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5-(4-Bromophenoxy)-1-methyl-3-methyl-1H-pyrazole-4-carbaldehyde-O-[(5-methoxy-1,3,4-thiadiazol-2-yl)-methyl]-oxime

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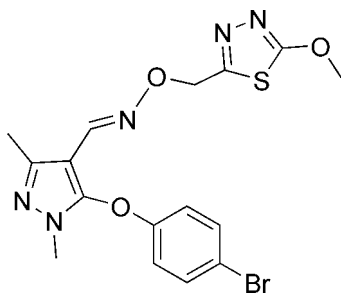
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Key indicators: single-crystal X-ray study; $T = 113$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.029; wR factor = 0.054; data-to-parameter ratio = 13.3.

In the title molecule, $\text{C}_{16}\text{H}_{16}\text{BrN}_5\text{O}_3\text{S}$, the 1,3,4-thiadiazole ring is situated under the benzene ring, forming a dihedral angle of $86.6(2)^\circ$, and with an $\text{S}\cdots\text{C}_g$ (where C_g is the centroid of the benzene ring) distance of $3.312(3)$ Å. The benzene and 1,3,4-thiadiazole rings form dihedral angles of $83.8(3)$ and $57.7(2)^\circ$, respectively, with the central pyrazole ring. In the absence of classical hydrogen bonds, the crystal packing is stabilized by a $\text{C}-\text{H}\cdots\pi$ interaction.

Related literature

For a related structure, see: Dai *et al.* (2011).



Experimental

Crystal data

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

$M_r = 438.31$

Triclinic, $P\bar{1}$
 $a = 9.732(3)$ Å
 $b = 9.832(2)$ Å
 $c = 11.166(3)$ Å
 $\alpha = 64.55(2)^\circ$
 $\beta = 69.62(2)^\circ$
 $\gamma = 75.33(3)^\circ$

$V = 897.5(4)$ Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 2.43$ mm⁻¹
 $T = 113$ K
 $0.20 \times 0.18 \times 0.12$ mm

Data collection

Rigaku Saturn724 CCD diffractometer
 Absorption correction: multi-scan (*CrystalClear*; Rigaku, 2008)
 $T_{\min} = 0.642$, $T_{\max} = 0.759$

7729 measured reflections
 3175 independent reflections
 2363 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.044$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.029$
 $wR(F^2) = 0.054$
 $S = 1.02$
 3175 reflections

238 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.36$ e Å⁻³
 $\Delta\rho_{\min} = -0.49$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

C_g is the centroid of the C1–C6 ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C11}-\text{H11A}\cdots\text{C}_g^i$	0.98	2.89	3.652(4)	125

Symmetry code: (i) $-x + 2, -y + 2, -z + 2$.

Data collection: *CrystalClear* (Rigaku, 2008); 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: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CV5348).

References

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 Rigaku (2008). *CrystalClear*. Rigaku Corporation, Tokyo, Japan.
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.

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5-(4-Bromophenoxy)-1-methyl-3-methyl-1*H*-pyrazole-4-carbaldehyde-*O*-[(5-methoxy-1,3,4-thiadiazol-2-yl)-methyl]oxime

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S1. Comment

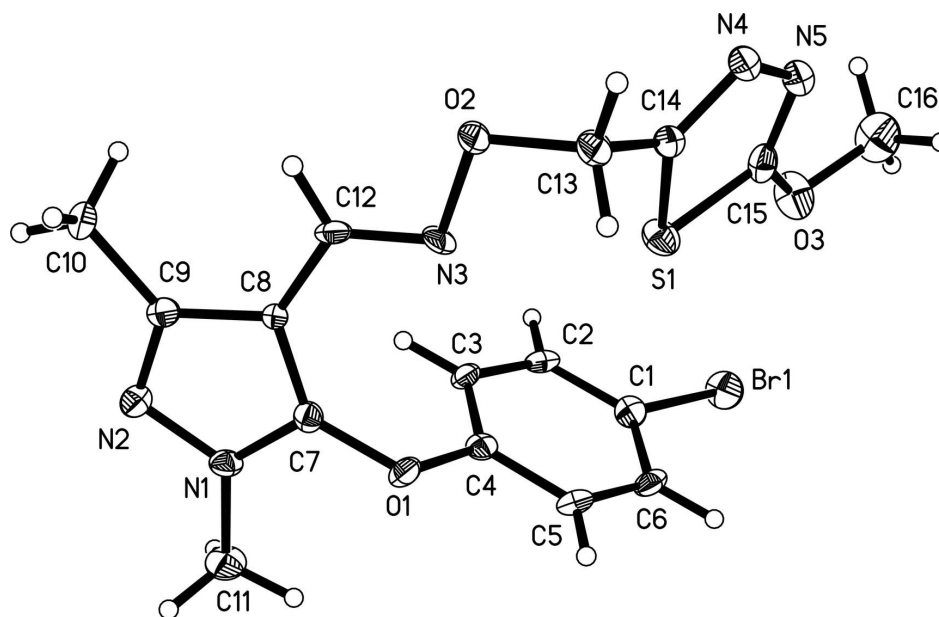
In a continuation of our structural study of pyrazole oxime derivatives (Dai *et al.*, 2011), we report here the crystal structure of the title compound, (I). In (I) (Fig. 1), all bonds lengths and angles are similar to those observed in the related compound (Dai *et al.*, 2011). The dihedral angles between the substituted phenyl ring and the pyrazole ring and between the 1,3,4-thiadiazole ring and the pyrazole ring are 83.8 (3)° and 57.7 (2)°, respectively. The crystal packing displays short intermolecular C...C contacts of 3.203 (4) Å.

