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## N-(4-Chlorobutanoyl)-N'-[2-(trifluoromethyl)phenyl]thiourea

Mohd Sukeri Mohd Yusof,<sup>a</sup> Nur Farhana Embong,<sup>a</sup> Suhana Arshad<sup>b</sup> and Ibrahim Abdul Razak<sup>b\*†</sup><sup>a</sup>Department of Chemical Sciences, Faculty of Science and Technology, Universiti Malaysia Terengganu, Mengabang Telipot, 21030 Kuala Terengganu, Malaysia, and <sup>b</sup>School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia

Correspondence e-mail: arazaki@usm.my

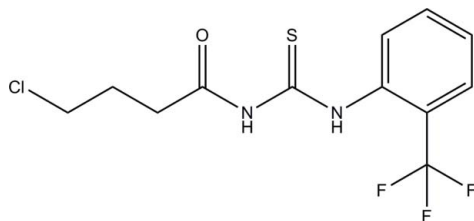
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Key indicators: single-crystal X-ray study;  $T = 100$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.030;  $wR$  factor = 0.078; data-to-parameter ratio = 25.8.

In the title compound,  $\text{C}_{12}\text{H}_{12}\text{ClF}_3\text{N}_2\text{OS}$ , the dihedral angle between the benzene ring and the thiourea fragment is  $69.41(5)^\circ$ . The thiourea N—H atoms adopt an *anti* conformation, such that one of them forms an intramolecular N—H $\cdots$ O hydrogen bond, generating an  $S(6)$  ring. In the crystal, both N—H groups form inversion dimers, one *via* a pair of N—H $\cdots$ S hydrogen bonds and one *via* a pair of N—H $\cdots$ O hydrogen bonds. These lead to  $R_2^2(8)$  and  $R_2^2(12)$  loops, respectively. Weak C—H $\cdots$ Cl, C—H $\cdots$ F, C—H $\cdots$ S and  $\pi$ — $\pi$  [centroid—centroid separation =  $3.7098(6)$  Å and slippage =  $1.853$  Å] interactions also occur.

## Related literature

For a related structure and background to thiourea derivatives, see: Yusof *et al.* (2011). For related structures, see: Khawar Rauf *et al.* (2006); Yusof *et al.* (2007). For hydrogen-bond motifs, see: Bernstein *et al.* (1995). For the stability of the temperature controller used in the data collection, see: Cosier & Glazer (1986).



## Experimental

## Crystal data

 $\text{C}_{12}\text{H}_{12}\text{ClF}_3\text{N}_2\text{OS}$   
 $M_r = 324.75$ Triclinic,  $P\bar{1}$   
 $a = 7.8622(1)$  Å

† Thomson Reuters ResearcherID: A-5599-2009

 $b = 8.9073(1)$  Å  
 $c = 11.0341(1)$  Å  
 $\alpha = 113.687(1)^\circ$   
 $\beta = 103.419(1)^\circ$   
 $\gamma = 95.653(1)^\circ$   
 $V = 672.18(2)$  Å<sup>3</sup> $Z = 2$   
Mo  $K\alpha$  radiation  
 $\mu = 0.47$  mm<sup>-1</sup>  
 $T = 100$  K  
 $0.41 \times 0.19 \times 0.15$  mm

## Data collection

Bruker SMART APEXII CCD diffractometer  
Absorption correction: multi-scan (SADABS; Bruker, 2009)  
 $T_{\min} = 0.829$ ,  $T_{\max} = 0.932$ 18148 measured reflections  
4884 independent reflections  
4304 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.021$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.030$   
 $wR(F^2) = 0.078$   
 $S = 0.97$   
4884 reflections  
189 parametersH atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.49$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.32$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

D—H $\cdots$ A	D—H	H $\cdots$ A	D $\cdots$ A	D—H $\cdots$ A
N1—H1N1 $\cdots$ O1	0.855 (16)	1.971 (17)	2.6486 (12)	135.4 (16)
N1—H1N1 $\cdots$ O1 <sup>i</sup>	0.855 (16)	2.514 (17)	3.2273 (12)	141.6 (14)
N2—H1N2 $\cdots$ S1 <sup>ii</sup>	0.849 (17)	2.682 (17)	3.5079 (10)	164.7 (14)
C2—H2A $\cdots$ Cl1 <sup>iii</sup>	0.95	2.82	3.5535 (11)	135
C3—H3A $\cdots$ F1 <sup>iv</sup>	0.95	2.47	3.2617 (13)	140
C9—H9A $\cdots$ S1 <sup>ii</sup>	0.99	2.84	3.7829 (10)	159

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

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6655).

