

3-{[6-(4-Chlorophenyl)imidazo[2,1-*b*]-[1,3,4]thiadiazol-2-yl]methyl}-1,2-benzoxazole

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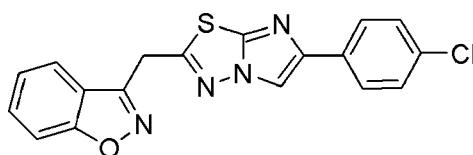
Received 1 January 2011; accepted 7 February 2011

Key indicators: single-crystal X-ray study; $T = 123\text{ K}$; mean $\sigma(\text{C-C}) = 0.005\text{ \AA}$; R factor = 0.053; wR factor = 0.185; data-to-parameter ratio = 15.0.

In the title compound, $\text{C}_{18}\text{H}_{11}\text{ClN}_4\text{OS}$, the benzoxazole and imidazothiadiazole rings are inclined at an angle of $23.81(7)^\circ$ with respect to each other. The imidazothiadiazole and chlorophenyl rings make a dihedral angle of $27.34(3)^\circ$. In the crystal, intermolecular C–H···N interactions generate a chain along the c axis and C–H···O interactions form centrosymmetric dimers resulting in an $R_2^2(26)$ graph-set motif. Moreover, the C–H···N and S···N [3.206(4) Å] interactions links the molecules into $R(7)$ ring motifs. The packing is further stabilized by π – π stacking interactions between the thiadiazole rings with a shortest centroid–centroid distance of 3.497(3) Å. In addition, C–H··· π interactions are observed in the crystal structure

Related literature

For the preparation of the title compound see: Lamani *et al.* (2009). For the biological activity of benzoxazole derivatives, see: Priya *et al.* (2005). For the use of five-membered heterocyclic ring 1,3,4-thiadiazoles in the design of compounds, see: Katritzky (1984); Diamond & Sevrain (2003a,b); Nakao *et al.* (2002a,b). For related structures, see: Sun & Zhang (2009). For hydrogen-bond motifs, see: Bernstein *et al.* (1995)



Experimental

Crystal data

$\text{C}_{18}\text{H}_{11}\text{ClN}_4\text{OS}$
 $M_r = 366.83$

Monoclinic, $C2/c$
 $a = 38.419(7)\text{ \AA}$

$b = 5.7761(10)\text{ \AA}$
 $c = 14.772(3)\text{ \AA}$
 $\beta = 108.004(5)^\circ$
 $V = 3117.5(10)\text{ \AA}^3$
 $Z = 8$

Mo $K\alpha$ radiation
 $\mu = 0.39\text{ mm}^{-1}$
 $T = 123\text{ K}$
 $0.18 \times 0.16 \times 0.16\text{ mm}$

Data collection

Bruker SMART APEX CCD
detector diffractometer
Absorption correction: multi-scan
Bruker Smart Apex
 $T_{\min} = 0.933$, $T_{\max} = 0.940$

8822 measured reflections
3379 independent reflections
2587 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.051$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.185$
 $S = 1.31$
3379 reflections

226 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.74\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.58\text{ e \AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

$Cg1$ and $Cg2$ are the centroids of the C7–C12 and C13–C18 rings, respectively.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C9–H9···O1 ⁱ	0.93	2.39	3.232 (4)	150
C2–H2B···N1 ⁱⁱ	0.97	2.49	3.352 (4)	148
C5–H5···N1 ⁱⁱⁱ	0.93	2.60	3.478 (4)	157
C17–H17···Cg1 ^{iv}	0.93	2.78	3.470 (4)	131
C11–H11···Cg2 ^{iv}	0.93	2.93	3.548 (4)	126

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

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT-Plus* (Bruker, 1998); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *CAMERON* (Watkin *et al.*, 1996); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

NSB is thankful to the University Grants Commission (UGC), India, for financial assistance and the Department of Science and Technology, (DST), India, for the data-collection facility under the IRHPA–DST program.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GW2098).

