

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

N-[4-(4-Bromophenyl)thiazol-2-yl]-4-(piperidin-1-yl)butanamide

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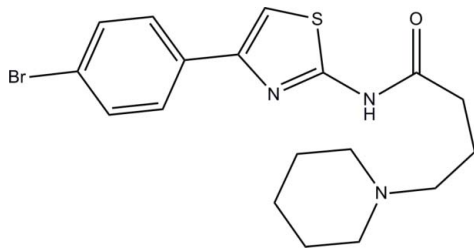
Received 28 April 2012; accepted 29 April 2012

Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.042; wR factor = 0.114; data-to-parameter ratio = 22.7.

In the title compound, $\text{C}_{18}\text{H}_{22}\text{BrN}_3\text{OS}$, the piperidine ring adopts a chair conformation. The mean plane of the thiazole ring forms dihedral angles of 23.97 (10) and 75.82 (10)° with the mean planes of its adjacent benzene and piperidine rings, respectively. An intramolecular $\text{N}-\text{H}\cdots\text{N}$ hydrogen bond generates an $S(7)$ ring motif in the molecule. In the crystal, no significant intermolecular hydrogen bonds are observed, but a weak $\pi-\pi$ interaction with a centroid-centroid distance of 3.8855 (13) Å occurs.

Related literature

For the pharmacological activity of 2-aminothiazole derivatives, see: Lednicer & Mitscher (1977); Vagdevi *et al.* (2006). For ring puckering parameters, see: Cremer & Pople (1975). For further synthetic details, see: El-Subbagh *et al.* (1999). For the stability of the temperature controller used in the data collection, see: Cosier & Glazer (1986).



Experimental

Crystal data

 $\text{C}_{18}\text{H}_{22}\text{BrN}_3\text{OS}$ $M_r = 408.36$

‡ Thomson Reuters ResearcherID: A-3561-2009.

Triclinic, $P\bar{1}$
 $a = 6.8276$ (7) Å
 $b = 9.2782$ (9) Å
 $c = 14.5907$ (14) Å
 $\alpha = 88.812$ (2)°
 $\beta = 86.085$ (3)°
 $\gamma = 75.394$ (2)°

$V = 892.33$ (15) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 2.43$ mm⁻¹
 $T = 100$ K
 $0.37 \times 0.14 \times 0.05$ mm

Data collection

Bruker APEX DUO CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2009)
 $T_{\min} = 0.467$, $T_{\max} = 0.890$

17762 measured reflections
 5019 independent reflections
 4117 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.042$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.114$
 $S = 1.07$
 5019 reflections
 221 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 1.80$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.86$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H1N2}\cdots\text{N3}$	0.93 (3)	1.83 (3)	2.742 (2)	167 (3)

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).

HAG and AAK thank the Deanship of Scientific Research and Research Center, College of Pharmacy, King Saud University. HKF and TSC thank Universiti Sains Malaysia (USM) for the Research University Grant (1001/PFIZIK/811160). TSC also thanks the Malaysian Government and USM for the award of a research fellowship.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6767).

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supplementary materials

Acta Cryst. (2012). E68, o1665 [doi:10.1107/S1600536812019204]

***N*-[4-(4-Bromophenyl)thiazol-2-yl]-4-(piperidin-1-yl)butanamide**

Hazem A. Ghabbour, Adnan A. Kadi, Hussein I. El-Subbagh, Tze Shyang Chia and Hoong-Kun Fun

Comment

2-Aminothiazole derivatives possess a wide range of pharmacological activities. Some of them have been used as anti-infective or anti-trichomonal agents. Those, having an aromatic substituent at C-4 position, exhibit some central nervous system (CNS) activities (Lednicer & Mitscher, 1977) and have been found to be potent biological response modifiers with significant immunosuppressant activity (Vagdevi *et al.*, 2006).