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## supplementary materials

*Acta Cryst.* (2012). E68, o1029 [doi:10.1107/S1600536812008859]

***N*-(4-Chlorobutanoyl)-*N'*-[2-(trifluoromethyl)phenyl]thiourea**

Mohd Sukeri Mohd Yusof, Nur Farhana Embong, Suhana Arshad and Ibrahim Abdul Razak

**Comment**

As part of our ongoing studies of thiourea derivatives we now describe the title compound. It is analogous to the previously reported *N*-(4-chlorobutanoyl)-*N'*-(2-fluorophenyl)thiourea (Yusof *et al.*, 2011) except the fluoro atom is replaced by trifluoromethyl atom.

In the molecular structure (Fig. 1), the benzene ring (C1–C6) is essentially planar with maximum deviation of 0.011 (1) Å at atom C5. The intramolecular N1—H1N1···O1 hydrogen bond (Table 1) generates *S*(6) ring motifs (Berstein *et al.*, 1995). The bond lengths and angles are within normal ranges and are comparable to the related structures (Khawar Rauf *et al.*, 2006; Yusof *et al.*, 2007).

The crystal packing is shown in Fig. 2.  $R^1_2(6)$ ,  $R^2_2(8)$ ,  $R^2_2(12)$  ring motifs (Berstein *et al.* 1995) are formed by intermolecular N2—H1N2···S1, N1—H1N1···O1 and C9—H9A···S1 (Table 1) hydrogen bonds, respectively. Intermolecular C2—H2A···Cl1 and C3—H3A···F1 (Table 1) interactions linked the molecules into three-dimensional network.  $\pi$ – $\pi$  interaction [ $Cg1 \cdots Cg1$  ( $-1 - x, 1 - y, 1 - z$ ) = 3.7098 (6) Å;] are also observed [ $Cg1$ : C1–C6].

**Experimental**

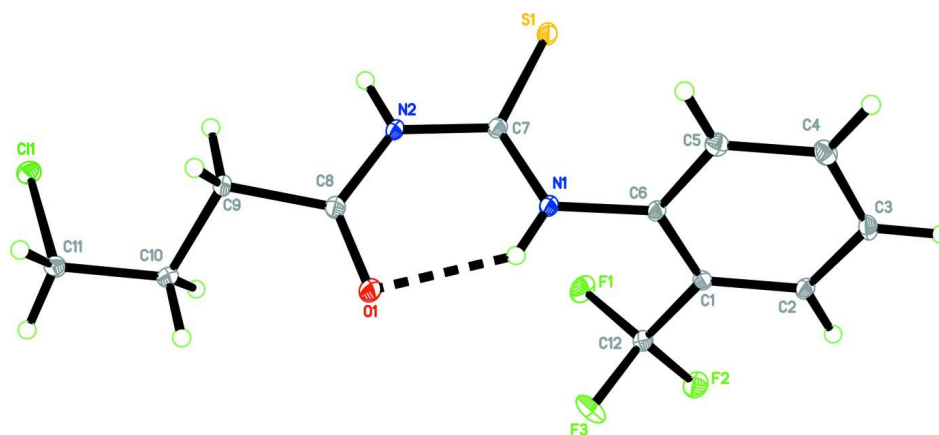
An equimolar amount of 2-(trifluoromethyl)aniline (1.14 g, 7.09 mmol) in 20 ml acetone was added drop-wise into a stirring acetone solution (75 ml) containing 4-chlorobutanoylchloride (1.00 g, 7.09 mmol) and ammonium thiocyanate (0.54 g, 7.09 mmol). The mixture was refluxed for 1 h. Then, the solution was filtered-off and left to evaporate at room temperature to yield colourless needles.

