

{2-[(1,3-Benzothiazol-2-yl)methoxy]-5-bromophenyl}(phenyl)methanone

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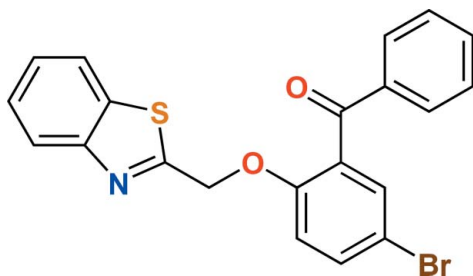
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Key indicators: single-crystal X-ray study; $T = 292$ K; mean $\sigma(\text{C}-\text{C}) = 0.007$ Å; R factor = 0.054; wR factor = 0.120; data-to-parameter ratio = 15.0.

In the title compound, $\text{C}_{21}\text{H}_{14}\text{BrNO}_2\text{S}$, the dihedral angle between the planes of the benzothiazole and phenyl-methanone groups is $63.4(2)^\circ$. In the crystal, pairs of $\text{C}-\text{H}\cdots\text{N}$ hydrogen bonds link the molecules to form inversion dimers, which are further linked by $\text{C}-\text{H}\cdots\text{O}$ interactions into chains along the c axis. $\text{C}-\text{H}\cdots\pi$ and $\pi-\pi$ interactions [centroid-centroid distance = $3.863(1)$ Å] further stabilize the molecular assembly.

Related literature

For background to the applications of benzothiazole derivatives, see: Kelarev *et al.* (2003); Rana *et al.* (2007); Telvekar *et al.* (2012); Saeed *et al.* (2010).



Experimental

Crystal data

$\text{C}_{21}\text{H}_{14}\text{BrNO}_2\text{S}$
 $M_r = 424.30$
Monoclinic, $P2_1/n$
 $a = 15.1475(6)$ Å
 $b = 7.6501(3)$ Å
 $c = 15.8339(6)$ Å
 $\beta = 102.105(3)^\circ$

$V = 1794.03(12)$ Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 2.42$ mm⁻¹
 $T = 292$ K
 $0.23 \times 0.21 \times 0.18$ mm

Data collection

Oxford Diffraction Xcalibur (Eos, Nova) diffractometer
Absorption correction: multi-scan (*CrysAlis PRO*; Oxford Diffraction, 2009)
 $T_{\min} = 0.606$, $T_{\max} = 0.670$

19116 measured reflections
3525 independent reflections
1971 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.088$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.054$
 $wR(F^2) = 0.120$
 $S = 0.98$
3525 reflections

235 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.34$ e Å⁻³
 $\Delta\rho_{\min} = -0.32$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the S1/C1/C6/N1/C7 ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C21}-\text{H21}\cdots\text{N1}^{\text{i}}$	0.93	2.55	3.398 (6)	152
$\text{C5}-\text{H5}\cdots\text{O2}^{\text{ii}}$	0.93	2.61	3.505 (6)	161
$\text{C20}-\text{H20}\cdots\text{Cg1}^{\text{iii}}$	0.93	2.68	3.459 (4)	142

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

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2009); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *PLATON* (Spek, 2009) and *PARST* (Nardelli, 1995).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FF2107).

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supplementary materials

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{2-[(1,3-Benzothiazol-2-yl)methoxy]-5-bromophenyl}(phenyl)methanone

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Comment

Substituted benzothiazole derivatives have been reported to exhibit various pharmacological properties such as analgesic, antibacterial, antifungal, antidepressant, antitumor, antihypertensive, anthelmintic, and herbicidal activity (Kelarev *et al.*, 2003). However, the variety of biological features of new benzothiazole derivatives is of great scientific interest (Telvekar *et al.*, 2012; Saeed *et al.*, 2010). Here, we report the single-crystal structure of the title compound.

