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## Structure Reports

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## 4-(Thiophen-2-yl)-2-[4-(trifluoromethyl)-phenyl]-2,3-dihydro-1,5-benzothiazepine

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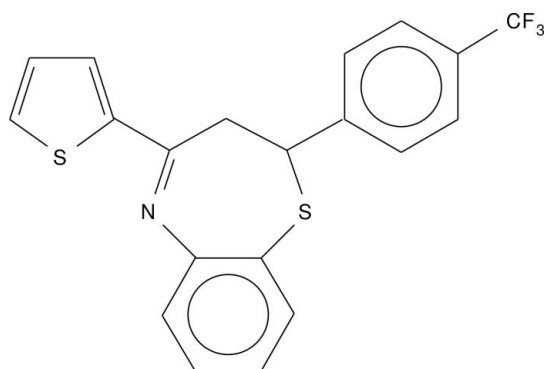
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Key indicators: single-crystal X-ray study;  $T = 296$  K; mean  $\sigma(\text{C}-\text{C}) = 0.009$  Å;  $R$  factor = 0.085;  $wR$  factor = 0.239; data-to-parameter ratio = 12.8.

In the title compound,  $\text{C}_{20}\text{H}_{14}\text{F}_3\text{NS}_2$ , the seven-membered thiazepine ring adopts a slightly distorted twist-boat conformation. The mean plane of the five-membered thiophene ring fused to the thiazepine ring is twisted by  $32.3$  (3) and  $55.6$  (4)° from the benzene and phenyl rings, respectively. In the crystal, inversion dimers linked by pairs of weak  $\text{C}-\text{H}\cdots\text{N}$  interactions are observed.

## Related literature

For the biological activity of 1, 4-thiazepines, see: Skiles *et al.* (1986); Zeng & Alper (2010). For a related structure, see: Manjula *et al.* (2013). For standard bond lengths, see: Allen *et al.* (1987).



## Experimental

## Crystal data

 $\text{C}_{20}\text{H}_{14}\text{F}_3\text{NS}_2$   
 $M_r = 389.46$   
Orthorhombic,  $P2_12_12_1$   
 $a = 9.847$  (2) Å  
 $b = 10.492$  (3) Å  
 $c = 17.819$  (4) Å  
 $V = 1841.0$  (8) Å<sup>3</sup>  
 $Z = 4$   
Cu  $K\alpha$  radiation  
 $\mu = 2.92$  mm<sup>-1</sup>  
 $T = 296$  K  
 $0.20 \times 0.19 \times 0.18$  mm

## Data collection

Bruker X8 Proteum diffractometer  
Absorption correction: multi-scan  
(SADABS; Bruker, 2013)  
 $T_{\min} = 0.593$ ,  $T_{\max} = 0.622$   
12103 measured reflections  
3028 independent reflections  
2547 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.056$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.085$   
 $wR(F^2) = 0.239$   
 $S = 1.03$   
3028 reflections  
237 parameters  
H-atom parameters constrained  
 $\Delta\rho_{\text{max}} = 0.94$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.51$  e Å<sup>-3</sup>  
Absolute structure: Flack (1983),  
1270 Friedel pairs  
Absolute structure parameter:  
0.0 (3)

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C21}-\text{H21}\cdots\text{N12}^i$	0.93	2.52	3.262 (7)	137

Symmetry code: (i)  $-x, y - \frac{1}{2}, -z + \frac{1}{2}$ .

Data collection: *APEX2* (Bruker, 2013); cell refinement: *SAINT* (Bruker, 2013); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *Mercury*.

The authors thank the IOE and the University of Mysore for providing the single-crystal X-ray diffractometer facility.

Supporting information for this paper is available from the IUCr electronic archives (Reference: JJ2182).

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## supplementary materials

*Acta Cryst.* (2014). E70, o261 [doi:10.1107/S1600536814002529]

## 4-(Thiophen-2-yl)-2-[4-(trifluoromethyl)phenyl]-2,3-dihydro-1,5-benzothiazepine

B. C. Manjunath, M. Manjula, K. R. Raghavendra, K. Ajay Kumar and N. K. Lokanath

### 1. Comment

1,4-Thiazepine is a privileged structure because of its presence in a number of pharmacologically important compounds. 1, 4-thiazepine moieties represent an important class of heterocyclic compounds with diverse potential and pharmacological activities. They are used as calcium channel blockers, HIV-1 integrase and reverse transcriptase inhibitors, antitumor, antiplatelet and antidepressant agents (Zeng *et al.*, 2010). Compounds containing the 1, 4-thiazepine moiety are important targets in synthetic and medicinal chemistry because this fragment is a key motif in a wide number of natural and synthetic biologically active agents (Skiles *et al.*, 1986).

