

6-(Adamantan-1-yl)-3-(3-fluorophenyl)-1,2,4-triazolo[3,4-*b*][1,3,4]thiadiazole

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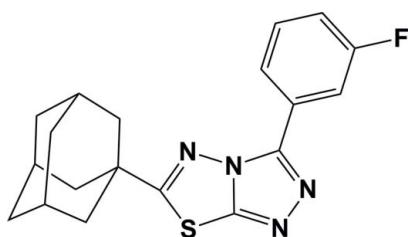
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Key indicators: single-crystal X-ray study; $T = 173\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; disorder in main residue; R factor = 0.030; wR factor = 0.074; data-to-parameter ratio = 11.1.

The title molecule, $\text{C}_{19}\text{H}_{19}\text{FN}_4\text{S}$, displays C_s molecular symmetry, in which the crystallographic mirror plane bisects the adamantan-1-yl unit while the 3-fluorophenyl triazole ring is located on the mirror plane. The F atom of the 3-fluorophenyl ring is positionally disordered [occupancy ratio 0.9:0.1]. In the crystal, $\pi-\pi$ interactions between the triazole and phenyl rings occur [centroid–centroid distance = 3.5849 (7) \AA] and weak C–H···F interactions form a ribbon propagating in [010].

Related literature

For the biological significance of fused heterocycles, see: Khan *et al.* (2010a,b); Demirbas *et al.* (2005); Amir *et al.* (2007); Ashok *et al.* (2007); Palekar *et al.* (2009); Serwar *et al.* (2009); Akhtar *et al.* (2007, 2008a,b). For the activity of adamantyl derivatives, see: Kadi *et al.* (2007); Kouatly *et al.* (2009); Zahid *et al.* (2009). For a related structure, see: Khan *et al.* (2009). For standard bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{19}\text{H}_{19}\text{FN}_4\text{S}$

$M_r = 354.44$

Monoclinic, $P2_1/m$
 $a = 11.6385$ (16) \AA

$b = 6.6555$ (5) \AA

$c = 11.6634$ (16) \AA

$\beta = 117.379$ (14) $^\circ$

$V = 802.25$ (17) \AA^3

$Z = 2$
Mo $K\alpha$ radiation
 $\mu = 0.22\text{ mm}^{-1}$

$T = 173\text{ K}$
 $0.34 \times 0.30 \times 0.11\text{ mm}$

Data collection

Stoe IPDS-2 diffractometer
6164 measured reflections
1708 independent reflections

1223 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.031$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.030$
 $wR(F^2) = 0.074$
 $S = 0.95$
1708 reflections
154 parameters
4 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.20\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.26\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
C11—H11A···F1A ⁱ	0.99	2.67	3.526 (3)	145

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

Data collection: *X-AREA* (Stoe & Cie, 2006); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2006); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009) and *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *SHELXL97* (Sheldrick, 2008), *PLATON* and *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: KP2286).

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supporting information

Acta Cryst. (2010). E66, o3215–o3216 [https://doi.org/10.1107/S1600536810046428]

6-(Adamantan-1-yl)-3-(3-fluorophenyl)-1,2,4-triazolo[3,4-*b*][1,3,4]thiadiazole

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S1. Comment

Fused heterocycles have received widespread attention in the recent years due to their outstanding biological applications [Khan *et al.*, 2010a; Demirbas *et al.*, 2005; Amir *et al.*, 2007; Ashok *et al.*, 2007; Palekar *et al.*, 2009]. Moreover, compounds bearing an adamantyl moiety are gaining prominence due to their biological and pharmaceutical potential [Kadi *et al.*, 2007; Kouatly *et al.*, 2009; Zahid *et al.*, 2009]. The title compound was synthesised in continuation of our previous studies on the synthesis and biological screening of five membered heterocycles [Akhtar *et al.*, 2007, 2008a, 2008b; Serwar *et al.*, 2009; Khan *et al.*, 2009, 2010b].