The title compound prefers the conformation with the dihedral angle $63.4(2)^\circ$ between the planes of benzothiazole and phenylmethanone group (Fig. 1). The weak C—H \cdots N hydrogen bonds lead to dimer formation, whereas C—H \cdots O hydrogen bonds connect the molecules into infinite chains (Fig. 2a), which leads to formation of layers parallel to (-101). Further, the C—H \cdots π interactions involving the five membered ring S1/C1/C6/N1/C7 and π - π [$Cg2\cdots Cg3 = 3.863(1) \text{ \AA}$, $Cg2$ is the centroid of the six membered ring C9—C14 and $Cg3$ is the centroid of the six membered ring C16—C21] stabilize the criss-cross molecular assembly (Fig 2 b).

Experimental

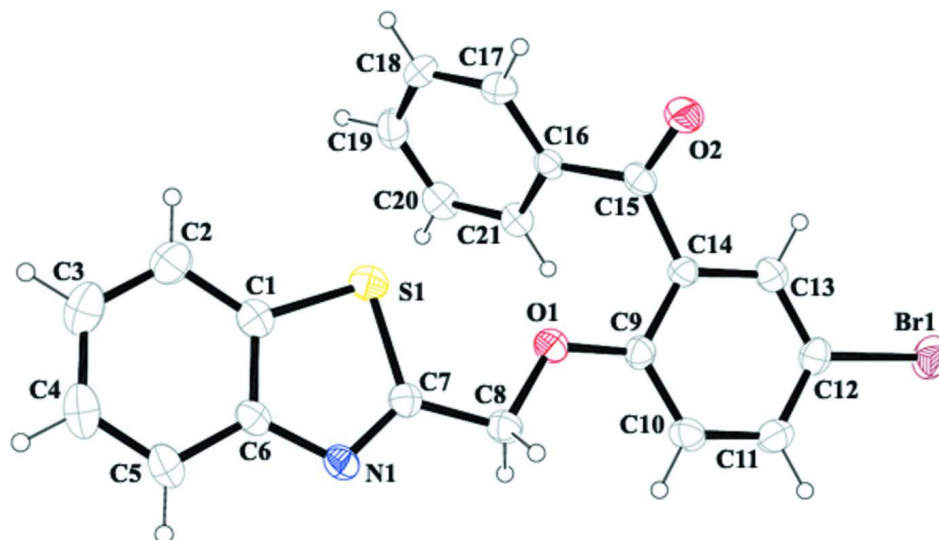
To a mixture of (2-chloromethyl)benzo[*d*]thiazole (1 mmol) and (5-bromo-2-hydroxyphenyl)(phenyl)methanone (1 mmol) in dry THF, dry potassium carbonate (1 mmol) was added and the reaction mixture was stirred at room temperature for 14 h. The reaction mixture was concentrated to remove the solvent, diluted with ethyl acetate, washed with water, brine solution and dried over anhydrous sodium sulfate. The organic layer was concentrated to yield a residue which was purified by column chromatography using ethyl acetate and n-hexane as eluent (7:3, $R_f = 0.71$) to afford the product in 77% as a white solid (m. p. $407(2) \text{ K}$). Suitable crystals for single-crystal X-ray study were obtained from acetonitrile solvent using slow evaporation technique at room temperature.

