 (7)
C12	0.0489 (8)	0.0362 (7)	0.0559 (8)	0.0048 (6)	-0.0032 (7)	0.0064 (6)
C13	0.0373 (7)	0.0336 (6)	0.0417 (7)	0.0013 (5)	-0.0006 (5)	-0.0005 (5)
N1	0.0442 (6)	0.0345 (5)	0.0368 (5)	0.0048 (4)	0.0009 (5)	0.0007 (4)
N2	0.0552 (8)	0.0416 (6)	0.0476 (7)	0.0074 (6)	-0.0130 (6)	-0.0025 (5)
O1	0.0702 (8)	0.0496 (7)	0.0822 (9)	-0.0119 (6)	-0.0263 (7)	-0.0049 (6)
O2	0.1183 (13)	0.0836 (10)	0.0414 (6)	0.0026 (9)	-0.0075 (7)	-0.0013 (6)
S1	0.0620 (3)	0.0411 (2)	0.0428 (2)	0.00900 (16)	0.01624 (17)	0.00421 (14)

Geometric parameters (Å, °)

C1—N1	1.2906 (17)	C7—S1	1.7247 (14)
C1—C8	1.4752 (18)	C8—C13	1.3844 (19)
C1—S1	1.7335 (13)	C8—C9	1.391 (2)
C2—N1	1.3858 (17)	C9—C10	1.381 (2)
C2—C3	1.3940 (19)	C9—H9	0.9300
C2—C7	1.400 (2)	C10—C11	1.371 (2)
C3—C4	1.371 (2)	C10—H10	0.9300
C3—H3	0.9300	C11—C12	1.380 (2)
C4—C5	1.381 (3)	C11—H11	0.9300
C4—H4	0.9300	C12—C13	1.3770 (18)
C5—C6	1.371 (3)	C12—H12	0.9300
C5—H5	0.9300	C13—N2	1.4617 (18)
C6—C7	1.395 (2)	N2—O2	1.2103 (18)
C6—H6	0.9300	N2—O1	1.2142 (19)

N1—C1—C8	124.15 (12)	C13—C8—C1	122.09 (12)
N1—C1—S1	116.61 (10)	C9—C8—C1	120.68 (12)
C8—C1—S1	119.24 (10)	C10—C9—C8	120.72 (14)
N1—C2—C3	125.64 (13)	C10—C9—H9	119.6
N1—C2—C7	115.00 (11)	C8—C9—H9	119.6
C3—C2—C7	119.36 (13)	C11—C10—C9	120.44 (15)
C4—C3—C2	118.84 (15)	C11—C10—H10	119.8
C4—C3—H3	120.6	C9—C10—H10	119.8
C2—C3—H3	120.6	C10—C11—C12	120.29 (14)
C3—C4—C5	121.16 (15)	C10—C11—H11	119.9
C3—C4—H4	119.4	C12—C11—H11	119.9
C5—C4—H4	119.4	C13—C12—C11	118.56 (14)
C6—C5—C4	121.63 (15)	C13—C12—H12	120.7
C6—C5—H5	119.2	C11—C12—H12	120.7
C4—C5—H5	119.2	C12—C13—C8	122.75 (13)
C5—C6—C7	117.59 (16)	C12—C13—N2	117.52 (13)
C5—C6—H6	121.2	C8—C13—N2	119.58 (11)
C7—C6—H6	121.2	C1—N1—C2	110.13 (11)
C6—C7—C2	121.41 (14)	O2—N2—O1	124.46 (15)
C6—C7—S1	129.23 (12)	O2—N2—C13	118.28 (14)
C2—C7—S1	109.36 (10)	O1—N2—C13	117.25 (13)
C13—C8—C9	117.22 (12)	C7—S1—C1	88.91 (6)
N1—C2—C3—C4	179.67 (15)	C10—C11—C12—C13	-1.0 (3)
C7—C2—C3—C4	-0.8 (2)	C11—C12—C13—C8	1.4 (2)
C2—C3—C4—C5	0.3 (3)	C11—C12—C13—N2	-174.13 (15)
C3—C4—C5—C6	0.4 (3)	C9—C8—C13—C12	-0.9 (2)
C4—C5—C6—C7	-0.5 (3)	C1—C8—C13—C12	178.36 (14)
C5—C6—C7—C2	0.0 (2)	C9—C8—C13—N2	174.55 (14)
C5—C6—C7—S1	179.49 (13)	C1—C8—C13—N2	-6.2 (2)
N1—C2—C7—C6	-179.77 (14)	C8—C1—N1—C2	178.70 (12)
C3—C2—C7—C6	0.7 (2)	S1—C1—N1—C2	-0.41 (15)
N1—C2—C7—S1	0.64 (16)	C3—C2—N1—C1	179.35 (14)
C3—C2—C7—S1	-178.90 (11)	C7—C2—N1—C1	-0.16 (17)
N1—C1—C8—C13	-47.3 (2)	C12—C13—N2—O2	-52.8 (2)
S1—C1—C8—C13	131.81 (12)	C8—C13—N2—O2	131.55 (16)
N1—C1—C8—C9	131.96 (15)	C12—C13—N2—O1	125.64 (16)
S1—C1—C8—C9	-48.95 (18)	C8—C13—N2—O1	-50.05 (19)
C13—C8—C9—C10	0.0 (2)	C6—C7—S1—C1	179.76 (16)
C1—C8—C9—C10	-179.32 (14)	C2—C7—S1—C1	-0.69 (11)
C8—C9—C10—C11	0.4 (3)	N1—C1—S1—C7	0.67 (12)
C9—C10—C11—C12	0.1 (3)	C8—C1—S1—C7	-178.49 (11)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C11—H11...O1 ⁱ	0.93	2.51	3.236 (2)	135
C9—H9...Cg1 ⁱⁱ	0.93	2.92	3.468 (2)	119

C10—H10...Cg2 ⁱⁱ	0.93	2.90	3.536 (2)	127
C3—H3...Cg3 ⁱⁱⁱ	0.93	2.99	3.673 (2)	132

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