Bond distances (Allen *et al.*, 1987) and bond angles are normal and similar to those reported for the 2-fluorophenyl derivative (Khan *et al.*, 2009). The 3-fluorophenyl moiety is positionally disordered, with atom F1 having occupancies of 0.45/0.05 (F1A/F1B), where F1A is bonded to atom C6, while atom F1B is bonded to atom C8 (Fig. 1).

The title molecule possesses crystallographic mirror symmetry; the mirror plane bisects the adamantane moiety while the rest of the molecule lies in the mirror plane. The 3-fluorophenyl moiety is positionally disordered. The 2-fluorophenyl derivative, however, crystallised in the orthorhombic space group Pbca and the molecule possess no crystallographic symmetry. The 2-fluorophenyl ring is inclined to the triazole ring by 48.6 (1) $^{\circ}$, to elivate steric hindrance.

In the crystal of the title compound there are weak C—H···F interactions [C11—H11A···F1Aⁱ, H···F 2.67 Å, C—H···F 145 $^{\circ}$, symmetry code (i) -*x*, *y* + 1/2, -*z* + 2)] leading to the formation of a one-dimensional ribbon-like structure running in [010] (Fig. 2). There are also π – π interactions with the centroid-to centroid distance between the mean planes of the triazole ring (N2,C2,N3,N4,C3) and symmetry related phenyl rings (C4—C9) being 3.5849 (7) Å [the perpendicular distance is 3.3278 (1) Å with a slippage of 1.333 Å].

S2. Experimental

A mixture of 4-amino-5-(3-fluorophenyl)-2*H*-1,2,4-triazole-3(4*H*)-thione (0.2 g, 0.56 mmol) and adamantan-1-carboxylic acid (0.10 g, 0.56 mmol) was refluxed in the presence of POCl₃ (5.0 mL) for 4 h [Khan *et al.* 2010a]. The reaction mixture was cooled to RT, poured into crushed ice and neutralized using solid potassium carbonate until pH 8. The precipitated solid was filtered off, washed with excess water and recrystallised from chloroform to give colourless plate-like crystals, suitable for X-ray analysis.

S3. Refinement

A region of electron density was located near atom C8 indicating a positional disorder of the 3-fluorophenyl moiety. In the final cycles of least-squares refinement atom F1 was split to give F1a (occupancy 0.45) bonded to atom C6 and atom F1b (occupancy 0.05) bonded to atom C8. Bond length C8—F1b was refined with a distance restraint of 1.36 (2) Å. H-atoms H6a, H8b and H7 were refined with distance restraints of 0.95 (2) Å% with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The remaining H-atoms were included in calculated positions and treated as riding atoms: C—H = 0.95, 0.99, and 1.00 Å for CH-aromatic,

CH_2 and CH -methine H-atoms, respectively, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}$ (parent C-atom).