Refinement

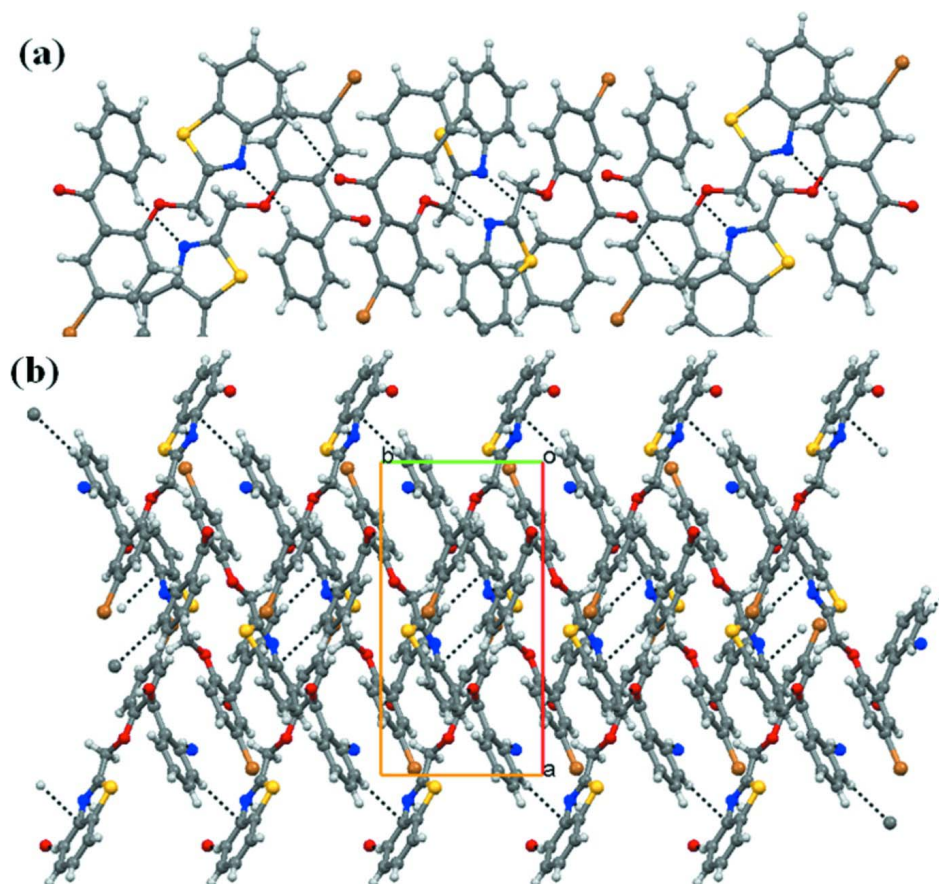
All H atoms were positioned geometrically and refined using a riding model with $U_{iso}(H) = 1.2 U_{eq}(C)$.

Computing details

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2009); cell refinement: *CrysAlis PRO* (Oxford Diffraction, 2009); data reduction: *CrysAlis PRO* (Oxford Diffraction, 2009); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and Mercury (Macrae *et al.*, 2008); software used to prepare material for publication: *PLATON* (Spek, 2009) and *PARST* (Nardelli, 1995).

**Figure 1**

Molecular structure shows the atom labelling scheme with displacement ellipsoids for non-H atoms at 30% probability level, hydrogen atoms are arbitrary circles.

**Figure 2**

(a) The dimer formation (C—H...N bonds) and their interaction by C—H...O hydrogen bonds. (b) additional C—H... π and π - π interactions stabilize the criss-cross molecular assembly; view along the *c* axis.

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Crystal data

$C_{21}H_{14}BrNO_2S$	$F(000) = 856$
$M_r = 424.30$	$D_x = 1.571 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	Melting point: 407(2) K
Hall symbol: -P 2yn	Mo $K\alpha$ radiation, $\lambda = 0.7107 \text{ \AA}$
$a = 15.1475 (6) \text{ \AA}$	Cell parameters from 350 reflections
$b = 7.6501 (3) \text{ \AA}$	$\theta = 1.0\text{--}28.0^\circ$
$c = 15.8339 (6) \text{ \AA}$	$\mu = 2.42 \text{ mm}^{-1}$
$\beta = 102.105 (3)^\circ$	$T = 292 \text{ K}$
$V = 1794.03 (12) \text{ \AA}^3$	Block, colourless
$Z = 4$	$0.23 \times 0.21 \times 0.18 \text{ mm}$