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supporting information

Acta Cryst. (2011). E67, o617–o618 [doi:10.1107/S1600536811004582]

3-{{6-(4-Chlorophenyl)imidazo[2,1-*b*][1,3,4]thiadiazol-2-yl)methyl}-1,2-benzoxazole

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S1. Comment

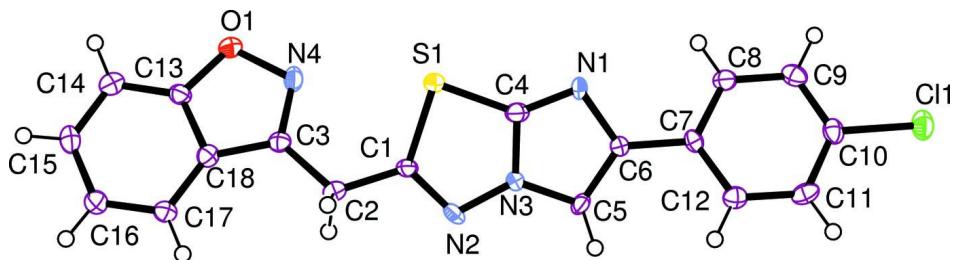
Imidazo[2,1-*b*][1,3,4]thiadiazole derivatives are reported to possess diverse pharmacological properties such as anticancer, antitubercular, antibacterial, antifungal, anticonvulsant, analgesic and antisecretory activities. Moreover, they are known to possess important biological activities (Priya *et al.*, 2005) and are useful in different therapies. Amongst them, five membered heterocyclic ring 1,3,4-thiadiazoles find wide application in designing compounds possessing useful properties (Katritzky *et al.*, 1984; Diamond & Sevrain, 2003*a,b*; Nakao *et al.*, 2002*a,b*). Due to the increasing importance of these heterocycles in biological and pharmaceutical fields, new chemical entities were synthesized by incorporating active pharmacophores in a single molecular frame work so as to enhance their biological activities. In the title compound, the benzisoxazole (O1/N4/C3/C13–18) and imidazothiadiazole (S1/N1—N3/C1/C4—C6) rings are individually planar similar to those reported earlier (Sun & Zhang, 2009) with maximum deviations of 0.038 (3) Å for C1 and 0.016 (3) Å for C3 respectively. The mean planes of the benzisoxazole and imidazothiadiazole are inclined at an angle 23.81 (7)° with each other. The imidazothiadiazole and chlorophenyl rings make a dihedral angle of 27.34 (3)°. The thiadiazole moiety displays differences in the bond lengths between S1—C1/S1—C4 [1.756 (3)/1.736 (3)]. This can be attributed to the resonance effects of the imidazole ring which is stronger than that due to thiadiazole group. The crystal structure is stabilized by intermolecular C—H···N, C—H···O and S···N interactions. The C—H···N interaction generates chain like pattern along c axis. The C—H···O interaction forms centrosymmetric head-to-head dimers about inversion centers corresponding to R22(26) graph set motif (Bernstein *et al.*, 1995). The C—H···N interaction along with S···N [3.206 (4) Å] interaction results in a ring motif with a graph set R(7). The molecular packing is further stabilized by π – π stacking interactions between thiadiazole rings (Cg_3 : centroid of S1/C1/N2/N3/C4) with the shortest centroid–centroid distance 3.497 (3) Å. In addition, π -ring interactions of the type C—H···Cg (Cg being the centroids of rings C7—C12 and C13—C18) are also observed in the crystal structure; details have been given in Table 1.

S2. Experimental

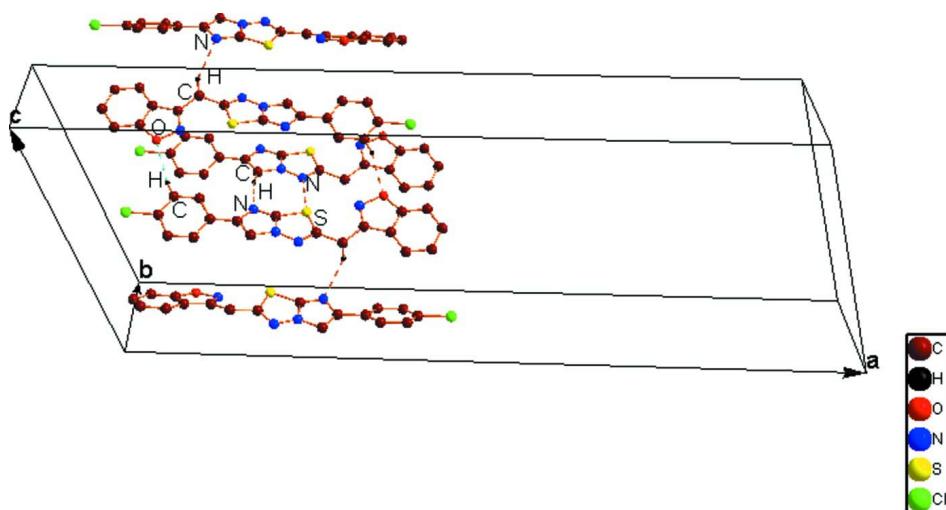
The title compound was synthesized by following the procedure reported earlier (Lamani *et al.*, 2009).