In the title compound, (I), the thiazepine ring adopts a slightly distorted twist-boat conformation. The mean plane of the five-membered thiophene ring fused to the thiazepine ring is twisted by 32.3 (3)° and 55.6 (4)° from the mean planes of the two benzene rings. In the crystal, the molecules are linked by weak C—H···N intermolecular interactions. The seven membered thiazepine ring adopts a twist boat conformation with the pairs of atoms N12/C11 and S9/C10 being oppositely oriented with respect to the C6/C7/C8 mean plane. The torsion angles around C6—C7 and C7—C8 are very close to the value of 52° reported for the corresponding torsion angle in the ideal twist boat conformation of cycloheptane. The bond lengths between C6—N12 is 1.29 (3)Å and C11—N12 is 1.39 (3)Å which is slightly less than the standard C—N value. The bond length between C8—S9 is 1.83 (9)Å and S9—C10 is 1.76 (3) Å. The sp<sup>2</sup> and sp<sup>3</sup> hybridization states of C10 and C8, respectively, account for the difference in the S—C bond lengths in (I). The bond lengths and angles do not show large deviations and are comparable with those reported for a similar structure (Manjula *et al.*, 2013).

### 2. Experimental

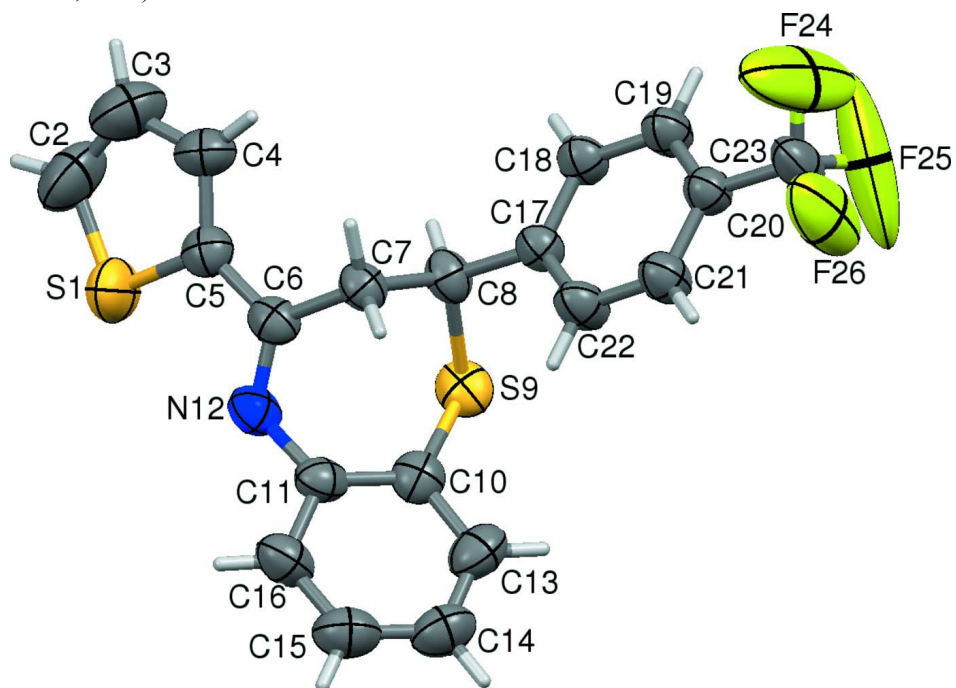
A mixture of (*Z*)-1-(thiophen-2-yl)-3-(4-(trifluoromethyl) phenyl)prop-2-en-1- one (3 mmol, 1 g) and 2-aminithiophenol (3 mmol, 0.4 g) with 3–4 drops of conc. HCl in methanol (10 ml) was heated with stirring at 433 K for 4 h. The reaction was monitored by thin-layer chromatography (hexane/ethyl acetate). After the completion of the reaction, the mixture was extracted in chloroform (30 ml), washed successively with dilute hydrochloric acid and water. The solvent was evaporated to dryness. The solid obtained was crystallized from 95° ethyl alcohol to get pale yellow needles of 4-(thiophen-2-yl)-2-(4-(trifluoromethyl) phenyl)-2,3-dihydrobenzo [*b*] [1,4] thiazepine in 80° yield.