**Refinement**

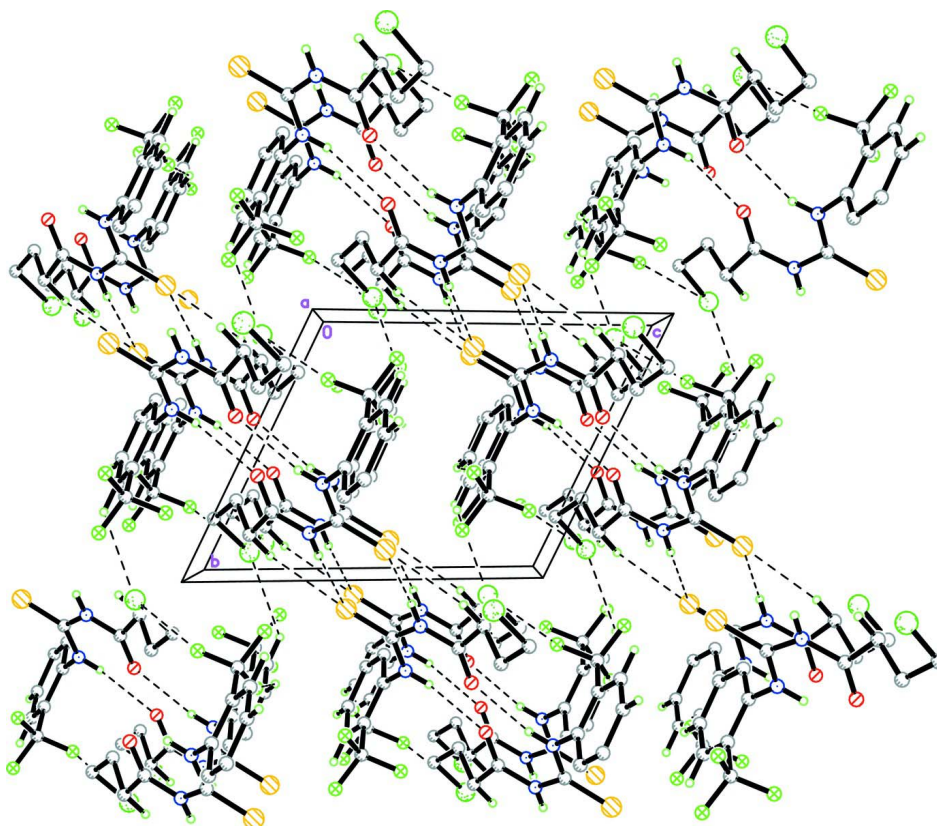
N-bound H atoms was located from the difference map and refined freely, [N–H = 0.856 (17) and 0.849 (15) Å]. The remaining H atoms were positioned geometrically [C–H = 0.95 or 0.99 Å] and refined using a riding model with  $U_{iso}(H) = 1.2 U_{eq}(C)$ .

**Computing details**

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINTE* (Bruker, 2009); data reduction: *SAINTE* (Bruker, 2009); program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008) and *PLATON* (Spek, 2009).

**Figure 1**

The molecular structure of the title compound with 30% probability displacement ellipsoids.

**Figure 2**

The crystal packing of the title compound. The H atoms not involved in the intermolecular interactions (dashed lines) have been omitted for clarity.

### *N*-(4-Chlorobutanoyl)-*N'*-[2-(trifluoromethyl)phenyl]thiourea

#### *Crystal data*

$C_{12}H_{12}ClF_3N_2OS$   
 $M_r = 324.75$

Triclinic,  $P\bar{1}$   
Hall symbol: -P 1

$a = 7.8622$  (1) Å  
 $b = 8.9073$  (1) Å  
 $c = 11.0341$  (1) Å  
 $\alpha = 113.687$  (1)°  
 $\beta = 103.419$  (1)°  
 $\gamma = 95.653$  (1)°  
 $V = 672.18$  (2) Å<sup>3</sup>  
 $Z = 2$   
 $F(000) = 332$