The asymmetric unit of the title compound is shown in Fig. 1. The piperidine ring (N3/C14–C18) adopts a chair conformation with puckering parameters (Cremer & Pople, 1975), $Q = 0.572$ (2) Å, $\theta = 2.6$ (2)° and $\varphi = 354$ (6)°. The atoms N3 and C16 are deviated from the mean plane of C14/C15/C17/C18 by -0.6758 (16) and 0.6567 (18) Å, respectively. The mean plane of central thiazole ring (S1/N1/C7–C9) forms dihedral angles of 23.97 (10) and 75.82 (10)° with the mean planes of adjacent benzene ring (C1–C6) and piperidine ring, respectively. An intramolecular N2—H1N2···N3 hydrogen bond generates an S(7) ring motif in the molecule.

In the crystal, no significant intermolecular hydrogen bondings are observed. The crystal packing is stabilized by Cg3—Cg3 interaction with centroid–centroid distance of 3.8855 (13) Å [symmetry code: 1-*X*,1-*Y*,1-*Z*; Cg3 is the centroid of C1–C6 ring].

Experimental

A mixture of *N*-(4-(4-bromophenyl)thiazol-2-yl)-4-chlorobutanamide (359 mg, 1 mmol) and piperidine (340 mg, 4 mmol) in dry toluene was stirred and heated to reflux. After 12 h, the mixture was cooled down to room temperature and the solvent was removed in vacuum. The residue was purified by chromatotron using CHCl₃:EtOAc (9:1) as eluting system and the title compound was then crystallized from ethanol (El-Subbagh *et al.*, 1999) as colourless plates.

Refinement

The atom H1N2 was located from difference fourier map and refined freely [N2—H1N2 = 0.93 (3) Å]. The remaining H atoms were positioned geometrically [C—H = 0.95 and 0.99 Å] and refined using a riding model with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$. Twelve outliers (4 5 5), (4 4 5), (4 3 6), (-3 - 2 7), (-3 - 3 7), (4 5 4), (4 4 6), (-2 - 2 5), (4 1 5), (-3 - 2 8), (1 1 2) and (3 1 0) were omitted.

Computing details

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

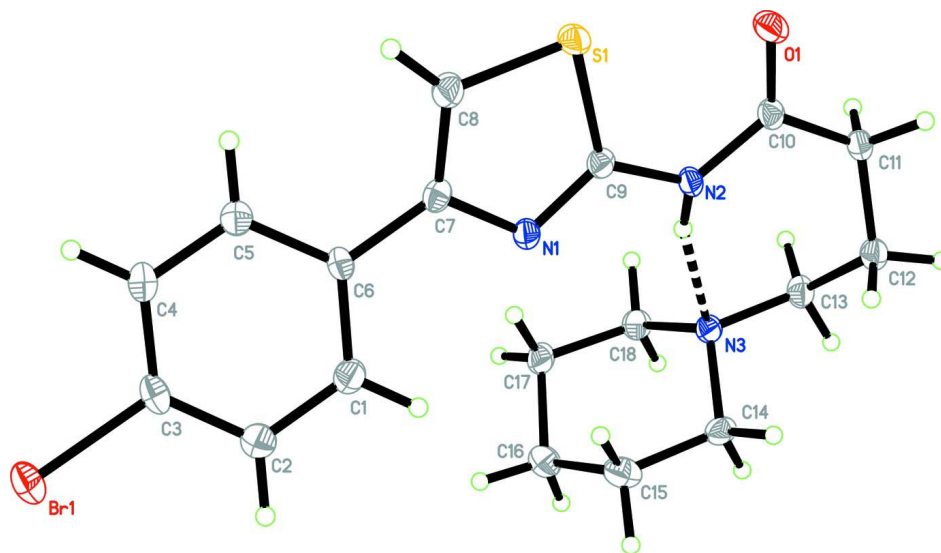


Figure 1

The molecular structure of the title compound with 50% probability displacement ellipsoids. The hydrogen bond is indicated by a dashed line.