### 3. Refinement

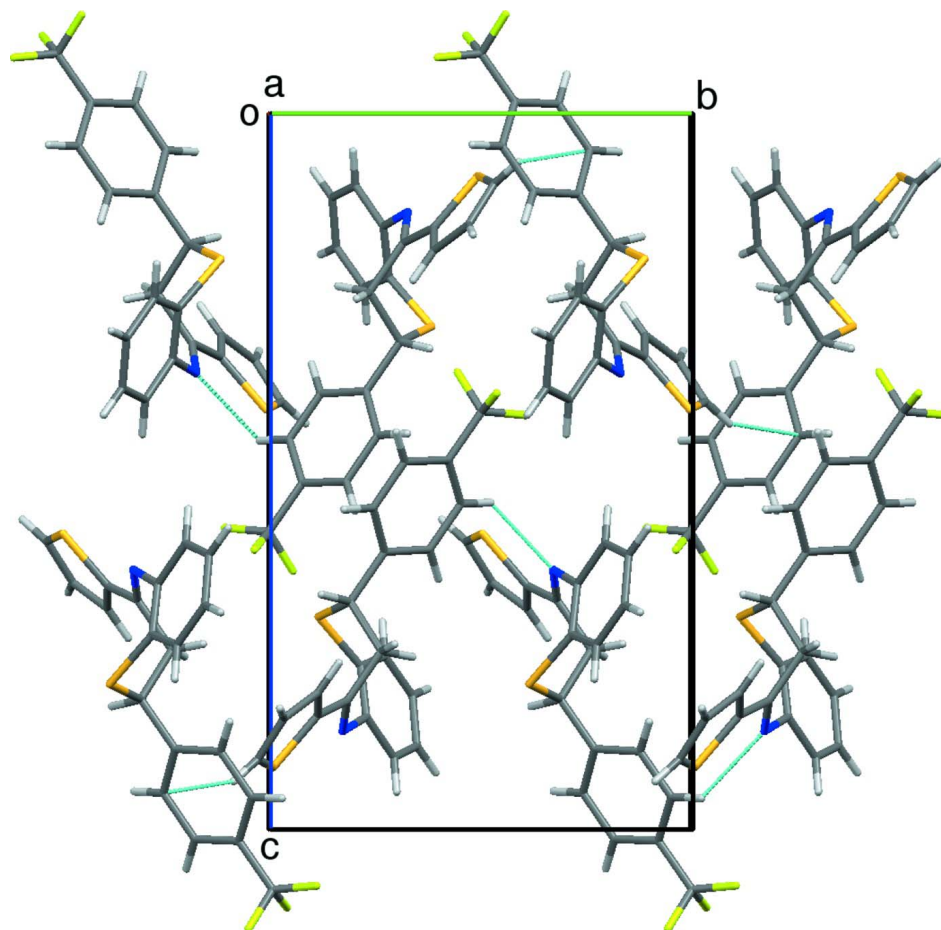
All hydrogen atoms were located geometrically with C—H = 0.93–0.97) Å and allowed to ride on their parent atoms with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{aromatic C})$ .

**Computing details**

Data collection: *APEX2* (Bruker, 2013); cell refinement: *SAINTE* (Bruker, 2013); data reduction: *SAINTE* (Bruker, 2013); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *Mercury* (Macrae *et al.*, 2006).

**Figure 1**

ORTEP diagram of the title molecule with 50% probability ellipsoids.

**Figure 2**

Packing diagram of molecule, viewed along the  $a$  axis. Dotted lines represent weak C—H...N intermolecular interactions.

