Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

4-{2-[(3,4-Dichlorophenyl)(methyl)amino]-4-methyl-1,3-thiazol-5-yl}-*N*-(3-methylphenyl)pyrimidin-2-amine

Hai-Bo Li,[a] Hai-Bo Shi[b,c]* and Wei-Xiao Hu[c]

[a]Nantong Center for Disease Control and Prevention, Nantong 226007, People's Republic of China, [b]Zhejiang Pharmaceutical College, Ningbo 315100, People's Republic of China, and [c]College of Pharmaceutical Science, Zhejiang University of Technology, Hangzhou 310032, People's Republic of China
Correspondence e-mail: nbblconba_shb@sohu.com

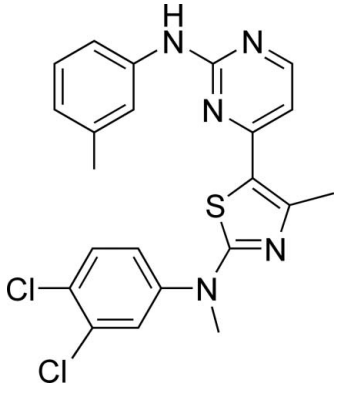
Received 29 November 2010; accepted 21 December 2010

Key indicators: single-crystal X-ray study; T = 103 K; mean σ(C–C) = 0.002 Å; R factor = 0.033; wR factor = 0.091; data-to-parameter ratio = 16.8.

In the title compound, $C_{22}H_{19}Cl_2N_5S$, the thiazole and pyrimidine rings are almost co-planar, making a dihedral angle of 6.48 (7)°. In the crystal, intermolecular N—H···N hydrogen bonds link pairs of molecules into centrosymmetric dimers..

Related literature

For general background to the biological activity of thiazole derivatives, see: Narayana *et al.* (2004). For the synthesis of the title compound, see: Bredereck *et al.* (1964).



Experimental

Crystal data

$C_{22}H_{19}Cl_2N_5S$
M_r = 456.38
Triclinic, $P\overline{1}$
a = 7.9679 (15) Å
b = 9.4042 (19) Å
c = 14.166 (3) Å
α = 85.421 (6)°
β = 76.727 (5)°
γ = 86.002 (6)°
V = 1028.4 (3) Å^3
Z = 2
Mo $K\alpha$ radiation
μ = 0.44 mm^{-1}
T = 103 K
0.53 × 0.50 × 0.40 mm

Data collection

Rigaku AFC10/Saturn724+ diffractometer
Absorption correction: multi-scan (*CrystalClear*; Rigaku/MSC, 2008)
T_{min} = 0.801, T_{max} = 0.844
9828 measured reflections
4626 independent reflections
3906 reflections with $I > 2\sigma(I)$
R_{int} = 0.021

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.033
$wR(F^2)$ = 0.091
S = 1.05
4626 reflections
275 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.37 e Å^{-3}
$\Delta\rho_{min}$ = −0.25 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
N5—H5*N*···N3^{i}	0.88	2.20	3.078 (2)	177

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

Data collection: *CrystalClear* (Rigaku/MSC, 2008); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *publCIF* (Westrip, 2010).

We are very thankful to the Natural Science Foundation of Ningbo City (grant No. 2009 A610185) for financial support. We are also grateful to the Beijing Institute of Technology for the X-ray diffraction measurements.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NG5079).

References

Bredereck, H., Effenberger, F. & Botsch, H. (1964). *Chem. Ber.* **97**, 3397–3406.
Narayana, B., Raj, K. K. V., Ashalatha, B. V., Kumari, N. S. & Sarojini, B. K. (2004). *Eur. J. Med. Chem.* **39**, 867–872.
Rigaku/MSC (2008). *CrystalClear*. Rigaku/MSC Inc., The Woodlands, Texas, USA.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.
Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.

supplementary materials

Acta Cryst. (2011). E**67**, o260 [**doi:10.1107/S1600536810053547**]

4-{2-[(3,4-Dichlorophenyl)(methyl)amino]-4-methyl-1,3-thiazol-5-yl}-*N*-(3-methylphenyl)pyrimidin-2-amine

H.-B. Li, H.-B. Shi and W.-X. Hu

Comment

Thiazole derivatives are found to be associated with various biological activities (Narayana *et al.*, 2004). In order to further study the structure-activity relationship (SAR) of the thiazolyl-pyrimidine derivatives, we introduced arylamino group into 2-position of thiazole ring of thiazolyl-pyrimidine according to the general pyrimidine condensation method of Bredereck (Bredereck *et al.*, 1964). But, it was found that the obtained compound was not desired compound that confirmed by ^{1}H NMR, MS. So, the structure of (I) was further determined using single-crystal X-ray diffraction.