N-[4-(4-Bromophenyl)thiazol-2-yl]-4-(piperidin-1-yl)butanamide

Crystal data

$C_{18}H_{22}BrN_3OS$
 $M_r = 408.36$
 Triclinic, $P\bar{1}$
 Hall symbol: $-P\ 1$
 $a = 6.8276$ (7) Å
 $b = 9.2782$ (9) Å
 $c = 14.5907$ (14) Å
 $\alpha = 88.812$ (2)°
 $\beta = 86.085$ (3)°
 $\gamma = 75.394$ (2)°
 $V = 892.33$ (15) Å³

$Z = 2$
 $F(000) = 420$
 $D_x = 1.520$ Mg m⁻³
 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 5955 reflections
 $\theta = 3.1$ – 29.8 °
 $\mu = 2.43$ mm⁻¹
 $T = 100$ K
 Plate, colourless
 $0.37 \times 0.14 \times 0.05$ mm

Data collection

Bruker APEX DUO CCD area-detector
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 φ and ω scans
 Absorption correction: multi-scan
 (*SADABS*; Bruker, 2009)
 $T_{\min} = 0.467$, $T_{\max} = 0.890$

17762 measured reflections
 5019 independent reflections
 4117 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.042$
 $\theta_{\max} = 29.9$ °, $\theta_{\min} = 1.4$ °
 $h = -9 \rightarrow 9$
 $k = -12 \rightarrow 12$
 $l = -20 \rightarrow 20$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.114$
 $S = 1.07$
 5019 reflections

221 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

$$w = 1/[\sigma^2(F_o^2) + (0.0683P)^2 + 0.091P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 1.80 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.86 \text{ e } \text{\AA}^{-3}$

Special details

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

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.82800 (4)	0.81297 (2)	0.549438 (15)	0.02561 (9)
S1	0.28942 (9)	0.30257 (5)	0.19656 (4)	0.01885 (13)
O1	-0.0331 (3)	0.31993 (16)	0.09887 (11)	0.0220 (4)
N1	0.2084 (3)	0.56893 (18)	0.26349 (12)	0.0153 (3)
N2	-0.0331 (3)	0.54008 (18)	0.16321 (12)	0.0154 (4)
N3	-0.2166 (3)	0.83694 (18)	0.14310 (12)	0.0154 (4)
C1	0.3846 (4)	0.6922 (2)	0.40953 (15)	0.0197 (4)
H1A	0.2429	0.7288	0.4038	0.024*
C2	0.4825 (4)	0.7643 (2)	0.46664 (15)	0.0221 (5)
H2A	0.4083	0.8490	0.5007	0.027*
C3	0.6911 (4)	0.7117 (2)	0.47378 (15)	0.0187 (4)
C4	0.8022 (4)	0.5877 (2)	0.42470 (14)	0.0191 (4)
H4A	0.9445	0.5532	0.4294	0.023*
C5	0.7013 (4)	0.5153 (2)	0.36861 (14)	0.0180 (4)
H5A	0.7760	0.4298	0.3354	0.022*
C6	0.4927 (3)	0.5654 (2)	0.35987 (13)	0.0142 (4)
C7	0.3892 (3)	0.4898 (2)	0.29872 (14)	0.0155 (4)
C8	0.4539 (4)	0.3451 (2)	0.26968 (15)	0.0197 (4)
H8A	0.5741	0.2770	0.2876	0.024*
C9	0.1412 (3)	0.4853 (2)	0.20874 (14)	0.0146 (4)
C10	-0.1143 (4)	0.4531 (2)	0.11071 (13)	0.0154 (4)
C11	-0.3088 (4)	0.5259 (2)	0.06591 (15)	0.0177 (4)
H11A	-0.3945	0.4541	0.0698	0.021*
H11B	-0.2725	0.5386	-0.0001	0.021*
C12	-0.4417 (4)	0.6759 (2)	0.10078 (15)	0.0189 (4)
H12A	-0.5764	0.6917	0.0754	0.023*
H12B	-0.4621	0.6698	0.1684	0.023*
C13	-0.3601 (4)	0.8121 (2)	0.07711 (15)	0.0190 (4)
H13A	-0.2907	0.7993	0.0149	0.023*
H13B	-0.4754	0.9015	0.0758	0.023*