#### 4-(Thiophen-2-yl)-2-[4-(trifluoromethyl)phenyl]-2,3-dihydro-1,5-benzothiazepine

##### Crystal data

$C_{20}H_{14}F_3NS_2$

$M_r = 389.46$

Orthorhombic,  $P2_12_12_1$

Hall symbol:  $P\ 2ac\ 2ab$

$a = 9.847\ (2)\ \text{\AA}$

$b = 10.492\ (3)\ \text{\AA}$

$c = 17.819\ (4)\ \text{\AA}$

$V = 1841.0\ (8)\ \text{\AA}^3$

$Z = 4$

$F(000) = 800$

$D_x = 1.405\ \text{Mg m}^{-3}$

Cu  $K\alpha$  radiation,  $\lambda = 1.54178\ \text{\AA}$

Cell parameters from 12103 reflections

$\theta = 4.9\text{--}64.8^\circ$

$\mu = 2.92\ \text{mm}^{-1}$

$T = 296\ \text{K}$

Needle, light yellow

$0.20 \times 0.19 \times 0.18\ \text{mm}$

##### Data collection

Bruker X8 Proteum

diffractometer

Radiation source: Bruker MicroStar microfocus

rotating anode

Helios multilayer optics monochromator

Detector resolution:  $10.7\ \text{pixels mm}^{-1}$

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2013)

$T_{\min} = 0.593$ ,  $T_{\max} = 0.622$

12103 measured reflections

3028 independent reflections

2547 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.056$

$\theta_{\max} = 64.8^\circ$ ,  $\theta_{\min} = 4.9^\circ$   
 $h = -11 \rightarrow 11$

$k = -8 \rightarrow 12$   
 $l = -20 \rightarrow 20$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.085$   
 $wR(F^2) = 0.239$   
 $S = 1.03$   
 3028 reflections  
 237 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  
 $w = 1/[\sigma^2(F_o^2) + (0.1979P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.94 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.51 \text{ e } \text{\AA}^{-3}$   
 Extinction correction: *SHELXL*,  
 $FC^* = KFC[1 + 0.001XFC^2\Lambda^3/\text{SIN}(2\Theta)]^{-1/4}$   
 Extinction coefficient: 0.017 (3)  
 Absolute structure: Flack (1983), 1270 Friedel  
 pairs  
 Absolute structure parameter: 0.0 (3)

*Special details*

**Experimental.** m.p. 405 K, <sup>1</sup>H NMR (CDCl<sub>3</sub>):  $\delta$  2.0 (d, 2H, C3—H), 3.80 (t, 1H, C2—H), 7.24 (dd, 2H, Ar—H), 7.58 (dd, 2H, Ar—H), 7.18 (t, 1H, C4—H 5 m ring), 7.46 (d, 1H, C3—H 5 m ring), 7.72 (d, 1H, C5—H 5 m ring), 7.32–7.46 (m, 4H, Ar—H). Anal. Calcd. for C<sub>20</sub>H<sub>14</sub>F<sub>3</sub>NS<sub>2</sub>: C 62.14, H 3.56, N 3.58%; Found C 61.68, H 3.62, N 3.60%. Mass FAB+ (NBA): 390 ( $M + 1$ , 100%).

**Geometry.** Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

**Refinement.** Refinement on  $F^2$  for ALL reflections except those flagged by the user for potential systematic errors. Weighted  $R$ -factors  $wR$  and all goodnesses 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 observed criterion of  $F^2 > \sigma(F^2)$  is used only for calculating  $-R$ -factor-obs *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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.3240 (2)	0.99443 (19)	0.41340 (10)	0.0841 (7)
S9	-0.00510 (15)	0.88544 (12)	0.20297 (9)	0.0608 (5)
F24	0.2469 (9)	0.4640 (10)	-0.1093 (6)	0.257 (6)
F25	0.0847 (19)	0.5559 (8)	-0.1388 (4)	0.289 (10)
F26	0.0674 (7)	0.3951 (6)	-0.0787 (3)	0.139 (3)
N12	0.1053 (5)	0.8288 (4)	0.3632 (3)	0.0550 (16)
C2	0.4905 (8)	1.0319 (9)	0.4023 (6)	0.104 (4)
C3	0.5488 (8)	0.9699 (9)	0.3449 (7)	0.101 (4)
C4	0.4592 (6)	0.8887 (6)	0.3089 (4)	0.0650 (19)
C5	0.3308 (5)	0.8882 (5)	0.3403 (3)	0.0533 (17)
C6	0.2105 (5)	0.8181 (5)	0.3202 (3)	0.0500 (16)
C7	0.2105 (5)	0.7370 (5)	0.2509 (3)	0.0477 (16)
C8	0.1595 (5)	0.8108 (4)	0.1818 (3)	0.0510 (16)
C10	-0.0758 (6)	0.7848 (5)	0.2719 (3)	0.0523 (16)
C11	-0.0168 (5)	0.7715 (5)	0.3428 (3)	0.0517 (14)
C13	-0.2004 (6)	0.7269 (6)	0.2580 (4)	0.0680 (19)
C14	-0.2705 (6)	0.6610 (6)	0.3132 (5)	0.071 (2)