S2. Experimental

To a stirred solution of 1-methyl-3-methyl-5-(4-bromophenoxy)-1*H*-pyrazole -4-carbaldehyde oxime (3 mmol), and powdered potassium carbonate (9 mmol) in 30 ml of anhydrous acetonitrile, was added 2-chloromethyl-5-methoxy-1,3,4-thiadiazole (4.2 mmol) at room temperature. The mixture was heated to reflux for 13 h. After removal of the solvent, the residue was separated by column chromatography on silica gel using a mixture of petroleum ether/ethyl acetate to obtain colourless crystals.

S3. Refinement

All H atoms were placed in calculated positions, with C–H = 0.95–0.99 Å, and included in the final cycles of refinement using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

**Figure 1**

The molecular structure of (I) showing the atomic numbering and 50% probability displacement ellipsoids.

5-(4-Bromophenoxy)-1-methyl-3-methyl-1*H*-pyrazole-4-carbaldehyde- *O*-[(5-methoxy-1,3,4-thiadiazol-2-yl)-methyl]oxime

Crystal data

$C_{16}H_{16}BrN_5O_3S$
 $M_r = 438.31$
 Triclinic, $P\bar{1}$
 Hall symbol: $-P\ 1$
 $a = 9.732\ (3)\ \text{\AA}$
 $b = 9.832\ (2)\ \text{\AA}$
 $c = 11.166\ (3)\ \text{\AA}$
 $\alpha = 64.55\ (2)^\circ$
 $\beta = 69.62\ (2)^\circ$
 $\gamma = 75.33\ (3)^\circ$
 $V = 897.5\ (4)\ \text{\AA}^3$

$Z = 2$
 $F(000) = 444$
 $D_x = 1.622\ \text{Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$
 Cell parameters from 3391 reflections
 $\theta = 2.1\text{--}27.9^\circ$
 $\mu = 2.43\ \text{mm}^{-1}$
 $T = 113\ \text{K}$
 Prism, colourless
 $0.20 \times 0.18 \times 0.12\ \text{mm}$

Data collection

Rigaku Saturn724 CCD
 diffractometer
 Radiation source: rotating anode
 Multilayer monochromator
 Detector resolution: $14.22\ \text{pixels mm}^{-1}$
 ω and φ scans
 Absorption correction: multi-scan
 (*CrystalClear*; Rigaku, 2008)
 $T_{\min} = 0.642$, $T_{\max} = 0.759$

7729 measured reflections
 3175 independent reflections
 2363 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.044$
 $\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.1^\circ$
 $h = -11 \rightarrow 11$
 $k = -11 \rightarrow 11$
 $l = -13 \rightarrow 13$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.029$
 $wR(F^2) = 0.054$
 $S = 1.02$
 3175 reflections
 238 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
 $w = 1/[\sigma^2(F_o^2) + (0.P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.36 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.49 \text{ e } \text{\AA}^{-3}$