C14	-0.3271 (4)	0.9117 (2)	0.22672 (15)	0.0189 (4)
H14A	-0.4143	0.8503	0.2553	0.023*
H14B	-0.4158	1.0092	0.2098	0.023*
C15	-0.1798 (4)	0.9352 (2)	0.29574 (15)	0.0231 (5)
H15A	-0.1006	0.8373	0.3172	0.028*
H15B	-0.2573	0.9897	0.3497	0.028*
C16	-0.0344 (4)	1.0234 (2)	0.25308 (16)	0.0231 (5)
H16A	-0.1106	1.1268	0.2402	0.028*
H16B	0.0691	1.0272	0.2967	0.028*
C17	0.0692 (4)	0.9490 (2)	0.16389 (16)	0.0211 (5)
H17A	0.1518	1.0120	0.1332	0.025*
H17B	0.1613	0.8515	0.1782	0.025*
C18	-0.0854 (4)	0.9263 (2)	0.09965 (15)	0.0195 (4)
H18A	-0.1701	1.0244	0.0814	0.023*
H18B	-0.0140	0.8754	0.0434	0.023*
H1N2	-0.083 (5)	0.643 (3)	0.163 (2)	0.031 (8)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.02680 (16)	0.02875 (14)	0.02536 (13)	-0.01282 (10)	-0.00715 (10)	-0.00372 (9)
S1	0.0190 (3)	0.0121 (2)	0.0252 (3)	-0.0019 (2)	-0.0067 (2)	-0.00195 (18)
O1	0.0235 (9)	0.0153 (7)	0.0278 (8)	-0.0044 (7)	-0.0060 (7)	-0.0037 (6)
N1	0.0127 (9)	0.0143 (7)	0.0193 (8)	-0.0041 (7)	-0.0024 (7)	-0.0009 (6)
N2	0.0123 (9)	0.0126 (7)	0.0220 (8)	-0.0037 (7)	-0.0043 (7)	-0.0018 (6)
N3	0.0147 (9)	0.0136 (7)	0.0187 (8)	-0.0050 (7)	-0.0002 (7)	-0.0003 (6)
C1	0.0156 (11)	0.0191 (9)	0.0237 (10)	-0.0031 (9)	-0.0020 (8)	-0.0008 (8)
C2	0.0205 (12)	0.0209 (9)	0.0248 (10)	-0.0050 (9)	-0.0012 (9)	-0.0036 (8)
C3	0.0215 (12)	0.0210 (9)	0.0175 (9)	-0.0117 (9)	-0.0053 (8)	0.0019 (7)
C4	0.0167 (11)	0.0223 (9)	0.0199 (9)	-0.0070 (9)	-0.0053 (8)	0.0029 (8)
C5	0.0172 (11)	0.0184 (9)	0.0187 (9)	-0.0049 (8)	-0.0020 (8)	0.0003 (7)
C6	0.0145 (10)	0.0143 (8)	0.0156 (8)	-0.0062 (8)	-0.0038 (8)	0.0018 (7)
C7	0.0125 (10)	0.0161 (8)	0.0181 (9)	-0.0043 (8)	-0.0004 (8)	0.0022 (7)
C8	0.0170 (12)	0.0169 (9)	0.0247 (10)	-0.0022 (8)	-0.0061 (9)	0.0004 (7)
C9	0.0133 (10)	0.0131 (8)	0.0178 (8)	-0.0044 (8)	-0.0001 (8)	0.0008 (6)
C10	0.0151 (11)	0.0169 (8)	0.0160 (8)	-0.0076 (8)	0.0003 (8)	0.0000 (7)
C11	0.0147 (11)	0.0174 (9)	0.0225 (9)	-0.0064 (8)	-0.0031 (8)	-0.0014 (7)
C12	0.0158 (11)	0.0190 (9)	0.0222 (10)	-0.0043 (8)	-0.0036 (8)	-0.0011 (7)
C13	0.0168 (11)	0.0171 (9)	0.0240 (10)	-0.0050 (8)	-0.0053 (9)	0.0017 (7)
C14	0.0178 (11)	0.0163 (9)	0.0226 (10)	-0.0061 (8)	0.0050 (9)	-0.0026 (7)
C15	0.0300 (14)	0.0195 (9)	0.0209 (10)	-0.0090 (10)	0.0031 (9)	-0.0032 (8)
C16	0.0243 (13)	0.0198 (9)	0.0269 (11)	-0.0087 (9)	-0.0015 (9)	-0.0039 (8)
C17	0.0156 (11)	0.0171 (9)	0.0322 (11)	-0.0075 (8)	0.0013 (9)	-0.0029 (8)
C18	0.0193 (12)	0.0190 (9)	0.0218 (9)	-0.0088 (9)	0.0022 (8)	-0.0003 (7)