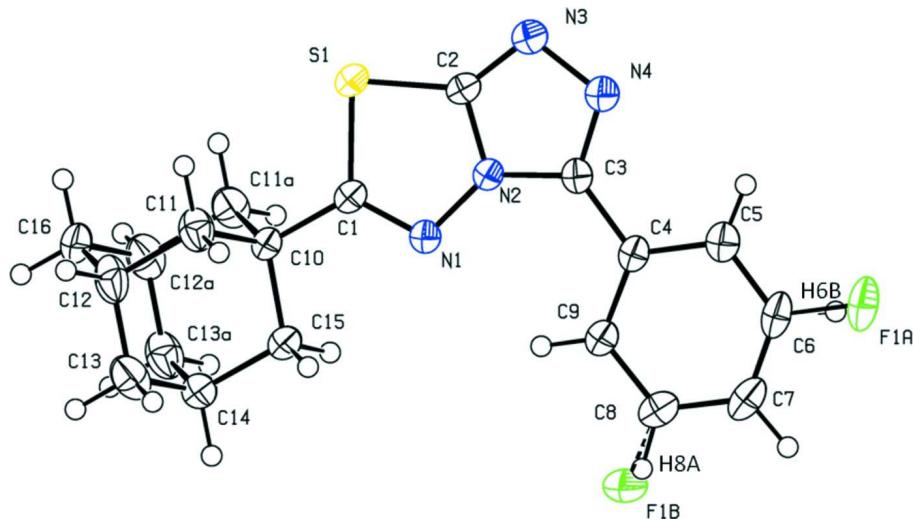
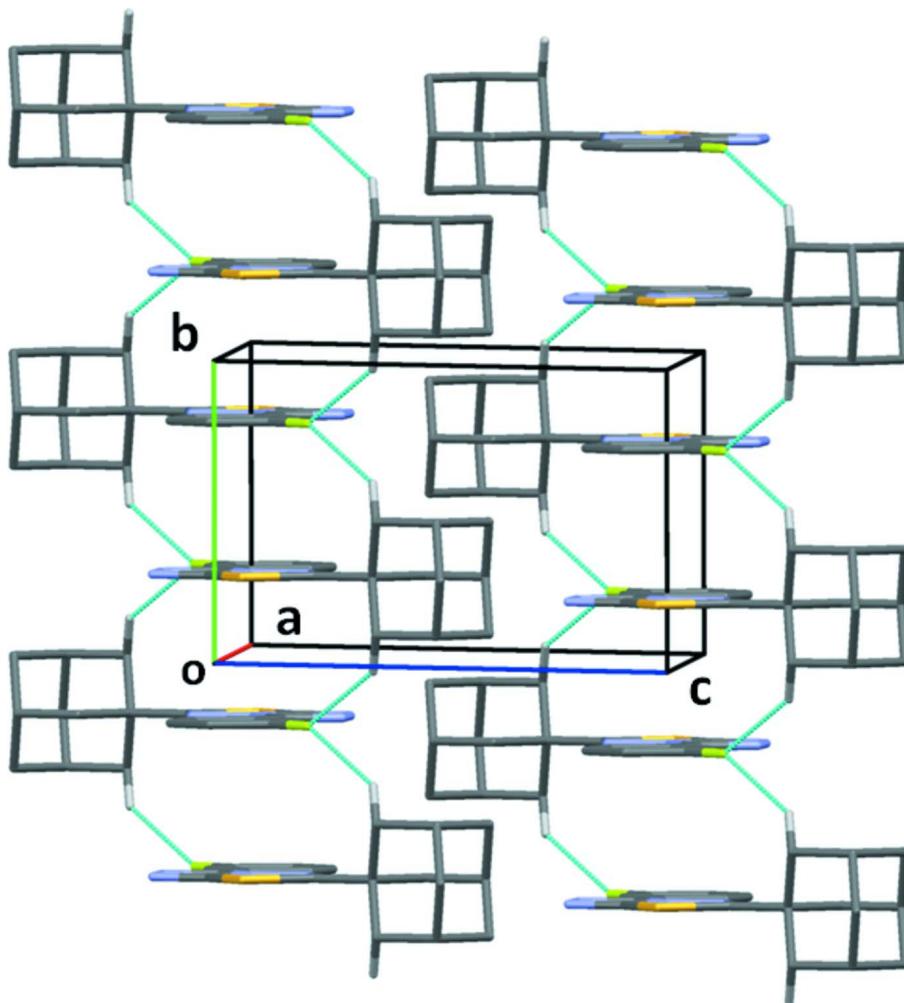


Figure 1

A view of the molecular structure of the title molecule with the displacement ellipsoids drawn at the 50% probability level [symmetry code (*a*) = $x, 3/2 - y, z$; The positional disorder of the 3-fluorophenyl group is also illustrated: atoms F1A and H8A (occupancy 0.45) and atoms F1B and H6B (occupancy 0.05)].

**Figure 2**

The crystal packing of the title compound along the *a* axis. The C—H···F interactions are shown as dashed cyan lines.