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	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.51369 (3)	0.66558 (3)	1.32176 (3)	0.02651 (10)
S1	0.54286 (8)	0.73735 (8)	0.87231 (7)	0.02150 (19)
O1	0.96309 (18)	0.88211 (18)	0.76885 (17)	0.0166 (5)
O2	0.63830 (19)	0.93453 (19)	0.53541 (18)	0.0202 (5)
O3	0.3297 (2)	0.6561 (2)	1.10298 (18)	0.0291 (5)
N1	1.0439 (2)	1.1205 (2)	0.6791 (2)	0.0157 (5)
N2	1.0308 (2)	1.2623 (2)	0.5802 (2)	0.0173 (6)
N3	0.7358 (2)	0.9392 (2)	0.6035 (2)	0.0166 (5)
N4	0.3648 (2)	0.7120 (2)	0.7625 (2)	0.0196 (6)
N5	0.2883 (2)	0.6764 (2)	0.9015 (2)	0.0202 (6)
C1	0.6530 (3)	0.7382 (3)	1.1473 (3)	0.0174 (7)
C2	0.6437 (3)	0.8902 (3)	1.0675 (3)	0.0161 (7)
H2	0.5676	0.9587	1.0999	0.019*
C3	0.7454 (3)	0.9447 (3)	0.9390 (3)	0.0155 (7)
H3	0.7395	1.0497	0.8827	0.019*
C4	0.8547 (3)	0.8423 (3)	0.8958 (3)	0.0143 (6)
C5	0.8662 (3)	0.6893 (3)	0.9768 (3)	0.0168 (7)
H5	0.9440	0.6213	0.9455	0.020*
C6	0.7646 (3)	0.6357 (3)	1.1029 (3)	0.0176 (7)
H6	0.7705	0.5305	1.1587	0.021*
C7	0.9571 (3)	1.0319 (3)	0.6821 (3)	0.0144 (7)
C8	0.8804 (3)	1.1148 (3)	0.5847 (3)	0.0114 (6)
C9	0.9324 (3)	1.2589 (3)	0.5235 (3)	0.0150 (7)
C10	0.8892 (3)	1.3946 (3)	0.4084 (2)	0.0211 (7)
H10A	0.9228	1.4843	0.4016	0.032*

H10B	0.7816	1.4105	0.4264	0.032*
H10C	0.9350	1.3784	0.3216	0.032*
C11	1.1437 (3)	1.0799 (3)	0.7631 (3)	0.0239 (7)
H11A	1.1112	1.1413	0.8199	0.036*
H11B	1.2437	1.0986	0.7032	0.036*
H11C	1.1436	0.9721	0.8229	0.036*
C12	0.7799 (3)	1.0706 (3)	0.5427 (3)	0.0153 (6)
H12	0.7448	1.1421	0.4664	0.018*
C13	0.5995 (3)	0.7852 (3)	0.5917 (3)	0.0194 (7)
H13A	0.6904	0.7119	0.5950	0.023*
H13B	0.5533	0.7753	0.5301	0.023*
C14	0.4958 (3)	0.7459 (3)	0.7328 (3)	0.0151 (6)
C15	0.3691 (3)	0.6834 (3)	0.9682 (3)	0.0194 (7)
C16	0.1869 (3)	0.6019 (3)	1.1781 (3)	0.0360 (9)
H16A	0.1105	0.6772	1.1391	0.054*
H16B	0.1657	0.5860	1.2754	0.054*
H16C	0.1879	0.5058	1.1708	0.054*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.02695 (19)	0.03281 (19)	0.01571 (17)	-0.00625 (14)	-0.00270 (14)	-0.00650 (15)
S1	0.0216 (4)	0.0264 (4)	0.0212 (4)	-0.0059 (4)	-0.0073 (3)	-0.0105 (4)
O1	0.0170 (11)	0.0132 (10)	0.0140 (11)	0.0018 (8)	-0.0045 (8)	-0.0019 (9)
O2	0.0230 (12)	0.0190 (11)	0.0215 (11)	-0.0089 (9)	-0.0136 (9)	-0.0009 (10)
O3	0.0335 (13)	0.0366 (13)	0.0201 (12)	-0.0128 (11)	-0.0003 (10)	-0.0140 (11)
N1	0.0149 (14)	0.0174 (13)	0.0178 (13)	-0.0006 (11)	-0.0085 (11)	-0.0070 (11)
N2	0.0196 (14)	0.0154 (13)	0.0160 (13)	-0.0016 (11)	-0.0063 (11)	-0.0044 (11)
N3	0.0142 (13)	0.0213 (13)	0.0184 (13)	-0.0059 (11)	-0.0091 (11)	-0.0056 (12)
N4	0.0185 (14)	0.0196 (13)	0.0202 (14)	-0.0064 (11)	-0.0067 (11)	-0.0036 (12)
N5	0.0190 (15)	0.0195 (14)	0.0211 (14)	-0.0057 (11)	-0.0055 (11)	-0.0050 (12)
C1	0.0212 (17)	0.0228 (16)	0.0088 (15)	-0.0060 (14)	-0.0058 (13)	-0.0034 (14)
C2	0.0157 (16)	0.0167 (15)	0.0195 (17)	0.0050 (13)	-0.0105 (13)	-0.0095 (14)
C3	0.0175 (16)	0.0106 (14)	0.0180 (16)	0.0011 (13)	-0.0096 (13)	-0.0030 (13)
C4	0.0155 (16)	0.0176 (15)	0.0121 (15)	-0.0018 (13)	-0.0075 (12)	-0.0049 (13)
C5	0.0168 (16)	0.0161 (15)	0.0170 (16)	0.0052 (13)	-0.0087 (13)	-0.0066 (13)
C6	0.0220 (17)	0.0134 (15)	0.0184 (16)	0.0020 (13)	-0.0130 (13)	-0.0034 (14)
C7	0.0130 (16)	0.0143 (15)	0.0139 (15)	-0.0015 (13)	-0.0012 (12)	-0.0057 (13)
C8	0.0103 (16)	0.0103 (14)	0.0130 (15)	-0.0014 (12)	-0.0040 (12)	-0.0031 (13)
C9	0.0132 (16)	0.0184 (15)	0.0110 (15)	0.0002 (13)	-0.0023 (13)	-0.0054 (13)
C10	0.0225 (17)	0.0156 (15)	0.0202 (17)	-0.0059 (13)	-0.0059 (14)	-0.0003 (14)
C11	0.0274 (19)	0.0275 (18)	0.0245 (17)	-0.0033 (15)	-0.0169 (14)	-0.0091 (15)
C12	0.0100 (16)	0.0194 (16)	0.0137 (16)	0.0028 (13)	-0.0055 (12)	-0.0043 (14)
C13	0.0216 (17)	0.0196 (16)	0.0213 (17)	-0.0060 (14)	-0.0097 (14)	-0.0067 (14)
C14	0.0198 (17)	0.0106 (14)	0.0219 (17)	-0.0005 (13)	-0.0105 (13)	-0.0096 (13)
C15	0.0218 (18)	0.0133 (15)	0.0204 (17)	-0.0033 (13)	-0.0021 (14)	-0.0061 (14)
C16	0.034 (2)	0.039 (2)	0.0244 (19)	-0.0063 (17)	0.0040 (16)	-0.0103 (17)