C15	-0.2116 (7)	0.6511 (6)	0.3834 (5)	0.073 (2)
C16	-0.0877 (7)	0.7058 (6)	0.3996 (4)	0.0653 (19)
C17	0.1511 (5)	0.7293 (4)	0.1124 (3)	0.0447 (12)
C18	0.2330 (6)	0.7568 (5)	0.0509 (3)	0.0503 (16)
C19	0.2279 (6)	0.6822 (5)	-0.0137 (3)	0.0557 (17)
C20	0.1419 (5)	0.5784 (5)	-0.0162 (3)	0.0493 (16)
C21	0.0616 (6)	0.5477 (5)	0.0452 (3)	0.0580 (17)
C22	0.0667 (6)	0.6223 (5)	0.1088 (3)	0.0580 (17)
C23	0.1334 (7)	0.4983 (7)	-0.0842 (3)	0.071 (2)
H2	0.53640	1.08950	0.43280	0.1250*
H3	0.63920	0.98020	0.33100	0.1210*
H4	0.48310	0.83940	0.26760	0.0780*
H7A	0.15280	0.66340	0.25910	0.0570*
H7B	0.30190	0.70660	0.24150	0.0570*
H8	0.22470	0.87940	0.17180	0.0610*
H13	-0.23780	0.73260	0.21020	0.0820*
H14	-0.35470	0.62440	0.30330	0.0850*
H15	-0.25690	0.60620	0.42080	0.0880*
H16	-0.05120	0.69950	0.44760	0.0790*
H18	0.29200	0.82590	0.05300	0.0600*
H19	0.28190	0.70230	-0.05490	0.0660*
H21	0.00450	0.47720	0.04340	0.0700*
H22	0.01320	0.60120	0.14990	0.0700*

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
S1	0.0843 (11)	0.0942 (13)	0.0738 (11)	-0.0174 (9)	-0.0048 (8)	-0.0297 (9)
S9	0.0687 (9)	0.0503 (7)	0.0634 (9)	0.0168 (6)	0.0018 (7)	0.0040 (5)
F24	0.190 (8)	0.307 (12)	0.275 (12)	-0.134 (8)	0.144 (8)	-0.238 (11)
F25	0.64 (3)	0.152 (6)	0.076 (4)	0.034 (12)	-0.127 (9)	-0.021 (4)
F26	0.200 (7)	0.123 (4)	0.093 (4)	-0.086 (4)	0.037 (4)	-0.050 (3)
N12	0.063 (3)	0.050 (2)	0.052 (3)	-0.002 (2)	0.010 (2)	-0.0088 (18)
C2	0.073 (4)	0.106 (6)	0.133 (8)	-0.017 (4)	-0.026 (5)	-0.019 (6)
C3	0.067 (4)	0.104 (6)	0.132 (8)	-0.018 (4)	0.003 (5)	-0.001 (6)
C4	0.051 (3)	0.070 (3)	0.074 (4)	-0.003 (2)	0.011 (3)	0.004 (3)
C5	0.053 (3)	0.050 (3)	0.057 (3)	0.004 (2)	-0.001 (2)	0.003 (2)
C6	0.054 (3)	0.046 (2)	0.050 (3)	0.008 (2)	0.005 (2)	0.002 (2)
C7	0.055 (3)	0.042 (2)	0.046 (3)	0.004 (2)	0.004 (2)	-0.005 (2)
C8	0.064 (3)	0.045 (2)	0.044 (3)	0.004 (2)	0.001 (2)	-0.0035 (19)
C10	0.057 (3)	0.040 (2)	0.060 (3)	0.008 (2)	0.000 (2)	-0.005 (2)
C11	0.050 (2)	0.050 (2)	0.055 (3)	-0.002 (2)	0.012 (2)	-0.010 (2)
C13	0.049 (3)	0.067 (3)	0.088 (4)	0.005 (3)	-0.008 (3)	-0.008 (3)
C14	0.054 (3)	0.064 (3)	0.095 (5)	-0.010 (3)	0.002 (3)	-0.002 (3)
C15	0.063 (3)	0.067 (4)	0.089 (5)	-0.005 (3)	0.017 (4)	0.003 (3)
C16	0.075 (4)	0.064 (3)	0.057 (3)	-0.007 (3)	0.018 (3)	-0.001 (3)
C17	0.052 (2)	0.039 (2)	0.043 (2)	-0.0020 (19)	0.0043 (19)	0.0005 (19)
C18	0.057 (3)	0.045 (2)	0.049 (3)	-0.007 (2)	0.004 (2)	0.005 (2)
C19	0.061 (3)	0.057 (3)	0.049 (3)	-0.014 (2)	0.008 (2)	0.000 (2)
C20	0.053 (3)	0.049 (2)	0.046 (3)	-0.004 (2)	0.005 (2)	0.000 (2)