6-(Adamantan-1-yl)-3-(3-fluorophenyl)-1,2,4-triazolo[3,4-*b*][1,3,4]thiadiazole

Crystal data

$C_{19}H_{19}FN_4S$

$M_r = 354.44$

Monoclinic, $P2_1/m$

Hall symbol: -P 2yb

$a = 11.6385 (16) \text{ \AA}$

$b = 6.6555 (5) \text{ \AA}$

$c = 11.6634 (16) \text{ \AA}$

$\beta = 117.379 (14)^\circ$

$V = 802.25 (17) \text{ \AA}^3$

$Z = 2$

$F(000) = 372$

$D_x = 1.467 \text{ Mg m}^{-3}$

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

Cell parameters from 4261 reflections

$\theta = 3.1\text{--}26.0^\circ$

$\mu = 0.22 \text{ mm}^{-1}$

$T = 173 \text{ K}$

Plate, colourless

$0.34 \times 0.30 \times 0.11 \text{ mm}$

Data collection

Stoe IPDS-2

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

6164 measured reflections

1708 independent reflections
 1223 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.031$
 $\theta_{\text{max}} = 26.1^\circ, \theta_{\text{min}} = 3.4^\circ$

$h = -14 \rightarrow 14$
 $k = -7 \rightarrow 8$
 $l = -14 \rightarrow 14$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.030$
 $wR(F^2) = 0.074$
 $S = 0.95$
 1708 reflections
 154 parameters
 4 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
 $w = 1/[\sigma^2(F_o^2) + (0.0449P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.20 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.26 \text{ e } \text{\AA}^{-3}$

Special details

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 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	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
S1	0.42509 (5)	0.75000	0.98501 (5)	0.0307 (2)	
F1A	-0.2195 (2)	0.75000	1.1429 (2)	0.0500 (6)	0.900
N1	0.17319 (15)	0.75000	0.85412 (15)	0.0251 (5)	
N2	0.20898 (15)	0.75000	0.98446 (16)	0.0243 (5)	
N3	0.35605 (17)	0.75000	1.18934 (17)	0.0325 (6)	
N4	0.23242 (17)	0.75000	1.18047 (16)	0.0311 (6)	
C1	0.27709 (18)	0.75000	0.84027 (19)	0.0249 (6)	
C2	0.33636 (19)	0.75000	1.06956 (19)	0.0268 (6)	
C3	0.14499 (19)	0.75000	1.05806 (19)	0.0257 (6)	
C4	0.00535 (19)	0.75000	1.0101 (2)	0.0248 (6)	
C5	-0.0419 (2)	0.75000	1.1000 (2)	0.0318 (7)	
C6	-0.1732 (2)	0.75000	1.0556 (2)	0.0342 (7)	
C7	-0.2599 (2)	0.75000	0.9272 (2)	0.0378 (8)	
C8	-0.2117 (2)	0.75000	0.8390 (2)	0.0367 (8)	
C9	-0.08020 (19)	0.75000	0.8790 (2)	0.0299 (7)	
C10	0.27522 (18)	0.75000	0.71113 (18)	0.0254 (6)	
C11	0.34448 (16)	0.9374 (3)	0.69694 (15)	0.0362 (5)	
C12	0.34071 (17)	0.9366 (3)	0.56371 (15)	0.0413 (6)	
C13	0.20148 (17)	0.9360 (3)	0.45854 (15)	0.0428 (6)	
C14	0.1337 (2)	0.75000	0.4720 (2)	0.0382 (8)	
C15	0.1355 (2)	0.75000	0.6040 (2)	0.0368 (7)	