Geometric parameters (Å, °)

Br1—C1	1.897 (2)	C3—H3	0.9500
S1—C15	1.730 (3)	C4—C5	1.381 (3)
S1—C14	1.735 (3)	C5—C6	1.376 (3)
O1—C7	1.371 (3)	C5—H5	0.9500
O1—C4	1.402 (3)	C6—H6	0.9500
O2—C13	1.420 (3)	C7—C8	1.378 (3)
O2—N3	1.426 (3)	C8—C9	1.422 (3)
O3—C15	1.337 (3)	C8—C12	1.442 (3)
O3—C16	1.449 (3)	C9—C10	1.492 (3)
N1—C7	1.342 (3)	C10—H10A	0.9800
N1—N2	1.365 (3)	C10—H10B	0.9800
N1—C11	1.448 (3)	C10—H10C	0.9800
N2—C9	1.333 (3)	C11—H11A	0.9800
N3—C12	1.279 (3)	C11—H11B	0.9800
N4—C14	1.293 (3)	C11—H11C	0.9800
N4—N5	1.395 (3)	C12—H12	0.9500
N5—C15	1.287 (3)	C13—C14	1.489 (3)
C1—C2	1.370 (3)	C13—H13A	0.9900
C1—C6	1.393 (3)	C13—H13B	0.9900
C2—C3	1.395 (3)	C16—H16A	0.9800
C2—H2	0.9500	C16—H16B	0.9800
C3—C4	1.376 (3)	C16—H16C	0.9800
C15—S1—C14	85.62 (13)	N2—C9—C10	121.1 (2)
C7—O1—C4	117.5 (2)	C8—C9—C10	127.0 (3)
C13—O2—N3	109.17 (19)	C9—C10—H10A	109.5
C15—O3—C16	114.5 (2)	C9—C10—H10B	109.5
C7—N1—N2	111.0 (2)	H10A—C10—H10B	109.5
C7—N1—C11	127.8 (2)	C9—C10—H10C	109.5
N2—N1—C11	121.2 (2)	H10A—C10—H10C	109.5
C9—N2—N1	105.0 (2)	H10B—C10—H10C	109.5
C12—N3—O2	108.0 (2)	N1—C11—H11A	109.5
C14—N4—N5	113.1 (2)	N1—C11—H11B	109.5
C15—N5—N4	110.5 (2)	H11A—C11—H11B	109.5
C2—C1—C6	121.0 (2)	N1—C11—H11C	109.5
C2—C1—Br1	119.8 (2)	H11A—C11—H11C	109.5
C6—C1—Br1	119.2 (2)	H11B—C11—H11C	109.5
C1—C2—C3	120.2 (2)	N3—C12—C8	122.9 (3)
C1—C2—H2	119.9	N3—C12—H12	118.5
C3—C2—H2	119.9	C8—C12—H12	118.5
C4—C3—C2	118.3 (2)	O2—C13—C14	112.9 (2)
C4—C3—H3	120.8	O2—C13—H13A	109.0
C2—C3—H3	120.8	C14—C13—H13A	109.0
C3—C4—C5	121.7 (2)	O2—C13—H13B	109.0
C3—C4—O1	124.0 (2)	C14—C13—H13B	109.0
C5—C4—O1	114.3 (2)	H13A—C13—H13B	107.8