$D_x = 1.605$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å  
 Cell parameters from 9937 reflections  
 $\theta = 2.5$ – $32.6$ °  
 $\mu = 0.47$  mm<sup>-1</sup>  
 $T = 100$  K  
 Needle, colourless  
 $0.41 \times 0.19 \times 0.15$  mm

*Data collection*

Bruker SMART APEXII CCD  
 diffractometer  
 Radiation source: fine-focus sealed tube  
 Graphite monochromator  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan  
 (SADABS; Bruker, 2009)  
 $T_{\min} = 0.829$ ,  $T_{\max} = 0.932$

18148 measured reflections  
 4884 independent reflections  
 4304 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.021$   
 $\theta_{\max} = 32.6$ °,  $\theta_{\min} = 2.1$ °  
 $h = -11 \rightarrow 11$   
 $k = -13 \rightarrow 12$   
 $l = -15 \rightarrow 16$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.030$   
 $wR(F^2) = 0.078$   
 $S = 0.97$   
 4884 reflections  
 189 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 atoms treated by a mixture of independent  
 and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0365P)^2 + 0.3295P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.49$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.32$  e Å<sup>-3</sup>

*Special details*

**Experimental.** The crystal was placed in the cold stream of an Oxford Cryosystems Cobra open-flow nitrogen cryostat (Cosier & Glazer, 1986) operating at 100.0 (1) K.

**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 (Å<sup>2</sup>)*

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	1.66943 (3)	0.08190 (3)	0.87826 (3)	0.01972 (6)
S1	0.77778 (4)	0.12238 (3)	0.48626 (3)	0.01851 (6)
F1	0.97525 (9)	0.59829 (9)	0.65758 (8)	0.02506 (15)
F2	0.86578 (9)	0.81937 (8)	0.70633 (8)	0.02360 (14)
F3	0.97036 (10)	0.74396 (10)	0.86623 (7)	0.02883 (16)
O1	1.11812 (11)	0.39045 (10)	0.93786 (8)	0.02277 (17)

