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4-(1*H*-Benzimidazol-2-ylmethyl)-2*H*-1,4-benzothiazin-3(4*H*)-one

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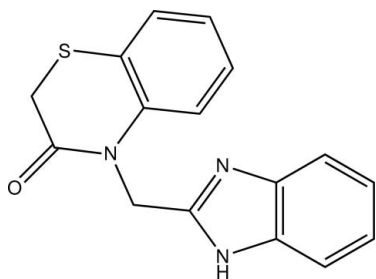
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.041; wR factor = 0.119; data-to-parameter ratio = 21.0.

In the title compound, $\text{C}_{16}\text{H}_{13}\text{N}_3\text{OS}$, the thiomorpholine ring exists in a screw boat conformation. The angle between the benzimidazole ring system and the benzene ring fused to the thiazine ring is $67.22(6)^\circ$. In the crystal, molecules form infinite chains along the a axis *via* intermolecular $\text{N}-\text{H}\cdots\text{N}$ interactions. $\text{C}-\text{H}\cdots\pi$ interactions also contribute to the stability of the crystal structure.

Related literature

For the biological activity of molecules containing 1*H*-benzimidazole, see: Sridhar & Ramesh (2001); Guven *et al.* (2007); Nofal *et al.* (2002); Pedini *et al.* (1994). For a related structure, see: Fun *et al.* (2009). For ring puckering parameters, see: Cremer & Pople (1975). For the stability of the temperature controller used for the data collection, see: Cosier & Glazer (1986).



Experimental

Crystal data

 $\text{C}_{16}\text{H}_{13}\text{N}_3\text{OS}$
 $M_r = 295.35$

 Orthorhombic, $Pbca$
 $a = 9.4498(8)$ Å
 $b = 17.0223(16)$ Å
 $c = 17.4454(16)$ Å
 $V = 2806.2(4)$ Å³
 $Z = 8$

 Mo $K\alpha$ radiation
 $\mu = 0.23$ mm⁻¹
 $T = 100$ K

 $0.50 \times 0.20 \times 0.13$ mm

Data collection

 Bruker APEXII DUO CCD area-detector diffractometer
 Absorption correction: multi-scan (*SADABS*; Bruker, 2009)
 $T_{\min} = 0.893$, $T_{\max} = 0.969$

 16727 measured reflections
 4075 independent reflections
 3185 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.040$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.119$
 $S = 1.03$
 4075 reflections
 194 parameters

H atoms treated by a mixture of independent and constrained refinement

 $\Delta\rho_{\text{max}} = 0.44$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.26$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

 Cg1 and Cg2 are the centroids of the $\text{C1}-\text{C6}$ and $\text{C11}-\text{C16}$ rings, respectively.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1N1}\cdots\text{N2}^i$	0.859 (19)	1.926 (19)	2.7800 (15)	173 (2)
$\text{C12}-\text{H12A}\cdots\text{Cg1}^{\text{ii}}$	0.93	2.97	3.6736 (16)	134
$\text{C3}-\text{H3A}\cdots\text{Cg2}^{\text{iii}}$	0.93	2.61	3.4750 (17)	155

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

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WN2387).

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4-(1*H*-Benzimidazol-2-ylmethyl)-2*H*-1,4-benzothiazin-3(4*H*)-one

Hoong-Kun Fun, Mohd Mustaqim Rosli, Janardhana Gowda, A. M. A. Khader and B. Kalluraya

S1. Comment

A number of molecules containing the 1*H*-benzimidazole nucleus exhibit a broad spectrum of biological activity, including anti-inflammatory (Sridhar *et al.*, 2001), antifungal (Güven *et al.*, 2007), antibacterial (Nofal *et al.*, 2002) and anthelmintic (Pedini *et al.*, 1994) properties. With these results in mind, we have paid particular attention to the preparation of derivatives of 1*H*-benzimidazole and we report here the crystal structure of the title compound, a 1*H*-benzimidazole derivative containing 2*H*-1,4-benzothiazin-3(4*H*)-one.

The bond lengths and angles are within normal ranges. The thiomorpholine ring (C1, C6-C8, N3, S1) adopts a screw boat confirmation with puckering parameters (Cremer & Pople, 1975) being $Q = 0.6563(13)$ Å; $\theta = 66.76(12)^\circ$ and $\varphi = 334.16(14)^\circ$. The angle between the benzimidazole ring system and the benzene ring fused to the thiazine ring is $67.22(6)^\circ$.