C16	0.4093 (2)	0.75000	0.5501 (2)	0.0441 (9)	
F1B	-0.2859 (18)	0.75000	0.7166 (11)	0.052 (5)	0.100
H7	-0.3504 (11)	0.75000	0.902 (2)	0.0450*	
H8A	-0.276 (4)	0.75000	0.7516 (18)	0.0440*	0.900
H9	-0.04830	0.75000	0.81730	0.0360*	
H11A	0.30140	1.05980	0.70660	0.0440*	
H11B	0.43540	0.93830	0.76530	0.0440*	
H12	0.38550	1.05930	0.55460	0.0490*	
H5	0.01550	0.75000	1.19000	0.0380*	
H13B	0.19930	0.93710	0.37270	0.0510*	
H14	0.04200	0.75000	0.40260	0.0460*	
H15A	0.08990	0.87050	0.61220	0.0440*	0.500
H15B	0.08990	0.62950	0.61220	0.0440*	0.500
H16A	0.40810	0.75000	0.46460	0.0530*	
H16B	0.50060	0.75000	0.61760	0.0530*	
H13A	0.15650	1.05780	0.46590	0.0510*	
H6B	-0.21 (3)	0.75000	1.11 (2)	0.0410*	0.100

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0210 (3)	0.0411 (4)	0.0298 (3)	0.0000	0.0116 (2)	0.0000
F1A	0.0514 (10)	0.0583 (11)	0.0639 (12)	0.0000	0.0467 (9)	0.0000
N1	0.0251 (8)	0.0263 (10)	0.0252 (8)	0.0000	0.0128 (7)	0.0000
N2	0.0218 (8)	0.0275 (9)	0.0245 (8)	0.0000	0.0115 (6)	0.0000
N3	0.0268 (9)	0.0403 (12)	0.0291 (9)	0.0000	0.0118 (7)	0.0000
N4	0.0285 (9)	0.0371 (12)	0.0285 (9)	0.0000	0.0139 (7)	0.0000
C1	0.0219 (9)	0.0230 (11)	0.0301 (10)	0.0000	0.0123 (8)	0.0000
C2	0.0207 (9)	0.0285 (12)	0.0289 (10)	0.0000	0.0095 (8)	0.0000
C3	0.0283 (10)	0.0249 (12)	0.0269 (10)	0.0000	0.0153 (9)	0.0000
C4	0.0260 (10)	0.0206 (11)	0.0320 (10)	0.0000	0.0169 (9)	0.0000
C5	0.0344 (12)	0.0298 (13)	0.0361 (11)	0.0000	0.0205 (9)	0.0000
C6	0.0368 (12)	0.0282 (12)	0.0520 (14)	0.0000	0.0329 (11)	0.0000
C7	0.0270 (11)	0.0337 (14)	0.0581 (15)	0.0000	0.0243 (11)	0.0000
C8	0.0254 (11)	0.0401 (15)	0.0414 (13)	0.0000	0.0126 (10)	0.0000
C9	0.0256 (10)	0.0336 (13)	0.0326 (11)	0.0000	0.0153 (9)	0.0000
C10	0.0217 (10)	0.0288 (12)	0.0276 (10)	0.0000	0.0131 (8)	0.0000
C11	0.0436 (9)	0.0339 (10)	0.0359 (8)	-0.0108 (7)	0.0223 (7)	-0.0049 (7)
C12	0.0535 (10)	0.0393 (10)	0.0405 (9)	-0.0168 (8)	0.0298 (8)	-0.0033 (7)
C13	0.0565 (11)	0.0416 (11)	0.0355 (8)	0.0138 (8)	0.0257 (8)	0.0081 (7)
C14	0.0291 (11)	0.0576 (17)	0.0279 (11)	0.0000	0.0132 (9)	0.0000
C15	0.0232 (10)	0.0586 (17)	0.0290 (10)	0.0000	0.0123 (9)	0.0000
C16	0.0313 (12)	0.073 (2)	0.0357 (12)	0.0000	0.0220 (10)	0.0000
F1B	0.030 (7)	0.084 (12)	0.031 (7)	0.0000	0.006 (7)	0.0000

Geometric parameters (\AA , $\text{^{\circ}}$)