Data collection

Oxford Diffraction Xcalibur (Eos, Nova) diffractometer	19116 measured reflections
Radiation source: Mova (Mo) X-ray Source	3525 independent reflections
Mirror monochromator	1971 reflections with $I > 2\sigma(I)$
Detector resolution: 16.0839 pixels mm^{-1}	$R_{\text{int}} = 0.088$
ω scans	$\theta_{\text{max}} = 26.0^\circ$, $\theta_{\text{min}} = 2.6^\circ$
Absorption correction: multi-scan (CrysAlis PRO; Oxford Diffraction, 2009)	$h = -18 \rightarrow 18$
$T_{\text{min}} = 0.606$, $T_{\text{max}} = 0.670$	$k = -9 \rightarrow 9$
	$l = -19 \rightarrow 19$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.054$	H-atom parameters constrained
$wR(F^2) = 0.120$	$w = 1/[\sigma^2(F_o^2) + (0.0408P)^2]$
$S = 0.98$	where $P = (F_o^2 + 2F_c^2)/3$
3525 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
235 parameters	$\Delta\rho_{\text{max}} = 0.34 \text{ e \AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.32 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

Special details

Experimental. CrysAlisPro (Oxford Diffraction, 2009) Empirical absorption correction using spherical harmonics, implemented in SCALE3 ABSPACK scaling algorithm.

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	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.47887 (4)	0.70720 (7)	0.07925 (3)	0.0671 (2)
S1	-0.04002 (8)	0.33280 (15)	0.17148 (7)	0.0513 (3)

O1	0.1169 (2)	0.4165 (4)	0.11024 (17)	0.0499 (8)
N1	-0.0810 (2)	0.1671 (4)	0.0260 (2)	0.0410 (9)
O2	0.2746 (2)	0.5691 (4)	0.31997 (18)	0.0587 (9)
C6	-0.1521 (3)	0.1423 (5)	0.0671 (3)	0.0389 (10)
C15	0.2206 (3)	0.5983 (5)	0.2529 (3)	0.0395 (10)
C8	0.0664 (3)	0.3039 (5)	0.0468 (3)	0.0424 (11)
H8A	0.1004	0.1981	0.0423	0.051*
H8B	0.0539	0.3614	-0.0090	0.051*
C16	0.1297 (3)	0.6683 (5)	0.2544 (2)	0.0351 (10)
C21	0.0809 (3)	0.7618 (5)	0.1858 (3)	0.0396 (11)
H21	0.1031	0.7750	0.1357	0.048*
C9	0.1982 (3)	0.4790 (5)	0.0998 (3)	0.0397 (10)
C7	-0.0190 (3)	0.2608 (5)	0.0734 (2)	0.0379 (10)
C14	0.2497 (3)	0.5708 (5)	0.1688 (2)	0.0368 (10)
C12	0.3650 (3)	0.6114 (5)	0.0873 (3)	0.0443 (11)
C4	-0.2920 (3)	0.0222 (6)	0.0832 (4)	0.0662 (14)
H4	-0.3432	-0.0445	0.0621	0.079*
C5	-0.2287 (3)	0.0427 (6)	0.0339 (3)	0.0535 (12)
H5	-0.2364	-0.0086	-0.0204	0.064*
C18	0.0136 (3)	0.7221 (6)	0.3328 (3)	0.0554 (13)
H18	-0.0090	0.7088	0.3827	0.066*
C2	-0.2067 (3)	0.1994 (6)	0.1977 (3)	0.0600 (14)
H2	-0.1998	0.2520	0.2517	0.072*
C20	-0.0003 (3)	0.8356 (5)	0.1912 (3)	0.0507 (12)
H20	-0.0325	0.8996	0.1449	0.061*
C1	-0.1411 (3)	0.2200 (5)	0.1478 (3)	0.0438 (11)
C19	-0.0345 (3)	0.8158 (6)	0.2644 (3)	0.0560 (13)
H19	-0.0898	0.8653	0.2676	0.067*
C11	0.3151 (3)	0.5212 (6)	0.0191 (3)	0.0494 (12)
H11	0.3365	0.5058	-0.0313	0.059*
C17	0.0947 (3)	0.6485 (6)	0.3280 (3)	0.0472 (12)
H17	0.1266	0.5846	0.3745	0.057*
C10	0.2324 (3)	0.4533 (5)	0.0261 (3)	0.0470 (12)
H10	0.1990	0.3892	-0.0195	0.056*
C13	0.3333 (3)	0.6361 (5)	0.1623 (3)	0.0436 (11)
H13	0.3682	0.6967	0.2084	0.052*
C3	-0.2813 (4)	0.0995 (7)	0.1649 (4)	0.0688 (15)
H3	-0.3252	0.0830	0.1972	0.083*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0546 (4)	0.0762 (4)	0.0768 (4)	-0.0116 (3)	0.0282 (3)	-0.0030 (3)
S1	0.0522 (8)	0.0533 (8)	0.0482 (7)	-0.0063 (6)	0.0098 (6)	-0.0133 (6)
O1	0.045 (2)	0.055 (2)	0.0503 (19)	-0.0147 (16)	0.0117 (15)	-0.0175 (15)
N1	0.038 (2)	0.043 (2)	0.038 (2)	0.0021 (18)	0.0004 (18)	0.0006 (17)
O2	0.050 (2)	0.079 (2)	0.0432 (19)	0.0160 (18)	0.0014 (16)	-0.0005 (17)
C6	0.036 (3)	0.034 (2)	0.042 (3)	0.005 (2)	-0.001 (2)	0.000 (2)
C15	0.041 (3)	0.034 (2)	0.040 (3)	-0.006 (2)	0.001 (2)	0.001 (2)
C8	0.045 (3)	0.040 (3)	0.042 (3)	-0.003 (2)	0.007 (2)	-0.006 (2)