The intermolecular interaction $N1-H1N1\cdots N2$ links the molecules to form infinite chains along the *a*-axis. The crystal structure is further stabilized by $C-H\cdots\pi$ interactions involving the C1-C6 (Cg1) and C11-C16 (Cg2) benzene rings (Table 1).

S2. Experimental

A mixture of 2-(3-oxo-2,3-dihydro-4*H*-1,4-benzothiazin-4-yl)acetic acid (3.3 mmol) (Fun *et al.*, 2009) and *o*-phenylenediamine (2.2 mmol) was heated at 140 °C under solvent-free conditions for 3 h and completion of the reaction was checked by TLC. The reaction mixture was cooled to room temperature and the solid product was washed with a saturated solution of sodium bicarbonate to yield 4-(1*H*-benzimidazol-2-ylmethyl)-2*H*-1,4-benzothiazin-3(4*H*)-one as a red solid. Single crystals suitable for X-ray analysis were obtained by crystallization from absolute ethanol under slow evaporation (M.p. 493 K).

S3. Refinement

The H atom attached to N1 was located in a difference map and refined isotropically; $N1-H1N1 = 0.86(2)$ Å. The carbon-bound H atoms were positioned geometrically [$C-H = 0.93$ or 0.97 Å] and were refined using a riding model, with $U_{iso}(H) = 1.2U_{eq}(C)$.

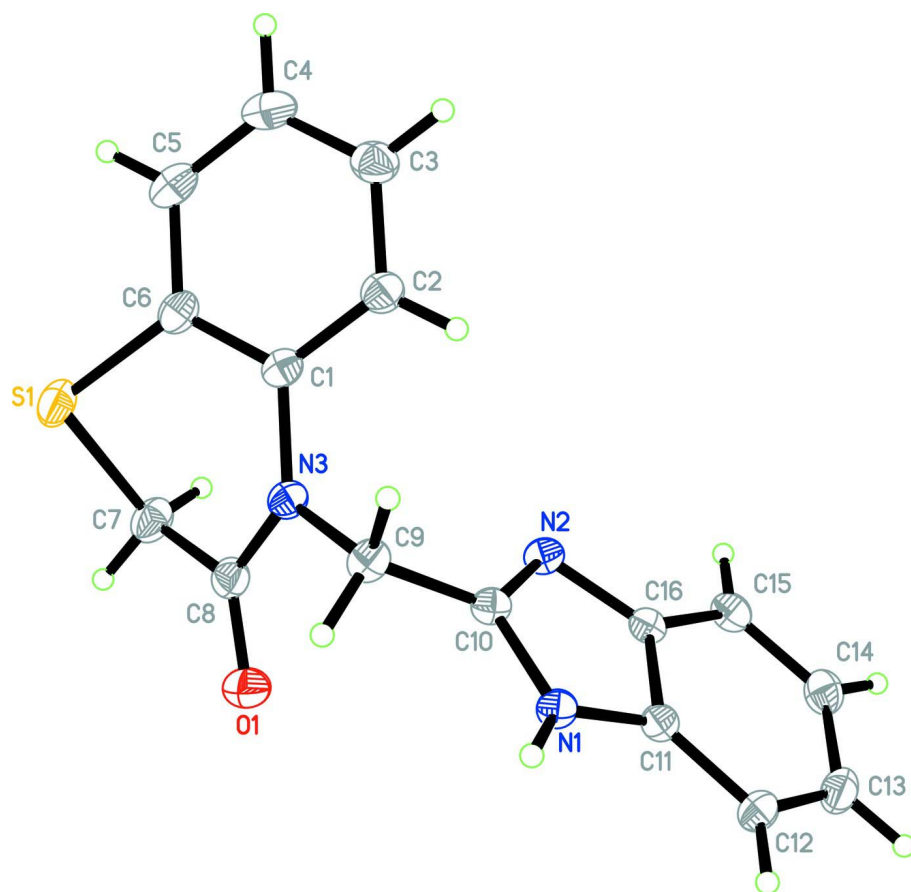


Figure 1

The molecular structure, showing 50% probability displacement ellipsoids and the atom-numbering scheme. Hydrogen atoms are shown as spheres of arbitrary radius.