N1	0.83210 (12)	0.36357 (10)	0.74177 (9)	0.01434 (15)
N2	1.02233 (11)	0.17737 (10)	0.71774 (9)	0.01377 (15)
C1	0.69392 (13)	0.58116 (12)	0.69974 (9)	0.01248 (16)
C2	0.54109 (13)	0.64135 (12)	0.66718 (10)	0.01409 (16)
H2A	0.5523	0.7491	0.6683	0.017*
C3	0.37240 (13)	0.54429 (13)	0.63303 (10)	0.01548 (17)
H3A	0.2685	0.5864	0.6123	0.019*
C4	0.35583 (14)	0.38496 (13)	0.62916 (10)	0.01655 (18)
H4A	0.2404	0.3175	0.6038	0.020*
C5	0.50789 (14)	0.32467 (12)	0.66239 (10)	0.01537 (17)
H5A	0.4962	0.2162	0.6598	0.018*
C6	0.67708 (13)	0.42288 (12)	0.69935 (9)	0.01262 (16)
C7	0.87834 (13)	0.22901 (12)	0.65719 (10)	0.01306 (16)
C8	1.13277 (14)	0.25600 (12)	0.85287 (10)	0.01538 (17)
C9	1.27517 (14)	0.16553 (12)	0.88836 (10)	0.01620 (17)
H9A	1.2920	0.0827	0.8021	0.019*
H9B	1.2370	0.1043	0.9385	0.019*
C10	1.45172 (14)	0.29236 (12)	0.97903 (10)	0.01529 (17)
H10A	1.4875	0.3522	0.9272	0.018*
H10B	1.4310	0.3765	1.0628	0.018*
C11	1.60482 (14)	0.21825 (13)	1.02413 (10)	0.01720 (18)
H11A	1.5689	0.1538	1.0726	0.021*
H11B	1.7087	0.3100	1.0902	0.021*
C12	0.87515 (13)	0.68537 (12)	0.73243 (11)	0.01615 (17)
H1N2	1.050 (2)	0.0918 (19)	0.6625 (16)	0.021 (4)*
H1N1	0.895 (2)	0.413 (2)	0.8270 (17)	0.026 (4)*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C11	0.01900 (12)	0.01807 (11)	0.01890 (11)	0.00744 (9)	0.00681 (9)	0.00360 (9)
S1	0.01973 (12)	0.01899 (12)	0.01193 (11)	0.00983 (9)	0.00168 (9)	0.00240 (9)
F1	0.0166 (3)	0.0231 (3)	0.0371 (4)	0.0074 (3)	0.0159 (3)	0.0098 (3)
F2	0.0204 (3)	0.0169 (3)	0.0348 (4)	0.0028 (2)	0.0088 (3)	0.0125 (3)
F3	0.0184 (3)	0.0350 (4)	0.0206 (3)	-0.0043 (3)	-0.0044 (3)	0.0078 (3)
O1	0.0251 (4)	0.0206 (4)	0.0152 (3)	0.0129 (3)	0.0010 (3)	0.0015 (3)
N1	0.0152 (4)	0.0144 (3)	0.0114 (3)	0.0069 (3)	0.0024 (3)	0.0038 (3)
N2	0.0141 (4)	0.0126 (3)	0.0131 (3)	0.0064 (3)	0.0030 (3)	0.0039 (3)
C1	0.0115 (4)	0.0135 (4)	0.0110 (4)	0.0036 (3)	0.0033 (3)	0.0038 (3)
C2	0.0140 (4)	0.0149 (4)	0.0128 (4)	0.0056 (3)	0.0042 (3)	0.0048 (3)
C3	0.0117 (4)	0.0216 (4)	0.0130 (4)	0.0064 (3)	0.0040 (3)	0.0066 (3)
C4	0.0130 (4)	0.0209 (4)	0.0150 (4)	0.0023 (3)	0.0051 (3)	0.0070 (3)
C5	0.0157 (4)	0.0157 (4)	0.0147 (4)	0.0034 (3)	0.0051 (3)	0.0064 (3)
C6	0.0126 (4)	0.0138 (4)	0.0105 (4)	0.0053 (3)	0.0033 (3)	0.0039 (3)
C7	0.0138 (4)	0.0128 (4)	0.0129 (4)	0.0048 (3)	0.0043 (3)	0.0053 (3)
C8	0.0155 (4)	0.0154 (4)	0.0138 (4)	0.0060 (3)	0.0030 (3)	0.0052 (3)
C9	0.0157 (4)	0.0146 (4)	0.0155 (4)	0.0065 (3)	0.0011 (3)	0.0049 (3)
C10	0.0171 (4)	0.0127 (4)	0.0138 (4)	0.0051 (3)	0.0037 (3)	0.0037 (3)
C11	0.0163 (4)	0.0174 (4)	0.0140 (4)	0.0061 (3)	0.0033 (3)	0.0031 (3)
C12	0.0127 (4)	0.0155 (4)	0.0175 (4)	0.0032 (3)	0.0037 (3)	0.0051 (3)

Geometric parameters (Å, °)