C16	0.033 (3)	0.034 (2)	0.036 (2)	-0.004 (2)	0.002 (2)	-0.0054 (19)
C21	0.044 (3)	0.030 (2)	0.044 (3)	-0.001 (2)	0.006 (2)	0.004 (2)
C9	0.035 (3)	0.039 (3)	0.045 (3)	0.002 (2)	0.009 (2)	-0.003 (2)
C7	0.037 (3)	0.037 (2)	0.038 (3)	0.007 (2)	0.004 (2)	0.004 (2)
C14	0.036 (3)	0.040 (3)	0.034 (2)	0.003 (2)	0.0060 (19)	-0.001 (2)
C12	0.042 (3)	0.038 (3)	0.055 (3)	0.005 (2)	0.014 (2)	0.002 (2)
C4	0.040 (3)	0.059 (3)	0.097 (4)	-0.002 (3)	0.010 (3)	0.008 (3)
C5	0.038 (3)	0.052 (3)	0.065 (3)	-0.003 (2)	-0.001 (2)	-0.004 (2)
C18	0.047 (3)	0.081 (4)	0.043 (3)	-0.002 (3)	0.019 (2)	-0.006 (3)
C2	0.058 (4)	0.062 (3)	0.064 (3)	0.006 (3)	0.020 (3)	-0.003 (3)
C20	0.053 (3)	0.036 (3)	0.059 (3)	0.005 (2)	0.002 (3)	0.001 (2)
C1	0.044 (3)	0.037 (3)	0.048 (3)	0.004 (2)	0.004 (2)	0.006 (2)
C19	0.039 (3)	0.057 (3)	0.072 (4)	0.000 (3)	0.011 (3)	-0.011 (3)
C11	0.053 (3)	0.053 (3)	0.044 (3)	0.006 (3)	0.016 (2)	-0.003 (2)
C17	0.051 (3)	0.053 (3)	0.037 (3)	0.001 (2)	0.007 (2)	0.003 (2)
C10	0.049 (3)	0.043 (3)	0.046 (3)	-0.003 (2)	0.006 (2)	-0.012 (2)
C13	0.041 (3)	0.041 (3)	0.047 (3)	-0.002 (2)	0.005 (2)	-0.002 (2)
C3	0.059 (4)	0.062 (4)	0.094 (4)	0.005 (3)	0.034 (3)	0.005 (3)