S3. Refinement

The H atoms were placed at calculated positions in the riding model approximation with aromatic C—H = 0.93 Å and methylene C—H = 0.97 Å, and $U_{iso}(\text{H}) = 1.2U_{eq}(\text{N/C})$.

**Figure 1**

ORTEP (Farrugia, 1997) view of the title compound, showing 50% probability ellipsoids and the atom numbering scheme.

**Figure 2**

A unit cell packing of the title compound depicting the C—H···N, C—H···O and S···N intermolecular interactions with dotted lines. H-atoms not involved in hydrogen bonding have been excluded.

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Crystal data



$M_r = 366.83$

Monoclinic, $C2/c$

Hall symbol: -C 2yc

$a = 38.419 (7)$ Å

$b = 5.7761 (10)$ Å

$c = 14.772 (3)$ Å

$\beta = 108.004 (5)^\circ$

$V = 3117.5 (10)$ Å³

$Z = 8$

$F(000) = 1504$

$D_x = 1.563 \text{ Mg m}^{-3}$

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

Cell parameters from 3379 reflections

$\theta = 2.2\text{--}27.0^\circ$

$\mu = 0.39 \text{ mm}^{-1}$

$T = 123$ K

Block, yellow

$0.18 \times 0.16 \times 0.16$ mm

Data collection

Bruker SMART APEX CCD detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan

Bruker Smart Apex

$T_{\min} = 0.933$, $T_{\max} = 0.940$

8822 measured reflections

3379 independent reflections

2587 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.051$
 $\theta_{\text{max}} = 27.0^\circ, \theta_{\text{min}} = 2.2^\circ$

$h = -46 \rightarrow 48$
 $k = -6 \rightarrow 7$
 $l = -18 \rightarrow 14$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.185$
 $S = 1.31$
3379 reflections
226 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-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0894P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.74 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.58 \text{ e } \text{\AA}^{-3}$

Special details

Experimental. The compound was synthesized by following the procedure given in Lamani *et al.*, (2009)

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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 $F^2 > 2\sigma(F^2)$ 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	0.19871 (8)	0.2236 (5)	0.1424 (2)	0.0151 (6)
C2	0.16451 (8)	0.1564 (5)	0.1657 (2)	0.0183 (7)
H2A	0.1565	0.0071	0.1366	0.022*
H2B	0.1711	0.1355	0.2341	0.022*
C3	0.13263 (8)	0.3171 (5)	0.1363 (2)	0.0166 (7)
C4	0.25323 (8)	0.4023 (5)	0.1246 (2)	0.0160 (7)
C5	0.28417 (8)	0.0835 (6)	0.1177 (2)	0.0167 (6)
H5	0.2910	-0.0704	0.1157	0.020*
C6	0.30436 (8)	0.2790 (5)	0.1142 (2)	0.0149 (6)
C7	0.34193 (8)	0.2930 (5)	0.1098 (2)	0.0148 (7)
C8	0.35464 (9)	0.4896 (5)	0.0746 (2)	0.0181 (7)
H8	0.3389	0.6139	0.0527	0.022*
C9	0.39034 (9)	0.5024 (5)	0.0718 (2)	0.0192 (7)
H9	0.3986	0.6340	0.0484	0.023*
C10	0.41322 (8)	0.3176 (6)	0.1039 (2)	0.0183 (7)
C11	0.40183 (9)	0.1192 (6)	0.1396 (2)	0.0206 (7)
H11	0.4178	-0.0038	0.1616	0.025*
C12	0.36610 (8)	0.1082 (6)	0.1418 (2)	0.0178 (7)
H12	0.3580	-0.0244	0.1649	0.021*
C13	0.07777 (9)	0.4766 (5)	0.1117 (2)	0.0176 (7)
C14	0.04157 (9)	0.5159 (6)	0.1076 (2)	0.0209 (7)

