

3,3-Bis[(4-chlorophenyl)sulfanyl]-1-methylpiperidin-2-one

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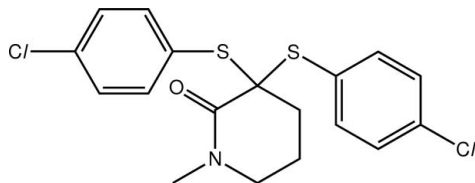
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Key indicators: single-crystal X-ray study; $T = 290$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.043; wR factor = 0.121; data-to-parameter ratio = 15.1.

The piperidone ring in the title compound, $\text{C}_{18}\text{H}_{17}\text{Cl}_2\text{NOS}_2$, has a distorted half-chair conformation. The S-bound benzene rings are approximately perpendicular to and splayed out of the mean plane through the piperidone ring [dihedral angles = 71.86 (13) and 46.94 (11)°]. In the crystal, C—H···O interactions link the molecules into [010] supramolecular chains with a helical topology. C—H···Cl and C—H··· π interactions are also present.

Related literature

For background to β -thiocarbonyl compounds, see: Vinhato (2007); Olivato *et al.* (2009). For related structures, see: Zukerman-Schpector *et al.* (2006, 2008). For ring conformational analysis, see: Cremer & Pople (1975). For further synthetic details, see: Hashmat & McDermott (2002); Zoretic & Soja (1976).



Experimental

Crystal data

$\text{C}_{18}\text{H}_{17}\text{Cl}_2\text{NOS}_2$
 $M_r = 398.37$
Monoclinic, $P2_1/n$
 $a = 8.0313$ (2) Å
 $b = 9.7460$ (2) Å
 $c = 24.2623$ (7) Å
 $\beta = 94.0767$ (12)°

$V = 1894.28$ (8) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.57$ mm⁻¹
 $T = 290$ K
 $0.33 \times 0.30 \times 0.29$ mm

Data collection

Nonius KappaCCD diffractometer
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.82$, $T_{\max} = 0.85$
12888 measured reflections
3288 independent reflections
2778 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.049$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.121$
 $S = 1.05$
3288 reflections
218 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.37$ e Å⁻³
 $\Delta\rho_{\min} = -0.38$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the C7–C12.

D—H···A	D—H	H···A	D···A	D—H···A
C9—H9···O1 ⁱ	0.93	2.32	3.218 (3)	164
C11—H11···Cl2 ⁱⁱ	0.93	2.83	3.708 (3)	157
C19—H19a···Cg1 ⁱⁱⁱ	0.96	2.95	3.676 (3)	133

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

Data collection: *COLLECT* (Nonius, 1999); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* (Otwinowski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5512).

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supporting information

Acta Cryst. (2010). E66, o1863 [doi:10.1107/S1600536810024347]

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S1. Comment

As part of our on-going research on the conformational and electronic interactions in β -thio-carbonyl compounds, *e.g.* *N,N*-diethyl-2-[(4'-substituted) phenylthio]acetamides, *N*-methoxy-*N*-methyl-2-[(4'-substituted) phenylthio]-propanamides, and 1-methyl-3-phenylsulfonyl-2-piperidone, utilizing spectroscopic, theoretical and X-ray diffraction methods (Vinhato, 2007; Olivato *et al.*, 2009; Zukerman-Schpector *et al.* 2008), the title compound, (I), was synthesized and its crystal structure determined.

In (I), Fig. 1, the piperidone ring has a distorted half-chair conformation: the ring-puckering parameters are $q_2 = 0.453(2) \text{ \AA}$, $q_3 = -0.271(2) \text{ \AA}$, $QT = 0.528(3) \text{ \AA}$, $\varphi_2 = 37.4(3)^\circ$ (Cremer & Pople, 1975). While the S2-bound benzene ring is orientated to be almost perpendicular to the plane through the piperidone ring [dihedral angle = $71.86(13)^\circ$], the S1-bound benzene ring is somewhat splayed with respect to the other rings, forming dihedral angles of $46.94(11)$ and $61.68(13)^\circ$ with those through the piperidone and S2-bound benzene rings, respectively.