The molecular structure of (I) is illustrated in Fig. 1. The thiazole ring (S1/C7/N2/C8/C9) and the pyrimidine ring (C10/C11/C12/N3/C13/N4) are almost planar, with a dihedral angle of 6.48 (7)°. The aniline rings (C1/C2/C3/C4/C5/C6/N1) and (C14/C15/C16/C17/C18/C19/N5) make dihedral angles of 73.76 (8) Å and 14.16 (7) Å with the thiazole ring, respectively. In the thiazole ring, the bond lengths S1—C7 [1.739 (15) Å], S1—C9 [1.749 (15) Å] and N2—C8 [1.377 (19) Å] correspond to typical single bond, and the C7—N2 [1.310 (2) Å], C8—C9 [1.372 (2) Å] belong to typical for double bonds. The crystal structure is stabilized by intermolecular weak N—H···N interactions (Fig. 2). Furthermore, every two molecules containing two N—H···N hydrogen bondings consists a dimer as octagon.

Experimental

A mixture of

3-dimethylamino-1-{2-[(3,4-Dichloro-phenyl)-methyl-amino]-4-methyl-thiazol-5-yl}-propenone (1.85 g, 5 mmol) and NaOH (0.2 g, 5 mmol) in 2-methoxylethanol (40 ml) was treated with *N*-*m*-tolyl-guanidine carbonate (1.58 g, 7.5 mmol). The reaction mixture was heated at 383 K under N_2 for 21 h. After concentration, the residue was filtered and washed liberally with ethanol and water. Recrystallization from THF affored the title compound as dark-brown crystals, 0.34 g, m.p.472–473 K, yield 15.0%. Since the crystal product was not found to be suitable for X-ray diffraction studies, a few crystals were dissolved in 2-butanone, which was allowed to evaporate slowly to give yellow crystals of (I) suitable for X-ray diffraction studies. ^{1}H NMR($CDCl_3$, TMS, 400 MHz, δp.p.m.): 8.30 (d, 1H, J =5.2 Hz, py—H), 7.61 (s, 1H, Ar—H), 7.57 (d, 1H, J = 2.8 Hz, Ar—H), 7.51 (d, 1H, J = 8.8 Hz, Ar—H), 7.33 (dd, 1H, J_1 = 2.4 Hz, J_2 = 2.4 Hz, Ar—H), 7.23–7.15 (m, 2H, Ar—H), 7.04 (s, 1H, Ar—H), 6.84 (d, 1H, J = 5.6 Hz, py—H), 3.54 (s, 3H, CH_3), 2.59 (s, 3H, CH_3), 2.27 (s, 3H, CH_3). EIMS m/z (%): 455 (M^+, 100), 440 (9), 295 (11), 256 (12), 222 (13), 213 (9), 185 (10),171 (8), 129 (19), 111 (9), 97 (16), 85 (19), 73 (41), 65 (8), 57 (48).

Refinement

All H atoms were placed in calculated positions (C—H 0.95–0.98 Å, N—H 0.87–0.89 Å) and refined as riding with U_{iso}(H) = 1.2–1.22Ueq of the parent atom.

Figures

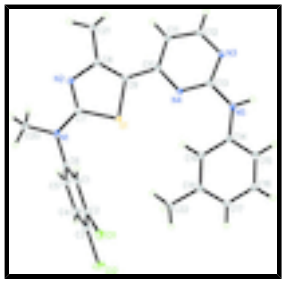


Fig. 1. The structure of (I), shown with 30% probability displacement ellipsoids.

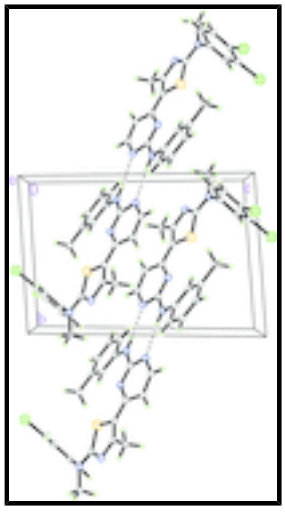


Fig. 2. Packing of the molecules down *a* axis. Dashed lines denote intermolecular N—H···N hydrogen bonds.