H14	0.0292	0.6519	0.0834	0.025*
C15	0.02526 (9)	0.3370 (6)	0.1424 (2)	0.0219 (7)
H15	0.0010	0.3519	0.1407	0.026*
C16	0.04440 (8)	0.1336 (6)	0.1800 (3)	0.0212 (7)
H16	0.0326	0.0188	0.2036	0.025*
C17	0.08036 (8)	0.1001 (6)	0.1828 (2)	0.0178 (7)
H17	0.0928	-0.0352	0.2074	0.021*
C18	0.09718 (8)	0.2764 (5)	0.1474 (2)	0.0166 (7)
O1	0.09977 (6)	0.6281 (4)	0.08271 (17)	0.0222 (5)
N1	0.28445 (7)	0.4813 (4)	0.11762 (19)	0.0161 (6)
N2	0.22129 (7)	0.0628 (5)	0.13533 (19)	0.0171 (6)
N3	0.25177 (7)	0.1673 (4)	0.12454 (19)	0.0158 (6)
N4	0.13496 (7)	0.5207 (5)	0.1006 (2)	0.0208 (6)
S1	0.21240 (2)	0.51078 (13)	0.13431 (6)	0.0182 (2)
Cl1	0.45843 (2)	0.33439 (15)	0.10073 (6)	0.0267 (3)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0141 (14)	0.0126 (15)	0.0175 (16)	0.0005 (12)	0.0032 (13)	0.0015 (12)
C2	0.0170 (15)	0.0154 (16)	0.0248 (18)	0.0016 (12)	0.0095 (14)	0.0034 (13)
C3	0.0197 (15)	0.0141 (15)	0.0155 (16)	-0.0028 (12)	0.0049 (13)	-0.0002 (12)
C4	0.0202 (15)	0.0121 (15)	0.0152 (16)	0.0008 (12)	0.0047 (13)	-0.0006 (12)
C5	0.0152 (14)	0.0138 (15)	0.0214 (16)	0.0032 (12)	0.0062 (13)	0.0002 (13)
C6	0.0131 (14)	0.0162 (16)	0.0153 (16)	-0.0011 (12)	0.0042 (13)	0.0011 (12)
C7	0.0137 (14)	0.0156 (16)	0.0159 (16)	-0.0034 (12)	0.0054 (12)	-0.0023 (12)
C8	0.0206 (16)	0.0157 (16)	0.0191 (17)	0.0021 (12)	0.0078 (14)	0.0006 (12)
C9	0.0252 (17)	0.0161 (17)	0.0176 (17)	-0.0052 (13)	0.0086 (14)	-0.0021 (12)
C10	0.0142 (14)	0.0231 (17)	0.0179 (17)	-0.0048 (12)	0.0051 (13)	-0.0056 (13)
C11	0.0231 (16)	0.0177 (17)	0.0220 (18)	0.0049 (13)	0.0084 (14)	-0.0019 (13)
C12	0.0193 (15)	0.0169 (16)	0.0178 (17)	0.0012 (13)	0.0067 (13)	0.0014 (13)
C13	0.0190 (16)	0.0158 (16)	0.0204 (17)	-0.0004 (12)	0.0096 (14)	0.0009 (13)
C14	0.0194 (16)	0.0200 (17)	0.0220 (18)	0.0035 (13)	0.0042 (14)	0.0000 (13)
C15	0.0177 (16)	0.0283 (19)	0.0209 (18)	-0.0017 (13)	0.0077 (14)	-0.0013 (14)
C16	0.0149 (15)	0.0216 (17)	0.0276 (19)	-0.0029 (13)	0.0075 (14)	0.0010 (14)
C17	0.0172 (15)	0.0163 (16)	0.0202 (17)	0.0005 (12)	0.0060 (13)	0.0023 (13)
C18	0.0136 (14)	0.0178 (16)	0.0183 (16)	0.0011 (12)	0.0049 (13)	-0.0003 (13)
O1	0.0200 (11)	0.0167 (12)	0.0325 (14)	0.0044 (9)	0.0120 (11)	0.0055 (10)
N1	0.0114 (12)	0.0167 (14)	0.0206 (14)	0.0006 (10)	0.0054 (11)	-0.0008 (11)
N2	0.0127 (12)	0.0178 (14)	0.0211 (15)	-0.0055 (10)	0.0059 (11)	0.0001 (11)
N3	0.0174 (13)	0.0115 (13)	0.0189 (15)	-0.0012 (10)	0.0059 (11)	0.0013 (10)
N4	0.0147 (13)	0.0244 (16)	0.0251 (16)	0.0015 (11)	0.0087 (12)	0.0010 (12)
S1	0.0184 (4)	0.0124 (4)	0.0257 (5)	-0.0008 (3)	0.0096 (4)	-0.0007 (3)
Cl1	0.0177 (4)	0.0329 (5)	0.0318 (5)	-0.0021 (3)	0.0110 (4)	-0.0057 (4)