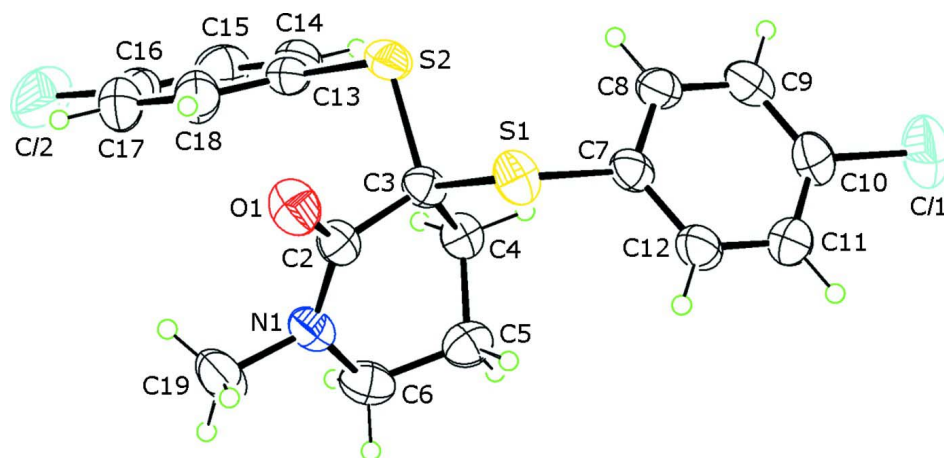
Supramolecular helical chains aligned along the *b* axis dominate the crystal packing, Fig. 2 and Table 1, and these are sustained in the crystal structure by C–H \cdots Cl and C–H \cdots π interactions, Table 1.