S1—C1	1.773 (2)	C10—C15	1.529 (3)
S1—C2	1.725 (2)	C11—C12	1.534 (2)
F1A—C6	1.353 (3)	C12—C13	1.517 (3)
F1B—C8	1.285 (12)	C12—C16	1.524 (3)
N1—N2	1.379 (2)	C13—C14	1.514 (3)
N1—C1	1.292 (3)	C14—C15	1.530 (3)
N2—C3	1.371 (3)	C5—H5	0.9500
N2—C2	1.353 (3)	C6—H6B	0.9 (3)
N3—N4	1.394 (3)	C7—H7	0.955 (17)
N3—C2	1.308 (3)	C8—H8A	0.95 (2)
N4—C3	1.318 (3)	C9—H9	0.9500
C1—C10	1.496 (3)	C11—H11A	0.9900
C3—C4	1.455 (3)	C11—H11B	0.9900
C4—C9	1.389 (3)	C12—H12	1.0000
C4—C5	1.390 (3)	C13—H13A	0.9900
C5—C6	1.370 (4)	C13—H13B	0.9900
C6—C7	1.368 (3)	C14—H14	1.0000
C7—C8	1.380 (3)	C15—H15A	0.9900
C8—C9	1.380 (4)	C15—H15B	0.9900
C10—C11	1.535 (2)	C16—H16A	0.9900
C10—C11 ⁱ	1.535 (2)	C16—H16B	0.9900
C1—S1—C2	88.25 (11)	C13—C14—C13 ⁱ	109.66 (19)
N2—N1—C1	108.20 (17)	C13 ⁱ —C14—C15	109.73 (12)
N1—N2—C2	118.84 (18)	C10—C15—C14	109.9 (2)
N1—N2—C3	135.60 (19)	C12—C16—C12 ⁱ	109.20 (19)
C2—N2—C3	105.56 (17)	C4—C5—H5	121.00
N4—N3—C2	104.65 (18)	C6—C5—H5	121.00
N3—N4—C3	109.67 (18)	C5—C6—H6B	122.00
S1—C1—N1	115.88 (15)	C7—C6—H6B	115.00
S1—C1—C10	121.10 (17)	C6—C7—H7	119.4 (13)
N1—C1—C10	123.02 (19)	C8—C7—H7	122.7 (13)
S1—C2—N2	108.83 (15)	C7—C8—H8A	114 (3)
S1—C2—N3	138.93 (19)	C9—C8—H8A	125 (3)
N2—C2—N3	112.2 (2)	C4—C9—H9	120.00
N2—C3—N4	107.9 (2)	C8—C9—H9	120.00
N2—C3—C4	126.26 (18)	C10—C11—H11A	110.00
N4—C3—C4	125.9 (2)	C10—C11—H11B	110.00
C3—C4—C5	117.99 (19)	C12—C11—H11A	110.00
C3—C4—C9	122.1 (2)	C12—C11—H11B	110.00
C5—C4—C9	119.9 (2)	H11A—C11—H11B	108.00
C4—C5—C6	118.3 (2)	C11—C12—H12	109.00
F1A—C6—C5	118.5 (2)	C13—C12—H12	109.00
F1A—C6—C7	118.4 (2)	C16—C12—H12	109.00
C5—C6—C7	123.2 (2)	C12—C13—H13A	110.00
C6—C7—C8	117.9 (2)	C12—C13—H13B	110.00