4-{2-[(3,4-Dichlorophenyl)(methyl)amino]-4-methyl-1,3-thiazol-5-yl}- *N*-(3-methylphenyl)pyrimidin-2-amine

Crystal data

$C_{22}H_{19}Cl_2N_5S$	$Z = 2$
$M_r = 456.38$	$F(000) = 472$
Triclinic, $P\overline{1}$	$D_x = 1.474$ Mg m^{-3}
Hall symbol: -P 1	Melting point = 473–472 K
$a = 7.9679$ (15) Å	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
$b = 9.4042$ (19) Å	Cell parameters from 3163 reflections
$c = 14.166$ (3) Å	$\theta = 3.3$–$27.5°$
$\alpha = 85.421$ (6)°	$\mu = 0.44$ mm^{-1}
$\beta = 76.727$ (5)°	$T = 103$ K
$\gamma = 86.002$ (6)°	Block, yellow
$V = 1028.4$ (3) Å^3	0.53 × 0.50 × 0.40 mm

Data collection

Rigaku AFC10/Saturn724+ diffractometer	4626 independent reflections
Radiation source: fine-focus sealed tube	3906 reflections with $I > 2\sigma(I)$
graphite	$R_{int} = 0.021$

Detector resolution: 28.5714 pixels mm^{-1}
phi and ω scans
Absorption correction: multi-scan (*CrystalClear*; Rigaku/MSC, 2008)
T_{min} = 0.801, T_{max} = 0.844
9828 measured reflections

θ_{max} = 27.5°, θ_{min} = 3.3°
h = −10→10
k = −12→12
l = −16→18

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.033
$wR(F^2)$ = 0.091
S = 1.05
4626 reflections
275 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
$w = 1/[\sigma^2(F_o^2) + (0.0486P)^2 + 0.1912P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max}$ = 0.002
$\Delta\rho_{max}$ = 0.37 e Å^{-3}
$\Delta\rho_{min}$ = −0.25 e Å^{-3}

Special details

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^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
Cl1	0.65348 (6)	0.39036 (4)	1.06940 (3)	0.02494 (11)
Cl2	1.02033 (6)	0.23777 (5)	0.99130 (3)	0.03270 (13)
S1	0.46927 (5)	0.38651 (4)	0.73411 (3)	0.01502 (10)
N1	0.42851 (17)	0.12266 (14)	0.82588 (10)	0.0177 (3)
N2	0.24614 (17)	0.20416 (14)	0.72255 (10)	0.0174 (3)
N3	0.40431 (17)	0.83042 (14)	0.50231 (10)	0.0166 (3)
N4	0.46548 (16)	0.64873 (13)	0.62050 (9)	0.0152 (3)
C1	0.5466 (2)	0.24613 (16)	0.93887 (11)	0.0167 (3)
H1	0.4370	0.2936	0.9610	0.020*
C2	0.6847 (2)	0.27394 (16)	0.97755 (11)	0.0172 (3)
C3	0.8457 (2)	0.20465 (18)	0.94480 (12)	0.0199 (3)
C4	0.8656 (2)	0.1063 (2)	0.87532 (13)	0.0260 (4)