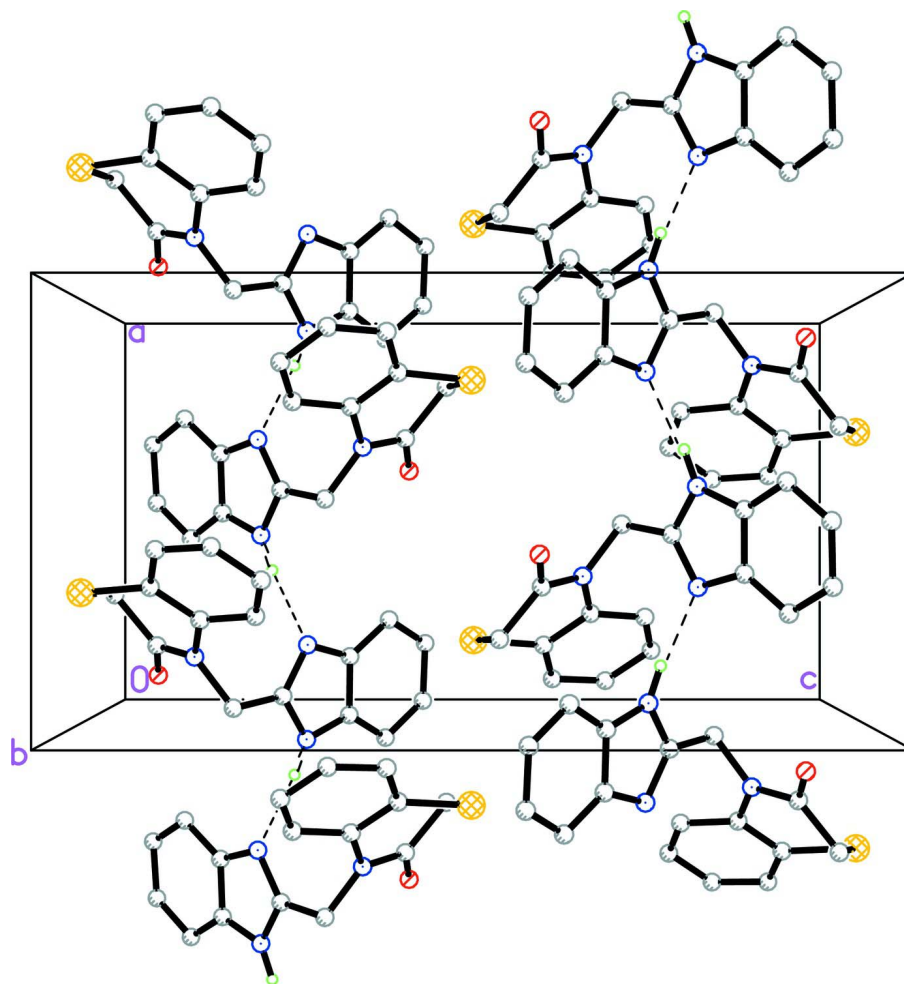


Figure 2

The crystal structure, showing infinite chains along the a-axis. Dashed lines indicate hydrogen bonds. H atoms not involved in the hydrogen bond interactions have been omitted for clarity.

4-(1*H*-Benzimidazol-2-ylmethyl)-2*H*-1,4-benzothiazin- 3(4*H*)-one

Crystal data

$C_{16}H_{13}N_3OS$

$M_r = 295.35$

Orthorhombic, *Pbca*

Hall symbol: -*P* 2ac 2ab

$a = 9.4498$ (8) Å

$b = 17.0223$ (16) Å

$c = 17.4454$ (16) Å

$V = 2806.2$ (4) Å³

$Z = 8$

$F(000) = 1232$

$D_x = 1.398$ Mg m⁻³

Mo *K*α radiation, $\lambda = 0.71073$ Å

Cell parameters from 4703 reflections

$\theta = 2.4$ – 31.6°

$\mu = 0.23$ mm⁻¹

$T = 100$ K

Block, red

$0.50 \times 0.20 \times 0.13$ mm

Data collection

Bruker APEXII DUO CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator
 φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2009)
 $T_{\min} = 0.893$, $T_{\max} = 0.969$
16727 measured reflections
4075 independent reflections
3185 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.040$
 $\theta_{\max} = 30.0^\circ$, $\theta_{\min} = 2.3^\circ$
 $h = -13 \rightarrow 13$
 $k = -23 \rightarrow 23$
 $l = -24 \rightarrow 21$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.119$
 $S = 1.03$
4075 reflections
194 parameters
0 restraints
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
 $w = 1/[\sigma^2(F_o^2) + (0.0605P)^2 + 1.2202P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.44 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.26 \text{ e } \text{\AA}^{-3}$

Special details

Experimental. The crystal was placed in the cold stream of an Oxford Cyrosystems Cobra open-flow nitrogen cryostat (Cosier & Glazer, 1986) operating at 100.0 (1) K.