S2. Experimental

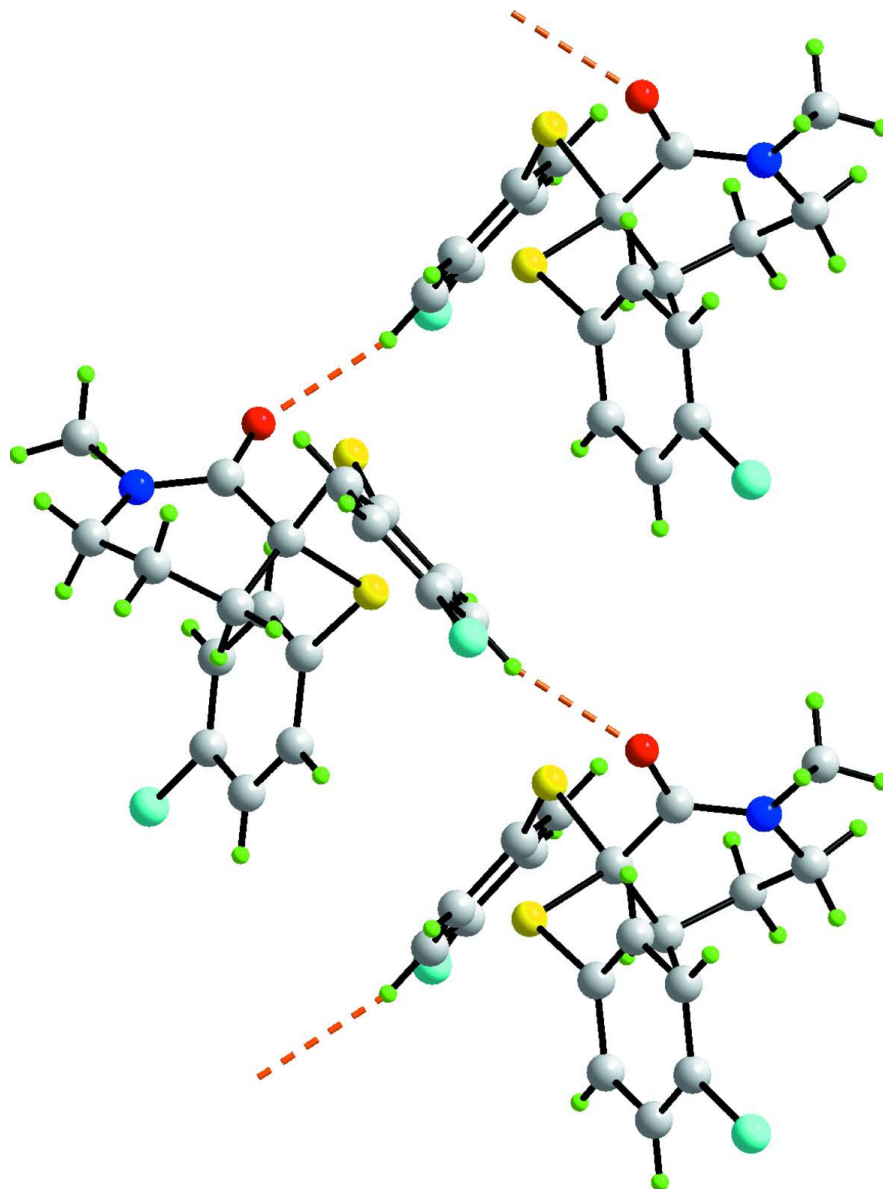
Firstly, 4-chlorothiophenol (5.8 g, 40 mmol) was reacted with bromine (1.1 ml, 40 mmol) in dichloromethane (250 ml) on hydrated silica gel support (25 g of SiO₂ and 12 ml of water) to give 4-chlorophenyl disulfide (5.3 g, yield = 93%). A yellow solid was obtained after filtration and evaporation without further purification (Hashmat & McDermott, 2002). 1-Methyl-2-piperidinone (2.0 g, 18 mmol) was added dropwise to a cooled (195 K) solution of hexamethylphosphoramide (HMPA) (3.3 ml, 18 mmol), diisopropylamine (2.6 ml, 18 mmol) and butyllithium (11.5 ml, 18 mmol) in THF (60 ml). After 20 minutes, 4-chlorophenyl disulfide (5.3 g, 18 mmol) dissolved in THF (10 ml) was added dropwise to the enolate solution (Zoretic & Soja, 1976). After stirring for 3 h at 195 K, water (80 ml) was added at room temperature and extraction with chloroform was performed. The organic layer was dried over anhydrous sodium sulfate. After evaporation of solvent, a crude solid was obtained. Purification through flash chromatography with a solution of hexane and ethyl acetate in a 7:3 ratio give the pure product (2.8 g, yield = 35%). Irregular lumps of (I) were obtained by vapour diffusion of *n*-hexane into a chloroform solution held at 283 K; m.p. 372–373 K. IR (cm⁻¹): $\nu(\text{C=O})$ 1663. NMR (CDCl₃, p.p.m.): δ 1.93–1.95 (2H, m), 1.97–1.99 (2H, m), 2.91 (3H, s), 3.21 (2H, triplet, $J = 6.0$ Hz), 7.31–7.33 (4H, m, Aryl-H), 7.55–7.57 (4H, m, Aryl-H). Analysis found: C 54.33, H 4.30, N 3.39%. C₁₈H₁₇OCl₂NS₂ requires: C 54.27, H 4.30, N 3.52%.

S3. Refinement

The H atoms were geometrically placed (C–H = 0.93–0.97 Å) and refined as riding with $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5U_{\text{eq}}(\text{C})$.

**Figure 1**

The molecular structure of (I) showing displacement ellipsoids at the 35% probability level (arbitrary spheres for the H atoms).

**Figure 2**

Supramolecular chain in (I) mediated by C–H···O interactions (orange dashed lines). The chain with helical topology is aligned along the *b* axis.