Geometric parameters (\AA , $^\circ$)

Br1—C3	1.898 (2)	C8—H8A	0.9500
S1—C8	1.720 (2)	C10—C11	1.515 (3)
S1—C9	1.7473 (19)	C11—C12	1.533 (3)

O1—C10	1.231 (2)	C11—H11A	0.9900
N1—C9	1.306 (3)	C11—H11B	0.9900
N1—C7	1.393 (3)	C12—C13	1.529 (3)
N2—C10	1.362 (3)	C12—H12A	0.9900
N2—C9	1.380 (3)	C12—H12B	0.9900
N2—H1N2	0.93 (3)	C13—H13A	0.9900
N3—C18	1.472 (3)	C13—H13B	0.9900
N3—C14	1.478 (3)	C14—C15	1.527 (4)
N3—C13	1.480 (3)	C14—H14A	0.9900
C1—C2	1.383 (3)	C14—H14B	0.9900
C1—C6	1.407 (3)	C15—C16	1.532 (4)
C1—H1A	0.9500	C15—H15A	0.9900
C2—C3	1.394 (4)	C15—H15B	0.9900
C2—H2A	0.9500	C16—C17	1.528 (3)
C3—C4	1.390 (3)	C16—H16A	0.9900
C4—C5	1.389 (3)	C16—H16B	0.9900
C4—H4A	0.9500	C17—C18	1.514 (3)
C5—C6	1.396 (3)	C17—H17A	0.9900
C5—H5A	0.9500	C17—H17B	0.9900
C6—C7	1.466 (3)	C18—H18A	0.9900
C7—C8	1.369 (3)	C18—H18B	0.9900
C8—S1—C9	88.45 (10)	H11A—C11—H11B	106.9
C9—N1—C7	110.70 (17)	C13—C12—C11	116.0 (2)
C10—N2—C9	122.88 (17)	C13—C12—H12A	108.3
C10—N2—H1N2	121 (2)	C11—C12—H12A	108.3
C9—N2—H1N2	116 (2)	C13—C12—H12B	108.3
C18—N3—C14	110.36 (17)	C11—C12—H12B	108.3
C18—N3—C13	109.99 (17)	H12A—C12—H12B	107.4
C14—N3—C13	110.74 (18)	N3—C13—C12	113.05 (17)
C2—C1—C6	120.7 (2)	N3—C13—H13A	109.0
C2—C1—H1A	119.7	C12—C13—H13A	109.0
C6—C1—H1A	119.7	N3—C13—H13B	109.0
C1—C2—C3	119.5 (2)	C12—C13—H13B	109.0
C1—C2—H2A	120.3	H13A—C13—H13B	107.8
C3—C2—H2A	120.3	N3—C14—C15	110.96 (19)
C4—C3—C2	121.1 (2)	N3—C14—H14A	109.4
C4—C3—Br1	119.07 (19)	C15—C14—H14A	109.4
C2—C3—Br1	119.76 (17)	N3—C14—H14B	109.4
C5—C4—C3	118.7 (2)	C15—C14—H14B	109.4
C5—C4—H4A	120.6	H14A—C14—H14B	108.0
C3—C4—H4A	120.6	C14—C15—C16	111.38 (19)
C4—C5—C6	121.5 (2)	C14—C15—H15A	109.4
C4—C5—H5A	119.2	C16—C15—H15A	109.4
C6—C5—H5A	119.2	C14—C15—H15B	109.4
C5—C6—C1	118.5 (2)	C16—C15—H15B	109.4
C5—C6—C7	120.87 (18)	H15A—C15—H15B	108.0
C1—C6—C7	120.6 (2)	C17—C16—C15	109.71 (19)
C8—C7—N1	114.3 (2)	C17—C16—H16A	109.7