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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.69098 (4)	0.04490 (2)	0.49857 (2)	0.02567 (12)
O1	0.90053 (12)	-0.12536 (7)	0.41611 (7)	0.0283 (2)
N1	1.05774 (11)	-0.12424 (7)	0.21967 (7)	0.0169 (2)
N2	0.82471 (11)	-0.10176 (7)	0.21708 (7)	0.0166 (2)
N3	0.84628 (12)	-0.01064 (7)	0.35843 (7)	0.0180 (2)
C1	0.75457 (14)	0.05425 (8)	0.34596 (8)	0.0177 (3)
C2	0.74342 (15)	0.08809 (8)	0.27346 (8)	0.0210 (3)
H2A	0.7935	0.0668	0.2325	0.025*
C3	0.65796 (16)	0.15354 (9)	0.26189 (9)	0.0251 (3)
H3A	0.6530	0.1764	0.2135	0.030*
C4	0.58032 (16)	0.18488 (9)	0.32169 (10)	0.0275 (3)
H4A	0.5232	0.2286	0.3137	0.033*
C5	0.58816 (16)	0.15078 (9)	0.39359 (10)	0.0256 (3)
H5A	0.5346	0.1711	0.4337	0.031*
C6	0.67593 (15)	0.08616 (8)	0.40635 (8)	0.0206 (3)
C7	0.70051 (17)	-0.05535 (9)	0.46596 (9)	0.0261 (3)

H7A	0.7100	-0.0900	0.5098	0.031*
H7B	0.6134	-0.0688	0.4396	0.031*
C8	0.82425 (15)	-0.06773 (8)	0.41241 (8)	0.0209 (3)
C9	0.96991 (14)	-0.02207 (8)	0.30913 (8)	0.0191 (3)
H9A	1.0499	-0.0374	0.3406	0.023*
H9B	0.9933	0.0274	0.2847	0.023*
C10	0.94729 (13)	-0.08304 (8)	0.24863 (8)	0.0159 (2)
C11	1.00375 (13)	-0.17492 (8)	0.16502 (8)	0.0165 (3)
C12	1.06703 (15)	-0.23109 (9)	0.11796 (8)	0.0213 (3)
H12A	1.1636	-0.2415	0.1201	0.026*
C13	0.97847 (17)	-0.27045 (9)	0.06787 (9)	0.0245 (3)
H13A	1.0162	-0.3086	0.0356	0.029*
C14	0.83220 (16)	-0.25414 (9)	0.06449 (9)	0.0234 (3)
H14A	0.7765	-0.2809	0.0291	0.028*
C15	0.76917 (15)	-0.19954 (8)	0.11228 (8)	0.0198 (3)
H15A	0.6724	-0.1898	0.1105	0.024*
C16	0.85770 (13)	-0.15965 (8)	0.16350 (8)	0.0158 (2)
H1N1	1.143 (2)	-0.1173 (12)	0.2353 (12)	0.033 (5)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0341 (2)	0.02221 (19)	0.02070 (19)	-0.00054 (14)	0.00590 (14)	-0.00461 (13)
O1	0.0333 (6)	0.0235 (5)	0.0283 (6)	0.0070 (4)	0.0003 (5)	0.0023 (4)
N1	0.0100 (5)	0.0214 (6)	0.0192 (5)	-0.0003 (4)	-0.0003 (4)	-0.0010 (4)
N2	0.0117 (5)	0.0174 (5)	0.0208 (6)	-0.0010 (4)	0.0004 (4)	-0.0003 (4)
N3	0.0166 (5)	0.0180 (5)	0.0194 (6)	0.0007 (4)	0.0020 (4)	-0.0006 (4)
C1	0.0157 (6)	0.0154 (6)	0.0221 (6)	-0.0018 (4)	0.0004 (5)	-0.0017 (5)
C2	0.0220 (6)	0.0180 (6)	0.0230 (7)	-0.0014 (5)	0.0003 (5)	-0.0005 (5)
C3	0.0282 (7)	0.0192 (7)	0.0279 (8)	-0.0014 (5)	-0.0046 (6)	0.0018 (6)
C4	0.0238 (7)	0.0185 (6)	0.0402 (9)	0.0032 (5)	-0.0036 (6)	-0.0013 (6)
C5	0.0220 (7)	0.0214 (7)	0.0334 (8)	0.0014 (5)	0.0045 (6)	-0.0069 (6)
C6	0.0201 (6)	0.0187 (6)	0.0230 (7)	-0.0025 (5)	0.0021 (5)	-0.0039 (5)
C7	0.0349 (8)	0.0197 (7)	0.0236 (7)	-0.0006 (6)	0.0088 (6)	-0.0004 (6)
C8	0.0247 (7)	0.0187 (6)	0.0192 (7)	-0.0005 (5)	-0.0001 (5)	-0.0014 (5)
C9	0.0133 (5)	0.0209 (6)	0.0232 (7)	-0.0019 (5)	0.0004 (5)	-0.0036 (5)
C10	0.0116 (5)	0.0173 (6)	0.0188 (6)	-0.0007 (4)	0.0018 (5)	0.0013 (5)
C11	0.0138 (5)	0.0191 (6)	0.0167 (6)	-0.0002 (4)	0.0006 (5)	0.0011 (5)
C12	0.0182 (6)	0.0245 (7)	0.0211 (7)	0.0030 (5)	0.0035 (5)	-0.0009 (5)
C13	0.0293 (7)	0.0238 (7)	0.0204 (7)	0.0013 (6)	0.0038 (6)	-0.0042 (5)
C14	0.0256 (7)	0.0241 (7)	0.0204 (7)	-0.0036 (5)	-0.0026 (5)	-0.0012 (5)
C15	0.0171 (6)	0.0214 (6)	0.0211 (7)	-0.0029 (5)	-0.0027 (5)	0.0019 (5)
C16	0.0134 (5)	0.0171 (6)	0.0168 (6)	-0.0011 (4)	0.0003 (5)	0.0030 (5)