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Crystal data

$C_{18}H_{17}Cl_2NOS_2$

$M_r = 398.37$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1/n$

$a = 8.0313\ (2)\ \text{\AA}$

$b = 9.7460\ (2)\ \text{\AA}$

$c = 24.2623\ (7)\ \text{\AA}$

$\beta = 94.0767\ (12)^\circ$

$V = 1894.28\ (8)\ \text{\AA}^3$

$Z = 4$

$F(000) = 824$

$D_x = 1.397\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 10679 reflections

$\theta = 2.9\text{--}27.5^\circ$

$\mu = 0.57\ \text{mm}^{-1}$

$T = 290$ K $0.33 \times 0.30 \times 0.29$ mm
 Irregular, colourless

Data collection

Nonius KappaCCD diffractometer	12888 measured reflections 3288 independent reflections
Radiation source: sealed tube	2778 reflections with $I > 2\sigma(I)$
Graphite monochromator	$R_{\text{int}} = 0.049$
CCD rotation images scans	$\theta_{\text{max}} = 25.0^\circ$, $\theta_{\text{min}} = 3.3^\circ$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$h = -9 \rightarrow 8$
$T_{\text{min}} = 0.82$, $T_{\text{max}} = 0.85$	$k = -11 \rightarrow 11$ $l = -28 \rightarrow 26$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.043$	H-atom parameters constrained
$wR(F^2) = 0.121$	$w = 1/[\sigma^2(F_o^2) + (0.0558P)^2 + 0.758P]$
$S = 1.05$	where $P = (F_o^2 + 2F_c^2)/3$
3288 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
218 parameters	$\Delta\rho_{\text{max}} = 0.37 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.38 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

Special details

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 $F^2 > \sigma(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
C2	0.3322 (3)	0.3195 (2)	0.66893 (9)	0.0514 (5)
C3	0.4451 (2)	0.2265 (2)	0.70636 (9)	0.0490 (5)
C4	0.3506 (3)	0.1218 (2)	0.73838 (9)	0.0527 (5)
H4A	0.3189	0.0448	0.7145	0.063*
H4B	0.4223	0.0876	0.7692	0.063*
C5	0.1959 (3)	0.1856 (3)	0.75971 (10)	0.0629 (6)
H5A	0.1407	0.1199	0.7823	0.075*
H5B	0.2264	0.2650	0.7823	0.075*
C6	0.0808 (3)	0.2277 (3)	0.71127 (11)	0.0697 (7)
H6A	0.0309	0.1464	0.6940	0.084*
H6B	-0.0085	0.2835	0.7243	0.084*
C7	0.6208 (2)	0.2649 (2)	0.81264 (9)	0.0520 (5)
C8	0.7382 (3)	0.1601 (2)	0.81591 (10)	0.0557 (5)
H8	0.7841	0.1296	0.7840	0.067*