C8—C7—C6	126.45 (19)	C15—C16—H16A	109.7
N1—C7—C6	119.23 (17)	C17—C16—H16B	109.7
C7—C8—S1	111.31 (17)	C15—C16—H16B	109.7
C7—C8—H8A	124.3	H16A—C16—H16B	108.2
S1—C8—H8A	124.3	C18—C17—C16	111.1 (2)
N1—C9—N2	121.60 (17)	C18—C17—H17A	109.4
N1—C9—S1	115.24 (16)	C16—C17—H17A	109.4
N2—C9—S1	123.16 (15)	C18—C17—H17B	109.4
O1—C10—N2	121.7 (2)	C16—C17—H17B	109.4
O1—C10—C11	120.4 (2)	H17A—C17—H17B	108.0
N2—C10—C11	117.80 (17)	N3—C18—C17	111.59 (18)
C10—C11—C12	120.40 (18)	N3—C18—H18A	109.3
C10—C11—H11A	107.2	C17—C18—H18A	109.3
C12—C11—H11A	107.2	N3—C18—H18B	109.3
C10—C11—H11B	107.2	C17—C18—H18B	109.3
C12—C11—H11B	107.2	H18A—C18—H18B	108.0
C6—C1—C2—C3	1.0 (3)	C10—N2—C9—N1	-175.4 (2)
C1—C2—C3—C4	-0.2 (3)	C10—N2—C9—S1	5.7 (3)
C1—C2—C3—Br1	178.34 (17)	C8—S1—C9—N1	-0.78 (18)
C2—C3—C4—C5	-0.7 (3)	C8—S1—C9—N2	178.24 (19)
Br1—C3—C4—C5	-179.21 (16)	C9—N2—C10—O1	-1.9 (3)
C3—C4—C5—C6	0.8 (3)	C9—N2—C10—C11	178.70 (19)
C4—C5—C6—C1	0.0 (3)	O1—C10—C11—C12	162.6 (2)
C4—C5—C6—C7	178.56 (19)	N2—C10—C11—C12	-18.0 (3)
C2—C1—C6—C5	-0.9 (3)	C10—C11—C12—C13	73.2 (3)
C2—C1—C6—C7	-179.45 (19)	C18—N3—C13—C12	157.54 (18)
C9—N1—C7—C8	-0.6 (3)	C14—N3—C13—C12	-80.2 (2)
C9—N1—C7—C6	178.28 (19)	C11—C12—C13—N3	-84.7 (2)
C5—C6—C7—C8	23.5 (3)	C18—N3—C14—C15	-58.9 (2)
C1—C6—C7—C8	-158.0 (2)	C13—N3—C14—C15	179.05 (17)
C5—C6—C7—N1	-155.2 (2)	N3—C14—C15—C16	56.3 (2)
C1—C6—C7—N1	23.3 (3)	C14—C15—C16—C17	-52.9 (3)
N1—C7—C8—S1	0.0 (3)	C15—C16—C17—C18	53.2 (2)
C6—C7—C8—S1	-178.77 (17)	C14—N3—C18—C17	59.7 (2)
C9—S1—C8—C7	0.39 (18)	C13—N3—C18—C17	-177.78 (17)
C7—N1—C9—N2	-178.11 (19)	C16—C17—C18—N3	-57.3 (2)
C7—N1—C9—S1	0.9 (2)		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2—H1N2 \cdots N3	0.93 (3)	1.83 (3)	2.742 (2)	167 (3)