Geometric parameters (Å, °)

S1—C6	1.7611 (16)	C5—C6	1.396 (2)
S1—C7	1.8011 (16)	C5—H5A	0.9300

O1—C8	1.2191 (18)	C7—C8	1.511 (2)
N1—C10	1.3552 (16)	C7—H7A	0.9700
N1—C11	1.3831 (17)	C7—H7B	0.9700
N1—H1N1	0.86 (2)	C9—C10	1.4955 (19)
N2—C10	1.3214 (16)	C9—H9A	0.9700
N2—C16	1.3936 (17)	C9—H9B	0.9700
N3—C8	1.3692 (19)	C11—C12	1.3950 (19)
N3—C1	1.4207 (17)	C11—C16	1.4046 (18)
N3—C9	1.4637 (17)	C12—C13	1.383 (2)
C1—C2	1.394 (2)	C12—H12A	0.9300
C1—C6	1.3990 (19)	C13—C14	1.411 (2)
C2—C3	1.391 (2)	C13—H13A	0.9300
C2—H2A	0.9300	C14—C15	1.383 (2)
C3—C4	1.382 (2)	C14—H14A	0.9300
C3—H3A	0.9300	C15—C16	1.3998 (19)
C4—C5	1.384 (2)	C15—H15A	0.9300
C4—H4A	0.9300		
C6—S1—C7	95.35 (7)	H7A—C7—H7B	108.0
C10—N1—C11	107.20 (11)	O1—C8—N3	121.18 (13)
C10—N1—H1N1	122.3 (14)	O1—C8—C7	122.47 (14)
C11—N1—H1N1	130.5 (14)	N3—C8—C7	116.35 (12)
C10—N2—C16	104.70 (11)	N3—C9—C10	113.14 (11)
C8—N3—C1	124.36 (12)	N3—C9—H9A	109.0
C8—N3—C9	115.55 (12)	C10—C9—H9A	109.0
C1—N3—C9	120.02 (11)	N3—C9—H9B	109.0
C2—C1—C6	118.86 (13)	C10—C9—H9B	109.0
C2—C1—N3	120.42 (12)	H9A—C9—H9B	107.8
C6—C1—N3	120.71 (13)	N2—C10—N1	113.27 (12)
C3—C2—C1	120.45 (14)	N2—C10—C9	125.91 (12)
C3—C2—H2A	119.8	N1—C10—C9	120.81 (11)
C1—C2—H2A	119.8	N1—C11—C12	132.45 (12)
C4—C3—C2	120.52 (15)	N1—C11—C16	105.08 (11)
C4—C3—H3A	119.7	C12—C11—C16	122.47 (13)
C2—C3—H3A	119.7	C13—C12—C11	116.39 (13)
C3—C4—C5	119.58 (14)	C13—C12—H12A	121.8
C3—C4—H4A	120.2	C11—C12—H12A	121.8
C5—C4—H4A	120.2	C12—C13—C14	121.59 (13)
C4—C5—C6	120.46 (14)	C12—C13—H13A	119.2
C4—C5—H5A	119.8	C14—C13—H13A	119.2
C6—C5—H5A	119.8	C15—C14—C13	121.92 (14)
C5—C6—C1	120.11 (14)	C15—C14—H14A	119.0
C5—C6—S1	120.53 (11)	C13—C14—H14A	119.0
C1—C6—S1	119.35 (11)	C14—C15—C16	116.96 (13)
C8—C7—S1	111.46 (10)	C14—C15—H15A	121.5
C8—C7—H7A	109.3	C16—C15—H15A	121.5
S1—C7—H7A	109.3	N2—C16—C15	129.61 (12)
C8—C7—H7B	109.3	N2—C16—C11	109.75 (11)