C21	0.066 (3)	0.055 (3)	0.053 (3)	-0.009 (2)	0.010 (3)	-0.003 (2)
C22	0.067 (3)	0.053 (3)	0.054 (3)	-0.016 (2)	0.014 (2)	-0.005 (2)
C23	0.087 (4)	0.080 (4)	0.047 (3)	-0.026 (4)	0.018 (3)	-0.012 (3)

*Geometric parameters (Å, °)*

S1—C2	1.698 (8)	C17—C18	1.391 (8)
S1—C5	1.716 (6)	C17—C22	1.398 (7)
S9—C8	1.839 (5)	C18—C19	1.393 (8)
S9—C10	1.763 (6)	C19—C20	1.380 (8)
F24—C23	1.256 (11)	C20—C21	1.388 (8)
F25—C23	1.242 (12)	C20—C23	1.477 (8)
F26—C23	1.267 (10)	C21—C22	1.378 (8)
N12—C6	1.293 (7)	C2—H2	0.9300
N12—C11	1.393 (7)	C3—H3	0.9300
C2—C3	1.341 (15)	C4—H4	0.9300
C3—C4	1.384 (12)	C7—H7A	0.9700
C4—C5	1.383 (8)	C7—H7B	0.9700
C5—C6	1.440 (7)	C8—H8	0.9800
C6—C7	1.500 (8)	C13—H13	0.9300
C7—C8	1.539 (7)	C14—H14	0.9300
C8—C17	1.506 (7)	C15—H15	0.9300
C10—C11	1.398 (8)	C16—H16	0.9300
C10—C13	1.391 (8)	C18—H18	0.9300
C11—C16	1.410 (9)	C19—H19	0.9300
C13—C14	1.386 (10)	C21—H21	0.9300
C14—C15	1.383 (12)	C22—H22	0.9300
C15—C16	1.379 (10)		
C2—S1—C5	91.4 (4)	F24—C23—F25	101.8 (10)
C8—S9—C10	103.6 (2)	F24—C23—F26	103.8 (8)
C6—N12—C11	120.0 (5)	F24—C23—C20	113.9 (7)
S1—C2—C3	113.0 (7)	F25—C23—F26	106.2 (8)
C2—C3—C4	112.3 (7)	F25—C23—C20	112.8 (7)
C3—C4—C5	113.4 (7)	F26—C23—C20	116.9 (5)
S1—C5—C4	109.9 (4)	S1—C2—H2	123.00
S1—C5—C6	119.3 (4)	C3—C2—H2	124.00
C4—C5—C6	130.8 (5)	C2—C3—H3	124.00
N12—C6—C5	117.9 (5)	C4—C3—H3	124.00
N12—C6—C7	122.5 (5)	C3—C4—H4	123.00
C5—C6—C7	119.7 (4)	C5—C4—H4	123.00
C6—C7—C8	111.9 (4)	C6—C7—H7A	109.00
S9—C8—C7	109.7 (4)	C6—C7—H7B	109.00
S9—C8—C17	111.2 (3)	C8—C7—H7A	109.00
C7—C8—C17	112.9 (4)	C8—C7—H7B	109.00
S9—C10—C11	121.7 (4)	H7A—C7—H7B	108.00
S9—C10—C13	119.1 (5)	S9—C8—H8	108.00
C11—C10—C13	118.9 (5)	C7—C8—H8	108.00
N12—C11—C10	123.5 (5)	C17—C8—H8	108.00
N12—C11—C16	116.8 (5)	C10—C13—H13	119.00