C9	0.7876 (3)	0.1008 (2)	0.86615 (11)	0.0632 (6)
H9	0.8659	0.0303	0.8681	0.076*
C10	0.7204 (3)	0.1464 (3)	0.91293 (11)	0.0678 (7)
C11	0.6026 (3)	0.2500 (3)	0.91080 (11)	0.0756 (7)
H11	0.5572	0.2800	0.9429	0.091*
C12	0.5534 (3)	0.3082 (3)	0.86065 (10)	0.0673 (7)
H12	0.4737	0.3776	0.8589	0.081*
C13	0.4783 (3)	0.0506 (2)	0.61552 (9)	0.0572 (5)
C14	0.4608 (3)	-0.0901 (3)	0.61981 (11)	0.0687 (6)
H14	0.5093	-0.1360	0.6505	0.082*
C15	0.3712 (4)	-0.1629 (3)	0.57846 (13)	0.0814 (8)
H15	0.3600	-0.2576	0.5811	0.098*
C16	0.2998 (3)	-0.0938 (4)	0.53379 (12)	0.0798 (8)
C17	0.3157 (4)	0.0450 (4)	0.52858 (12)	0.0869 (9)
H17	0.2660	0.0900	0.4979	0.104*
C18	0.4061 (4)	0.1180 (3)	0.56924 (11)	0.0744 (7)
H18	0.4188	0.2123	0.5657	0.089*
C19	0.0614 (4)	0.3812 (3)	0.62912 (13)	0.0835 (8)
H19A	0.0725	0.4778	0.6362	0.125*
H19B	-0.0530	0.3545	0.6314	0.125*
H19C	0.0954	0.3612	0.5928	0.125*
O1	0.3934 (2)	0.40242 (17)	0.63843 (7)	0.0719 (5)
Cl1	0.78388 (14)	0.07472 (10)	0.97627 (4)	0.1129 (3)
Cl2	0.19021 (13)	-0.18374 (14)	0.48074 (4)	0.1289 (4)
S1	0.56494 (8)	0.35278 (6)	0.75025 (3)	0.0615 (2)
S2	0.60276 (7)	0.14457 (7)	0.66590 (3)	0.0638 (2)
N1	0.1666 (2)	0.3052 (2)	0.67012 (8)	0.0589 (5)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C2	0.0504 (12)	0.0463 (11)	0.0566 (12)	-0.0034 (9)	-0.0033 (9)	-0.0035 (9)
C3	0.0412 (11)	0.0466 (11)	0.0588 (12)	-0.0035 (8)	-0.0002 (9)	-0.0038 (9)
C4	0.0553 (12)	0.0434 (11)	0.0585 (13)	-0.0066 (9)	-0.0020 (10)	-0.0008 (9)
C5	0.0609 (14)	0.0606 (13)	0.0688 (15)	-0.0116 (11)	0.0150 (11)	-0.0024 (11)
C6	0.0474 (13)	0.0693 (15)	0.0928 (19)	-0.0050 (11)	0.0079 (12)	-0.0031 (13)
C7	0.0425 (11)	0.0500 (11)	0.0621 (12)	-0.0023 (9)	-0.0056 (9)	-0.0108 (10)
C8	0.0446 (11)	0.0576 (12)	0.0646 (14)	0.0018 (9)	0.0013 (10)	-0.0100 (10)
C9	0.0493 (12)	0.0538 (13)	0.0846 (17)	0.0044 (10)	-0.0083 (11)	-0.0089 (12)
C10	0.0654 (15)	0.0707 (15)	0.0645 (15)	-0.0047 (12)	-0.0161 (12)	-0.0043 (12)
C11	0.0682 (16)	0.0962 (19)	0.0615 (15)	0.0109 (14)	-0.0027 (12)	-0.0207 (14)
C12	0.0574 (14)	0.0725 (15)	0.0699 (16)	0.0178 (12)	-0.0097 (11)	-0.0204 (12)
C13	0.0451 (11)	0.0698 (14)	0.0566 (13)	0.0102 (10)	0.0043 (9)	-0.0087 (10)
C14	0.0633 (15)	0.0689 (16)	0.0735 (16)	0.0100 (12)	0.0019 (12)	-0.0054 (12)
C15	0.0790 (19)	0.0766 (17)	0.090 (2)	-0.0058 (14)	0.0131 (15)	-0.0209 (15)
C16	0.0612 (16)	0.109 (2)	0.0690 (17)	-0.0075 (15)	0.0079 (13)	-0.0283 (16)
C17	0.086 (2)	0.112 (2)	0.0606 (16)	0.0151 (17)	-0.0100 (14)	-0.0096 (16)
C18	0.0828 (18)	0.0781 (17)	0.0609 (15)	0.0126 (14)	-0.0043 (13)	-0.0028 (12)