C7—C8—C9	121.1 (2)	C14—C13—H13A	110.00
F1B—C8—C7	122.2 (10)	C14—C13—H13B	110.00
F1B—C8—C9	116.8 (10)	H13A—C13—H13B	108.00
C4—C9—C8	119.6 (2)	C13—C14—H14	109.00
C1—C10—C11	110.08 (11)	C15—C14—H14	109.00
C1—C10—C15	109.90 (19)	C13 ⁱ —C14—H14	109.00
C1—C10—C11 ⁱ	110.08 (11)	C10—C15—H15A	110.00
C11—C10—C15	109.02 (11)	C10—C15—H15B	110.00
C11—C10—C11 ⁱ	108.71 (18)	C14—C15—H15A	110.00
C11 ⁱ —C10—C15	109.02 (11)	C14—C15—H15B	110.00
C10—C11—C12	109.55 (15)	H15A—C15—H15B	108.00
C11—C12—C13	109.96 (17)	C12—C16—H16A	110.00
C11—C12—C16	109.64 (15)	C12—C16—H16B	110.00
C13—C12—C16	109.24 (16)	H16A—C16—H16B	108.00
C12—C13—C14	109.39 (16)	C12 ⁱ —C16—H16A	110.00
C13—C14—C15	109.73 (12)	C12 ⁱ —C16—H16B	110.00
C2—S1—C1—N1	0.00 (1)	N4—C3—C4—C5	0.00 (1)
C2—S1—C1—C10	180.00 (1)	N4—C3—C4—C9	180.00 (1)
C1—S1—C2—N2	0.00 (1)	C3—C4—C5—C6	180.00 (1)
C1—S1—C2—N3	-180.00 (1)	C9—C4—C5—C6	0.00 (1)
C1—N1—N2—C2	0.00 (1)	C3—C4—C9—C8	180.00 (1)
C1—N1—N2—C3	180.00 (1)	C5—C4—C9—C8	0.00 (1)
N2—N1—C1—S1	0.00 (1)	C4—C5—C6—F1A	180.00 (1)
N2—N1—C1—C10	180.00 (1)	C4—C5—C6—C7	0.00 (1)
N1—N2—C3—C4	0.00 (1)	F1A—C6—C7—C8	180.00 (1)
C3—N2—C2—N3	0.00 (1)	C5—C6—C7—C8	0.00 (1)
N1—N2—C2—S1	0.00 (1)	C6—C7—C8—C9	0.00 (1)
C3—N2—C2—S1	180.00 (1)	C7—C8—C9—C4	0.00 (1)
N1—N2—C3—N4	-180.00 (1)	C1—C10—C11—C12	-179.45 (16)
C2—N2—C3—C4	180.00 (1)	C15—C10—C11—C12	-58.8 (2)
C2—N2—C3—N4	0.00 (1)	C11 ⁱ —C10—C11—C12	59.90 (19)
N1—N2—C2—N3	180.00 (1)	C1—C10—C15—C14	180.00 (1)
N4—N3—C2—S1	180.00 (1)	C11—C10—C15—C14	59.27 (12)
N4—N3—C2—N2	0.00 (1)	C10—C11—C12—C13	59.6 (2)
C2—N3—N4—C3	0.00 (1)	C10—C11—C12—C16	-60.5 (2)
N3—N4—C3—C4	-180.00 (1)	C11—C12—C13—C14	-60.1 (2)
N3—N4—C3—N2	0.00 (1)	C16—C12—C13—C14	60.2 (2)
N1—C1—C10—C11	120.09 (13)	C11—C12—C16—C12 ⁱ	60.3 (2)
N1—C1—C10—C15	0.00 (1)	C13—C12—C16—C12 ⁱ	-60.27 (19)
S1—C1—C10—C15	180.00 (1)	C12—C13—C14—C15	60.2 (2)
S1—C1—C10—C11	-59.91 (13)	C12—C13—C14—C13 ⁱ	-60.4 (2)
N2—C3—C4—C5	180.00 (1)	C13—C14—C15—C10	-60.28 (14)
N2—C3—C4—C9	0.00 (1)		

Symmetry code: (i) $x, -y+3/2, z$.

Hydrogen-bond geometry (Å, °)

<i>D—H···A</i>	<i>D—H</i>	<i>H···A</i>	<i>D···A</i>	<i>D—H···A</i>
C11—H11 <i>A</i> ···F1 <i>A</i> ⁱⁱ	0.99	2.67	3.526 (3)	145

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