Geometric parameters (Å, °)

Br1—C12	1.903 (4)	C12—C11	1.368 (6)
S1—C1	1.729 (5)	C12—C13	1.383 (5)
S1—C7	1.739 (4)	C4—C5	1.366 (6)
O1—C9	1.363 (4)	C4—C3	1.399 (6)
O1—C8	1.418 (4)	C4—H4	0.9300
N1—C7	1.289 (5)	C5—H5	0.9300
N1—C6	1.382 (5)	C18—C17	1.367 (6)
O2—C15	1.218 (4)	C18—C19	1.374 (6)
C6—C1	1.388 (6)	C18—H18	0.9300
C6—C5	1.395 (5)	C2—C3	1.373 (6)
C15—C16	1.482 (5)	C2—C1	1.401 (6)
C15—C14	1.502 (5)	C2—H2	0.9300
C8—C7	1.480 (5)	C20—C19	1.372 (6)
C8—H8A	0.9700	C20—H20	0.9300
C8—H8B	0.9700	C19—H19	0.9300
C16—C21	1.379 (5)	C11—C10	1.382 (5)
C16—C17	1.387 (5)	C11—H11	0.9300
C21—C20	1.373 (6)	C17—H17	0.9300
C21—H21	0.9300	C10—H10	0.9300
C9—C10	1.385 (5)	C13—H13	0.9300
C9—C14	1.393 (5)	C3—H3	0.9300
C14—C13	1.386 (5)		
C1—S1—C7	88.2 (2)	C3—C4—H4	119.3
C9—O1—C8	119.5 (3)	C4—C5—C6	118.4 (5)
C7—N1—C6	110.3 (3)	C4—C5—H5	120.8
N1—C6—C1	114.8 (4)	C6—C5—H5	120.8
N1—C6—C5	124.5 (4)	C17—C18—C19	120.3 (4)
C1—C6—C5	120.6 (4)	C17—C18—H18	119.8