C6—C5—C4	119.8 (3)	N4—C14—C13	123.1 (2)
C6—C5—H5	120.1	N4—C14—S1	114.4 (2)
C4—C5—H5	120.1	C13—C14—S1	122.53 (19)
C5—C6—C1	118.9 (2)	N5—C15—O3	126.0 (2)
C5—C6—H6	120.5	N5—C15—S1	116.3 (2)
C1—C6—H6	120.5	O3—C15—S1	117.6 (2)
N1—C7—O1	119.3 (3)	O3—C16—H16A	109.5
N1—C7—C8	109.1 (2)	O3—C16—H16B	109.5
O1—C7—C8	131.5 (2)	H16A—C16—H16B	109.5
C7—C8—C9	103.0 (2)	O3—C16—H16C	109.5
C7—C8—C12	130.8 (2)	H16A—C16—H16C	109.5
C9—C8—C12	126.0 (3)	H16B—C16—H16C	109.5
N2—C9—C8	111.9 (2)		
C7—N1—N2—C9	0.7 (3)	N1—C7—C8—C12	176.5 (2)
C11—N1—N2—C9	179.4 (2)	O1—C7—C8—C12	0.4 (5)
C13—O2—N3—C12	174.10 (19)	N1—N2—C9—C8	-0.1 (3)
C14—N4—N5—C15	-1.2 (3)	N1—N2—C9—C10	-179.3 (2)
C6—C1—C2—C3	0.8 (4)	C7—C8—C9—N2	-0.5 (3)
Br1—C1—C2—C3	-179.91 (19)	C12—C8—C9—N2	-176.4 (2)
C1—C2—C3—C4	-0.4 (4)	C7—C8—C9—C10	178.7 (2)
C2—C3—C4—C5	-0.7 (4)	C12—C8—C9—C10	2.8 (4)
C2—C3—C4—O1	179.6 (2)	O2—N3—C12—C8	-179.9 (2)
C7—O1—C4—C3	-1.3 (4)	C7—C8—C12—N3	6.2 (4)
C7—O1—C4—C5	179.0 (2)	C9—C8—C12—N3	-179.1 (2)
C3—C4—C5—C6	1.4 (4)	N3—O2—C13—C14	72.8 (3)
O1—C4—C5—C6	-178.9 (2)	N5—N4—C14—C13	178.9 (2)
C4—C5—C6—C1	-1.1 (4)	N5—N4—C14—S1	0.4 (3)
C2—C1—C6—C5	0.0 (4)	O2—C13—C14—N4	118.3 (3)
Br1—C1—C6—C5	-179.35 (19)	O2—C13—C14—S1	-63.3 (3)
N2—N1—C7—O1	175.63 (19)	C15—S1—C14—N4	0.4 (2)
C11—N1—C7—O1	-3.0 (4)	C15—S1—C14—C13	-178.2 (2)
N2—N1—C7—C8	-1.0 (3)	N4—N5—C15—O3	179.9 (2)
C11—N1—C7—C8	-179.6 (2)	N4—N5—C15—S1	1.5 (3)
C4—O1—C7—N1	98.1 (3)	C16—O3—C15—N5	5.5 (4)
C4—O1—C7—C8	-86.2 (3)	C16—O3—C15—S1	-176.12 (18)
N1—C7—C8—C9	0.9 (3)	C14—S1—C15—N5	-1.1 (2)
O1—C7—C8—C9	-175.2 (2)	C14—S1—C15—O3	-179.7 (2)

Hydrogen-bond geometry (Å, °)

Cg is the centroid of the C1—C6 ring.

<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 <i>A</i> ···Cg ⁱ	0.98	2.89	3.652 (4)	125

Symmetry code: (i) -x+2, -y+2, -z+2.