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

C1—N2	1.297 (4)	C9—C10	1.371 (4)
C1—C2	1.508 (4)	C9—H9	0.9300
C1—S1	1.756 (3)	C10—C11	1.387 (5)
C2—C3	1.490 (4)	C10—Cl1	1.755 (3)
C2—H2A	0.9700	C11—C12	1.384 (4)
C2—H2B	0.9700	C11—H11	0.9300
C3—N4	1.304 (4)	C12—H12	0.9300
C3—C18	1.440 (4)	C13—O1	1.374 (4)
C4—N1	1.316 (4)	C13—C14	1.392 (4)
C4—N3	1.358 (4)	C13—C18	1.389 (4)
C4—S1	1.736 (3)	C14—C15	1.387 (5)
C5—C6	1.380 (4)	C14—H14	0.9300
C5—N3	1.369 (4)	C15—C16	1.406 (5)
C5—H5	0.9300	C15—H15	0.9300
C6—N1	1.406 (4)	C16—C17	1.383 (4)
C6—C7	1.468 (4)	C16—H16	0.9300
C7—C12	1.398 (4)	C17—C18	1.392 (4)
C7—C8	1.398 (4)	C17—H17	0.9300
C8—C9	1.387 (4)	O1—N4	1.436 (3)
C8—H8	0.9300	N2—N3	1.370 (3)
N2—C1—C2	119.1 (3)	C12—C11—C10	118.5 (3)
N2—C1—S1	116.7 (2)	C12—C11—H11	120.7
C2—C1—S1	124.0 (2)	C10—C11—H11	120.7
C3—C2—C1	117.9 (3)	C11—C12—C7	121.1 (3)
C3—C2—H2A	107.8	C11—C12—H12	119.4
C1—C2—H2A	107.8	C7—C12—H12	119.4
C3—C2—H2B	107.8	O1—C13—C14	125.8 (3)
C1—C2—H2B	107.8	O1—C13—C18	109.8 (3)
H2A—C2—H2B	107.2	C14—C13—C18	124.4 (3)
N4—C3—C18	112.3 (3)	C13—C14—C15	115.0 (3)
N4—C3—C2	121.8 (3)	C13—C14—H14	122.5
C18—C3—C2	125.9 (3)	C15—C14—H14	122.5
N1—C4—N3	112.7 (3)	C14—C15—C16	121.9 (3)
N1—C4—S1	138.5 (3)	C14—C15—H15	119.1
N3—C4—S1	108.8 (2)	C16—C15—H15	119.1
C6—C5—N3	104.3 (3)	C17—C16—C15	121.6 (3)
C6—C5—H5	127.8	C17—C16—H16	119.2
N3—C5—H5	127.8	C15—C16—H16	119.2
C5—C6—N1	111.2 (3)	C16—C17—C18	117.5 (3)
C5—C6—C7	128.2 (3)	C16—C17—H17	121.3
N1—C6—C7	120.6 (3)	C18—C17—H17	121.3
C12—C7—C8	118.3 (3)	C17—C18—C13	119.7 (3)
C12—C7—C6	120.2 (3)	C17—C18—C3	136.7 (3)
C8—C7—C6	121.5 (3)	C13—C18—C3	103.7 (3)
C9—C8—C7	121.1 (3)	C13—O1—N4	107.6 (2)