S1—C7—H7B	109.3	C15—C16—C11	120.64 (13)
C8—N3—C1—C2	-150.61 (14)	C8—N3—C9—C10	76.17 (15)
C9—N3—C1—C2	25.96 (19)	C1—N3—C9—C10	-100.69 (14)
C8—N3—C1—C6	30.5 (2)	C16—N2—C10—N1	0.06 (15)
C9—N3—C1—C6	-152.88 (13)	C16—N2—C10—C9	178.82 (13)
C6—C1—C2—C3	1.4 (2)	C11—N1—C10—N2	-0.62 (16)
N3—C1—C2—C3	-177.44 (13)	C11—N1—C10—C9	-179.45 (12)
C1—C2—C3—C4	-1.5 (2)	N3—C9—C10—N2	28.7 (2)
C2—C3—C4—C5	0.1 (2)	N3—C9—C10—N1	-152.62 (12)
C3—C4—C5—C6	1.3 (2)	C10—N1—C11—C12	-179.20 (15)
C4—C5—C6—C1	-1.4 (2)	C10—N1—C11—C16	0.88 (14)
C4—C5—C6—S1	177.68 (12)	N1—C11—C12—C13	-178.56 (14)
C2—C1—C6—C5	0.0 (2)	C16—C11—C12—C13	1.3 (2)
N3—C1—C6—C5	178.85 (13)	C11—C12—C13—C14	0.3 (2)
C2—C1—C6—S1	-179.06 (10)	C12—C13—C14—C15	-1.7 (2)
N3—C1—C6—S1	-0.20 (18)	C13—C14—C15—C16	1.3 (2)
C7—S1—C6—C5	142.33 (13)	C10—N2—C16—C15	-178.42 (14)
C7—S1—C6—C1	-38.62 (13)	C10—N2—C16—C11	0.52 (15)
C6—S1—C7—C8	58.75 (12)	C14—C15—C16—N2	179.23 (13)
C1—N3—C8—O1	173.88 (13)	C14—C15—C16—C11	0.4 (2)
C9—N3—C8—O1	-2.8 (2)	N1—C11—C16—N2	-0.88 (15)
C1—N3—C8—C7	-5.6 (2)	C12—C11—C16—N2	179.20 (12)
C9—N3—C8—C7	177.65 (12)	N1—C11—C16—C15	178.18 (12)
S1—C7—C8—O1	137.50 (14)	C12—C11—C16—C15	-1.8 (2)
S1—C7—C8—N3	-42.98 (17)		

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

Cg1 and Cg2 are the centroids of the C1—C6 and C11—C16 rings, respectively.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1M1 \cdots N2 ⁱ	0.859 (19)	1.926 (19)	2.7800 (15)	173 (2)
C12—H12A \cdots Cg1 ⁱⁱ	0.93	2.97	3.6736 (16)	134
C3—H3A \cdots Cg2 ⁱⁱⁱ	0.93	2.61	3.4750 (17)	155

Symmetry codes: (i) $x+1/2, y, -z+1/2$; (ii) $x+5/2, -y-1/2, -z$; (iii) $x+1, -y-1/2, z-1/2$.