C19	0.0666 (17)	0.0893 (19)	0.0905 (19)	0.0192 (14)	-0.0237 (14)	-0.0028 (15)
O1	0.0693 (11)	0.0663 (10)	0.0790 (11)	-0.0096 (8)	-0.0020 (9)	0.0199 (9)
C11	0.1408 (8)	0.1145 (7)	0.0781 (5)	0.0100 (6)	-0.0305 (5)	0.0123 (5)
C12	0.1044 (7)	0.1885 (11)	0.0944 (6)	-0.0460 (7)	0.0104 (5)	-0.0639 (7)
S1	0.0640 (4)	0.0479 (3)	0.0702 (4)	-0.0118 (2)	-0.0113 (3)	-0.0013 (2)
S2	0.0407 (3)	0.0814 (4)	0.0688 (4)	0.0036 (3)	-0.0002 (3)	-0.0123 (3)
N1	0.0470 (10)	0.0599 (11)	0.0686 (12)	0.0035 (8)	-0.0055 (8)	-0.0025 (9)

Geometric parameters (Å, °)

C2—O1	1.222 (3)	C10—C11	1.382 (4)
C2—N1	1.340 (3)	C10—C11	1.731 (3)
C2—C3	1.533 (3)	C11—C12	1.375 (4)
C3—C4	1.518 (3)	C11—H11	0.9300
C3—S2	1.839 (2)	C12—H12	0.9300
C3—S1	1.851 (2)	C13—C14	1.383 (4)
C4—C5	1.513 (3)	C13—C18	1.391 (3)
C4—H4A	0.9700	C13—S2	1.777 (2)
C4—H4B	0.9700	C14—C15	1.388 (4)
C5—C6	1.500 (3)	C14—H14	0.9300
C5—H5A	0.9700	C15—C16	1.367 (4)
C5—H5B	0.9700	C15—H15	0.9300
C6—N1	1.463 (3)	C16—C17	1.366 (4)
C6—H6A	0.9700	C16—C12	1.743 (3)
C6—H6B	0.9700	C17—C18	1.380 (4)
C7—C12	1.385 (3)	C17—H17	0.9300
C7—C8	1.388 (3)	C18—H18	0.9300
C7—S1	1.769 (2)	C19—N1	1.460 (3)
C8—C9	1.381 (3)	C19—H19A	0.9600
C8—H8	0.9300	C19—H19B	0.9600
C9—C10	1.365 (4)	C19—H19C	0.9600
C9—H9	0.9300		
O1—C2—N1	121.6 (2)	C9—C10—C11	119.9 (2)
O1—C2—C3	120.21 (19)	C11—C10—C11	119.0 (2)
N1—C2—C3	118.18 (19)	C12—C11—C10	119.2 (2)
C4—C3—C2	113.84 (17)	C12—C11—H11	120.4
C4—C3—S2	111.66 (14)	C10—C11—H11	120.4
C2—C3—S2	109.95 (14)	C11—C12—C7	120.9 (2)
C4—C3—S1	114.26 (15)	C11—C12—H12	119.5
C2—C3—S1	102.08 (13)	C7—C12—H12	119.5
S2—C3—S1	104.26 (10)	C14—C13—C18	119.3 (2)
C5—C4—C3	110.55 (17)	C14—C13—S2	120.93 (19)
C5—C4—H4A	109.5	C18—C13—S2	119.6 (2)
C3—C4—H4A	109.5	C13—C14—C15	120.2 (3)
C5—C4—H4B	109.5	C13—C14—H14	119.9
C3—C4—H4B	109.5	C15—C14—H14	119.9
H4A—C4—H4B	108.1	C16—C15—C14	119.2 (3)