C9—C8—H8	119.4	C4—N1—C6	103.5 (2)
C7—C8—H8	119.4	C1—N2—N3	108.1 (3)
C10—C9—C8	118.8 (3)	C4—N3—N2	118.5 (3)
C10—C9—H9	120.6	C4—N3—C5	108.4 (3)
C8—C9—H9	120.6	N2—N3—C5	133.0 (3)
C9—C10—C11	122.1 (3)	C3—N4—O1	106.6 (2)
C9—C10—Cl1	118.8 (2)	C4—S1—C1	87.82 (14)
C11—C10—Cl1	119.1 (2)		
N2—C1—C2—C3	-155.9 (3)	O1—C13—C18—C3	-0.9 (3)
S1—C1—C2—C3	30.1 (4)	C14—C13—C18—C3	179.8 (3)
C1—C2—C3—N4	-7.1 (5)	N4—C3—C18—C17	-177.9 (4)
C1—C2—C3—C18	176.2 (3)	C2—C3—C18—C17	-0.9 (6)
N3—C5—C6—N1	-0.8 (3)	N4—C3—C18—C13	1.6 (4)
N3—C5—C6—C7	177.9 (3)	C2—C3—C18—C13	178.6 (3)
C5—C6—C7—C12	-22.8 (5)	C14—C13—O1—N4	179.2 (3)
N1—C6—C7—C12	155.8 (3)	C18—C13—O1—N4	0.0 (3)
C5—C6—C7—C8	157.7 (3)	N3—C4—N1—C6	-0.8 (3)
N1—C6—C7—C8	-23.7 (4)	S1—C4—N1—C6	178.9 (3)
C12—C7—C8—C9	-0.3 (5)	C5—C6—N1—C4	1.0 (3)
C6—C7—C8—C9	179.2 (3)	C7—C6—N1—C4	-177.8 (3)
C7—C8—C9—C10	0.2 (5)	C2—C1—N2—N3	-173.3 (3)
C8—C9—C10—C11	-0.3 (5)	S1—C1—N2—N3	1.1 (3)
C8—C9—C10—Cl1	-179.8 (2)	N1—C4—N3—N2	177.4 (3)
C9—C10—C11—C12	0.5 (5)	S1—C4—N3—N2	-2.5 (3)
Cl1—C10—C11—C12	180.0 (2)	N1—C4—N3—C5	0.3 (4)
C10—C11—C12—C7	-0.7 (5)	S1—C4—N3—C5	-179.5 (2)
C8—C7—C12—C11	0.6 (5)	C1—N2—N3—C4	0.9 (4)
C6—C7—C12—C11	-179.0 (3)	C1—N2—N3—C5	177.1 (3)
O1—C13—C14—C15	-179.3 (3)	C6—C5—N3—C4	0.3 (3)
C18—C13—C14—C15	-0.1 (5)	C6—C5—N3—N2	-176.1 (3)
C13—C14—C15—C16	0.9 (5)	C18—C3—N4—O1	-1.7 (4)
C14—C15—C16—C17	-1.1 (5)	C2—C3—N4—O1	-178.8 (3)
C15—C16—C17—C18	0.4 (5)	C13—O1—N4—C3	1.1 (3)
C16—C17—C18—C13	0.4 (5)	N1—C4—S1—C1	-177.4 (4)
C16—C17—C18—C3	179.8 (4)	N3—C4—S1—C1	2.4 (2)
O1—C13—C18—C17	178.7 (3)	N2—C1—S1—C4	-2.1 (3)
C14—C13—C18—C17	-0.6 (5)	C2—C1—S1—C4	172.1 (3)

Hydrogen-bond geometry (Å, °)

Cg1 and Cg2 are the centroids of the C7—C12 and C13—C18 rings, respectively.

D—H···A	D—H	H···A	D···A	D—H···A
C9—H9···O1 ⁱ	0.93	2.39	3.232 (4)	150
C2—H2B···N1 ⁱⁱ	0.97	2.49	3.352 (4)	148
C5—H5···N1 ⁱⁱⁱ	0.93	2.60	3.478 (4)	157

C17—H17···Cg1 ^{iv}	0.93	2.78	3.470 (4)	131
C11—H11···Cg2 ^{iv}	0.93	2.93	3.548 (4)	126

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