C10—C11—C16	119.5 (5)	C14—C13—H13	119.00
C10—C13—C14	122.0 (6)	C13—C14—H14	121.00
C13—C14—C15	118.1 (6)	C15—C14—H14	121.00
C14—C15—C16	122.0 (7)	C14—C15—H15	119.00
C11—C16—C15	119.4 (7)	C16—C15—H15	119.00
C8—C17—C18	119.8 (4)	C11—C16—H16	120.00
C8—C17—C22	121.8 (5)	C15—C16—H16	120.00
C18—C17—C22	118.4 (5)	C17—C18—H18	119.00
C17—C18—C19	120.9 (5)	C19—C18—H18	120.00
C18—C19—C20	119.5 (5)	C18—C19—H19	120.00
C19—C20—C21	120.5 (5)	C20—C19—H19	120.00
C19—C20—C23	120.7 (5)	C20—C21—H21	120.00
C21—C20—C23	118.8 (5)	C22—C21—H21	120.00
C20—C21—C22	119.7 (5)	C17—C22—H22	119.00
C17—C22—C21	121.0 (5)	C21—C22—H22	119.00
C5—S1—C2—C3	1.5 (8)	S9—C10—C11—C16	170.3 (4)
C2—S1—C5—C4	-1.9 (5)	C13—C10—C11—N12	-177.9 (5)
C2—S1—C5—C6	179.5 (5)	S9—C10—C13—C14	-171.0 (5)
C10—S9—C8—C7	-27.7 (4)	C11—C10—C13—C14	2.8 (9)
C10—S9—C8—C17	97.9 (4)	C13—C10—C11—C16	-3.3 (8)
C8—S9—C10—C11	65.2 (5)	S9—C10—C11—N12	-4.3 (8)
C8—S9—C10—C13	-121.2 (5)	N12—C11—C16—C15	177.7 (5)
C6—N12—C11—C16	135.8 (6)	C10—C11—C16—C15	2.7 (9)
C11—N12—C6—C5	174.6 (5)	C10—C13—C14—C15	-1.5 (10)
C11—N12—C6—C7	-4.9 (8)	C13—C14—C15—C16	0.8 (10)
C6—N12—C11—C10	-49.5 (7)	C14—C15—C16—C11	-1.5 (10)
S1—C2—C3—C4	-0.7 (11)	C8—C17—C18—C19	179.5 (5)
C2—C3—C4—C5	-0.7 (11)	C22—C17—C18—C19	2.1 (8)
C3—C4—C5—S1	1.8 (8)	C8—C17—C22—C21	-179.1 (5)
C3—C4—C5—C6	-179.8 (7)	C18—C17—C22—C21	-1.8 (8)
S1—C5—C6—N12	-6.9 (7)	C17—C18—C19—C20	-1.0 (8)
C4—C5—C6—N12	174.9 (6)	C18—C19—C20—C21	-0.4 (8)
C4—C5—C6—C7	-5.6 (9)	C18—C19—C20—C23	179.4 (5)
S1—C5—C6—C7	172.7 (4)	C19—C20—C21—C22	0.7 (8)
N12—C6—C7—C8	87.7 (6)	C23—C20—C21—C22	-179.1 (5)
C5—C6—C7—C8	-91.8 (6)	C19—C20—C23—F24	49.1 (9)
C6—C7—C8—C17	-176.8 (4)	C19—C20—C23—F25	-66.2 (11)
C6—C7—C8—S9	-52.1 (5)	C19—C20—C23—F26	170.3 (6)
C7—C8—C17—C18	-116.1 (5)	C21—C20—C23—F24	-131.0 (8)
S9—C8—C17—C18	120.0 (5)	C21—C20—C23—F25	113.6 (11)
S9—C8—C17—C22	-62.8 (5)	C21—C20—C23—F26	-9.9 (9)
C7—C8—C17—C22	61.1 (6)	C20—C21—C22—C17	0.4 (8)

Hydrogen-bond geometry (Å, °)

<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>
C21—H21...N12 <sup>i</sup>	0.93	2.52	3.262 (7)	137



Symmetry code: (i)  $-x, y-1/2, -z+1/2$ .