C6—C5—C4	108.67 (19)	C16—C15—H15	120.4
C6—C5—H5A	110.0	C14—C15—H15	120.4
C4—C5—H5A	110.0	C17—C16—C15	121.6 (3)
C6—C5—H5B	110.0	C17—C16—C12	118.4 (3)
C4—C5—H5B	110.0	C15—C16—C12	119.9 (3)
H5A—C5—H5B	108.3	C16—C17—C18	119.5 (3)
N1—C6—C5	112.44 (19)	C16—C17—H17	120.2
N1—C6—H6A	109.1	C18—C17—H17	120.2
C5—C6—H6A	109.1	C17—C18—C13	120.1 (3)
N1—C6—H6B	109.1	C17—C18—H18	119.9
C5—C6—H6B	109.1	C13—C18—H18	119.9
H6A—C6—H6B	107.8	N1—C19—H19A	109.5
C12—C7—C8	118.7 (2)	N1—C19—H19B	109.5
C12—C7—S1	118.79 (17)	H19A—C19—H19B	109.5
C8—C7—S1	122.34 (17)	N1—C19—H19C	109.5
C9—C8—C7	120.6 (2)	H19A—C19—H19C	109.5
C9—C8—H8	119.7	H19B—C19—H19C	109.5
C7—C8—H8	119.7	C7—S1—C3	105.06 (10)
C10—C9—C8	119.5 (2)	C13—S2—C3	102.48 (9)
C10—C9—H9	120.3	C2—N1—C19	117.4 (2)
C8—C9—H9	120.3	C2—N1—C6	125.80 (19)
C9—C10—C11	121.1 (2)	C19—N1—C6	116.7 (2)
O1—C2—C3—C4	-175.5 (2)	C14—C15—C16—C17	0.6 (4)
N1—C2—C3—C4	3.4 (3)	C14—C15—C16—C12	178.7 (2)
O1—C2—C3—S2	-49.4 (2)	C15—C16—C17—C18	0.0 (5)
N1—C2—C3—S2	129.55 (18)	C12—C16—C17—C18	-178.1 (2)
O1—C2—C3—S1	60.9 (2)	C16—C17—C18—C13	-0.9 (4)
N1—C2—C3—S1	-120.22 (18)	C14—C13—C18—C17	1.1 (4)
C2—C3—C4—C5	-40.9 (2)	S2—C13—C18—C17	177.0 (2)
S2—C3—C4—C5	-166.11 (15)	C12—C7—S1—C3	-113.91 (19)
S1—C3—C4—C5	75.9 (2)	C8—C7—S1—C3	70.55 (19)
C3—C4—C5—C6	63.6 (2)	C4—C3—S1—C7	29.71 (18)
C4—C5—C6—N1	-48.4 (3)	C2—C3—S1—C7	153.07 (14)
C12—C7—C8—C9	-0.4 (3)	S2—C3—S1—C7	-92.45 (12)
S1—C7—C8—C9	175.18 (17)	C14—C13—S2—C3	-104.0 (2)
C7—C8—C9—C10	-0.3 (3)	C18—C13—S2—C3	80.2 (2)
C8—C9—C10—C11	0.6 (4)	C4—C3—S2—C13	67.21 (17)
C8—C9—C10—C11	-178.95 (18)	C2—C3—S2—C13	-60.15 (16)
C9—C10—C11—C12	-0.3 (4)	S1—C3—S2—C13	-168.93 (11)
C11—C10—C11—C12	179.3 (2)	O1—C2—N1—C19	6.9 (3)
C10—C11—C12—C7	-0.3 (4)	C3—C2—N1—C19	-172.0 (2)
C8—C7—C12—C11	0.7 (4)	O1—C2—N1—C6	-168.6 (2)
S1—C7—C12—C11	-175.0 (2)	C3—C2—N1—C6	12.5 (3)
C18—C13—C14—C15	-0.4 (4)	C5—C6—N1—C2	11.0 (3)
S2—C13—C14—C15	-176.2 (2)	C5—C6—N1—C19	-164.5 (2)
C13—C14—C15—C16	-0.5 (4)		

Hydrogen-bond geometry (Å, °)

Cg1 is the centroid of the C7–C12.

<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>
C9—H9···O1 ⁱ	0.93	2.32	3.218 (3)	164
C11—H11···C12 ⁱⁱ	0.93	2.83	3.708 (3)	157
C19—H19a···Cg1 ⁱⁱⁱ	0.96	2.95	3.676 (3)	133

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