C11—C11	1.8038 (10)	C2—H2A	0.9500
S1—C7	1.6748 (10)	C3—C4	1.3947 (14)
F1—C12	1.3455 (12)	C3—H3A	0.9500
F2—C12	1.3404 (12)	C4—C5	1.3904 (14)
F3—C12	1.3430 (12)	C4—H4A	0.9500
O1—C8	1.2255 (12)	C5—C6	1.3905 (14)
N1—C7	1.3358 (12)	C5—H5A	0.9500
N1—C6	1.4298 (12)	C8—C9	1.5112 (14)
N1—H1N1	0.856 (17)	C9—C10	1.5305 (14)
N2—C8	1.3813 (13)	C9—H9A	0.9900
N2—C7	1.3921 (12)	C9—H9B	0.9900
N2—H1N2	0.849 (15)	C10—C11	1.5082 (14)
C1—C2	1.3935 (13)	C10—H10A	0.9900
C1—C6	1.4011 (13)	C10—H10B	0.9900
C1—C12	1.5013 (14)	C11—H11A	0.9900
C2—C3	1.3888 (14)	C11—H11B	0.9900
C7—N1—C6	123.49 (8)	O1—C8—N2	122.61 (9)
C7—N1—H1N1	118.1 (11)	O1—C8—C9	122.16 (9)
C6—N1—H1N1	118.3 (11)	N2—C8—C9	115.22 (8)
C8—N2—C7	127.83 (8)	C8—C9—C10	109.74 (8)
C8—N2—H1N2	116.8 (10)	C8—C9—H9A	109.7
C7—N2—H1N2	115.1 (10)	C10—C9—H9A	109.7
C2—C1—C6	119.79 (9)	C8—C9—H9B	109.7
C2—C1—C12	119.61 (9)	C10—C9—H9B	109.7
C6—C1—C12	120.59 (8)	H9A—C9—H9B	108.2
C3—C2—C1	120.22 (9)	C11—C10—C9	115.10 (8)
C3—C2—H2A	119.9	C11—C10—H10A	108.5
C1—C2—H2A	119.9	C9—C10—H10A	108.5
C2—C3—C4	119.88 (9)	C11—C10—H10B	108.5
C2—C3—H3A	120.1	C9—C10—H10B	108.5
C4—C3—H3A	120.1	H10A—C10—H10B	107.5
C5—C4—C3	120.14 (9)	C10—C11—C11	111.48 (7)
C5—C4—H4A	119.9	C10—C11—H11A	109.3
C3—C4—H4A	119.9	C11—C11—H11A	109.3
C4—C5—C6	120.12 (9)	C10—C11—H11B	109.3
C4—C5—H5A	119.9	C11—C11—H11B	109.3
C6—C5—H5A	119.9	H11A—C11—H11B	108.0
C5—C6—C1	119.81 (9)	F2—C12—F3	106.59 (8)
C5—C6—N1	119.42 (8)	F2—C12—F1	106.08 (8)
C1—C6—N1	120.72 (9)	F3—C12—F1	106.43 (8)
N1—C7—N2	116.42 (8)	F2—C12—C1	112.63 (8)
N1—C7—S1	124.76 (7)	F3—C12—C1	112.29 (8)
N2—C7—S1	118.81 (7)	F1—C12—C1	112.35 (8)
C6—C1—C2—C3	-0.66 (14)	C8—N2—C7—N1	-5.61 (15)
C12—C1—C2—C3	178.23 (9)	C8—N2—C7—S1	173.63 (8)
C1—C2—C3—C4	-1.03 (14)	C7—N2—C8—O1	-2.41 (17)

C2—C3—C4—C5	1.43 (14)	C7—N2—C8—C9	178.37 (9)
C3—C4—C5—C6	-0.13 (15)	O1—C8—C9—C10	-40.56 (14)
C4—C5—C6—C1	-1.56 (14)	N2—C8—C9—C10	138.66 (9)
C4—C5—C6—N1	176.03 (9)	C8—C9—C10—C11	178.93 (8)
C2—C1—C6—C5	1.95 (14)	C9—C10—C11—C11	65.16 (10)
C12—C1—C6—C5	-176.93 (9)	C2—C1—C12—F2	-9.09 (13)
C2—C1—C6—N1	-175.61 (8)	C6—C1—C12—F2	169.80 (8)
C12—C1—C6—N1	5.52 (13)	C2—C1—C12—F3	111.26 (10)
C7—N1—C6—C5	66.17 (13)	C6—C1—C12—F3	-69.86 (12)
C7—N1—C6—C1	-116.27 (11)	C2—C1—C12—F1	-128.80 (9)
C6—N1—C7—N2	-173.57 (9)	C6—C1—C12—F1	50.08 (12)
C6—N1—C7—S1	7.24 (14)		

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>
N1—H1N1...O1	0.855 (16)	1.971 (17)	2.6486 (12)	135.4 (16)
N1—H1N1...O1 <sup>i</sup>	0.855 (16)	2.514 (17)	3.2273 (12)	141.6 (14)
N2—H1N2...S1 <sup>ii</sup>	0.849 (17)	2.682 (17)	3.5079 (10)	164.7 (14)
C2—H2A...C11 <sup>iii</sup>	0.95	2.82	3.5535 (11)	135
C3—H3A...F1 <sup>iv</sup>	0.95	2.47	3.2617 (13)	140
C9—H9A...S1 <sup>ii</sup>	0.99	2.84	3.7829 (10)	159

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