O2—C15—C16	120.6 (4)	C19—C18—H18	119.8
O2—C15—C14	118.6 (4)	C3—C2—C1	118.2 (5)
C16—C15—C14	120.8 (4)	C3—C2—H2	120.9
O1—C8—C7	107.9 (3)	C1—C2—H2	120.9
O1—C8—H8A	110.1	C19—C20—C21	120.6 (4)
C7—C8—H8A	110.1	C19—C20—H20	119.7
O1—C8—H8B	110.1	C21—C20—H20	119.7
C7—C8—H8B	110.1	C6—C1—C2	120.6 (4)
H8A—C8—H8B	108.4	C6—C1—S1	110.0 (3)
C21—C16—C17	118.8 (4)	C2—C1—S1	129.4 (4)
C21—C16—C15	121.4 (4)	C20—C19—C18	119.5 (5)
C17—C16—C15	119.7 (4)	C20—C19—H19	120.3
C20—C21—C16	120.3 (4)	C18—C19—H19	120.3
C20—C21—H21	119.9	C12—C11—C10	119.1 (4)
C16—C21—H21	119.9	C12—C11—H11	120.5
O1—C9—C10	124.0 (4)	C10—C11—H11	120.5
O1—C9—C14	116.9 (4)	C18—C17—C16	120.5 (4)
C10—C9—C14	119.0 (4)	C18—C17—H17	119.7
N1—C7—C8	122.1 (4)	C16—C17—H17	119.7
N1—C7—S1	116.6 (3)	C11—C10—C9	121.3 (4)
C8—C7—S1	121.2 (3)	C11—C10—H10	119.3
C13—C14—C9	119.6 (4)	C9—C10—H10	119.3
C13—C14—C15	117.3 (4)	C12—C13—C14	120.1 (4)
C9—C14—C15	123.1 (4)	C12—C13—H13	119.9
C11—C12—C13	120.8 (4)	C14—C13—H13	119.9
C11—C12—Br1	120.0 (3)	C2—C3—C4	120.8 (5)
C13—C12—Br1	119.2 (3)	C2—C3—H3	119.6
C5—C4—C3	121.3 (5)	C4—C3—H3	119.6
C5—C4—H4	119.3		
C7—N1—C6—C1	-0.4 (5)	C1—C6—C5—C4	-0.6 (6)
C7—N1—C6—C5	-178.1 (4)	C16—C21—C20—C19	-0.7 (6)
C9—O1—C8—C7	-178.5 (3)	N1—C6—C1—C2	-177.6 (4)
O2—C15—C16—C21	155.4 (4)	C5—C6—C1—C2	0.2 (6)
C14—C15—C16—C21	-21.3 (6)	N1—C6—C1—S1	1.8 (4)
O2—C15—C16—C17	-20.9 (6)	C5—C6—C1—S1	179.5 (3)
C14—C15—C16—C17	162.4 (4)	C3—C2—C1—C6	0.5 (7)
C17—C16—C21—C20	0.9 (6)	C3—C2—C1—S1	-178.7 (4)
C15—C16—C21—C20	-175.5 (4)	C7—S1—C1—C6	-1.9 (3)
C8—O1—C9—C10	6.4 (6)	C7—S1—C1—C2	177.4 (4)
C8—O1—C9—C14	-172.2 (4)	C21—C20—C19—C18	0.5 (7)
C6—N1—C7—C8	178.2 (4)	C17—C18—C19—C20	-0.4 (7)
C6—N1—C7—S1	-1.2 (4)	C13—C12—C11—C10	-0.5 (6)
O1—C8—C7—N1	176.5 (4)	Br1—C12—C11—C10	-179.7 (3)
O1—C8—C7—S1	-4.2 (5)	C19—C18—C17—C16	0.5 (7)
C1—S1—C7—N1	1.8 (3)	C21—C16—C17—C18	-0.7 (6)
C1—S1—C7—C8	-177.5 (3)	C15—C16—C17—C18	175.7 (4)
O1—C9—C14—C13	179.4 (4)	C12—C11—C10—C9	1.8 (6)
C10—C9—C14—C13	0.7 (6)	O1—C9—C10—C11	179.5 (4)

O1—C9—C14—C15	1.6 (6)	C14—C9—C10—C11	-1.9 (6)
C10—C9—C14—C15	-177.1 (4)	C11—C12—C13—C14	-0.6 (6)
O2—C15—C14—C13	-45.6 (5)	Br1—C12—C13—C14	178.6 (3)
C16—C15—C14—C13	131.2 (4)	C9—C14—C13—C12	0.5 (6)
O2—C15—C14—C9	132.2 (4)	C15—C14—C13—C12	178.4 (4)
C16—C15—C14—C9	-50.9 (6)	C1—C2—C3—C4	-0.7 (7)
C3—C4—C5—C6	0.4 (7)	C5—C4—C3—C2	0.3 (8)
N1—C6—C5—C4	176.9 (4)		

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

Cg1 is the centroid of the S1/C1/C6/N1/C7 ring.

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C21—H21 \cdots N1 ⁱ	0.93	2.55	3.398 (6)	152
C5—H5 \cdots O2 ⁱⁱ	0.93	2.61	3.505 (6)	161
C20—H20 \cdots Cg1 ⁱⁱⁱ	0.93	2.68	3.459 (4)	142

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