H4	0.9749	0.0581	0.8535	0.031*
C5	0.7274 (2)	0.07723 (19)	0.83720 (12)	0.0227 (4)
H5	0.7415	0.0083	0.7901	0.027*
C6	0.5687 (2)	0.14890 (16)	0.86789 (11)	0.0158 (3)
C7	0.37223 (19)	0.22404 (16)	0.76455 (11)	0.0150 (3)
C8	0.2218 (2)	0.32085 (16)	0.66094 (11)	0.0161 (3)
C9	0.3305 (2)	0.42957 (16)	0.65541 (11)	0.0153 (3)
C10	0.3515 (2)	0.56427 (16)	0.59808 (11)	0.0144 (3)
C11	0.2660 (2)	0.60746 (16)	0.52437 (11)	0.0173 (3)
H11	0.1906	0.5473	0.5052	0.021*
C12	0.2970 (2)	0.74247 (17)	0.48071 (12)	0.0180 (3)
H12	0.2372	0.7750	0.4316	0.022*
C13	0.4895 (2)	0.77493 (16)	0.57055 (11)	0.0150 (3)
C14	0.7437 (2)	0.82794 (16)	0.63641 (11)	0.0155 (3)
C15	0.8818 (2)	0.91812 (16)	0.61470 (11)	0.0162 (3)
H15	0.8825	0.9958	0.5675	0.019*
C16	1.0172 (2)	0.89571 (16)	0.66104 (12)	0.0180 (3)
H16	1.1084	0.9595	0.6468	0.022*
C17	1.0204 (2)	0.78011 (17)	0.72849 (12)	0.0176 (3)
H17	1.1144	0.7642	0.7595	0.021*
C18	0.8856 (2)	0.68802 (17)	0.75029 (11)	0.0173 (3)
C19	0.7475 (2)	0.71314 (16)	0.70470 (11)	0.0179 (3)
H19	0.6546	0.6510	0.7205	0.021*
C20	0.3733 (2)	−0.02305 (17)	0.82886 (13)	0.0224 (4)
H20A	0.2555	−0.0291	0.8686	0.027*
H20B	0.4515	−0.0895	0.8573	0.027*
H20C	0.3759	−0.0482	0.7627	0.027*
C21	0.0762 (2)	0.31764 (18)	0.61070 (13)	0.0248 (4)
H21A	0.0086	0.4089	0.6166	0.030*
H21B	0.0020	0.2399	0.6408	0.030*
H21C	0.1229	0.3020	0.5418	0.030*
C22	0.8862 (2)	0.56045 (18)	0.82177 (13)	0.0243 (4)
H22A	0.8018	0.5787	0.8823	0.029*
H22B	1.0015	0.5437	0.8352	0.029*
H22C	0.8558	0.4761	0.7942	0.029*
N5	0.61031 (17)	0.86089 (14)	0.58696 (10)	0.0185 (3)
H5N	0.6034	0.9498	0.5630	0.032 (6)*

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0352 (2)	0.0191 (2)	0.0227 (2)	0.00067 (16)	−0.00980 (17)	−0.00635 (15)
Cl2	0.0238 (2)	0.0468 (3)	0.0331 (3)	−0.00328 (19)	−0.01516 (19)	−0.0099 (2)
S1	0.01644 (19)	0.01321 (18)	0.0177 (2)	−0.00395 (13)	−0.00857 (15)	0.00174 (14)
N1	0.0186 (7)	0.0134 (6)	0.0232 (7)	−0.0041 (5)	−0.0095 (6)	0.0033 (5)
N2	0.0165 (7)	0.0164 (7)	0.0209 (7)	−0.0034 (5)	−0.0075 (5)	0.0005 (5)
N3	0.0178 (7)	0.0153 (6)	0.0177 (7)	−0.0006 (5)	−0.0067 (5)	0.0004 (5)
N4	0.0166 (6)	0.0143 (6)	0.0163 (7)	−0.0020 (5)	−0.0068 (5)	−0.0005 (5)

C1	0.0175 (8)	0.0151 (7)	0.0167 (8)	0.0007 (6)	−0.0040 (6)	0.0021 (6)
C2	0.0249 (8)	0.0130 (7)	0.0140 (8)	−0.0015 (6)	−0.0057 (6)	0.0016 (6)
C3	0.0174 (8)	0.0255 (8)	0.0182 (8)	−0.0026 (6)	−0.0070 (6)	−0.0004 (6)
C4	0.0165 (8)	0.0362 (10)	0.0261 (10)	0.0036 (7)	−0.0055 (7)	−0.0100 (7)
C5	0.0197 (8)	0.0295 (9)	0.0201 (9)	0.0011 (7)	−0.0048 (7)	−0.0095 (7)
C6	0.0166 (7)	0.0159 (7)	0.0148 (8)	−0.0027 (6)	−0.0048 (6)	0.0043 (6)
C7	0.0149 (7)	0.0138 (7)	0.0161 (8)	−0.0026 (6)	−0.0028 (6)	−0.0007 (6)
C8	0.0158 (7)	0.0153 (7)	0.0186 (8)	−0.0011 (6)	−0.0066 (6)	−0.0016 (6)
C9	0.0155 (7)	0.0155 (7)	0.0167 (8)	−0.0001 (6)	−0.0073 (6)	−0.0018 (6)
C10	0.0160 (7)	0.0128 (7)	0.0156 (8)	0.0001 (5)	−0.0056 (6)	−0.0025 (6)
C11	0.0184 (8)	0.0172 (8)	0.0187 (8)	−0.0016 (6)	−0.0090 (6)	−0.0019 (6)
C12	0.0199 (8)	0.0181 (8)	0.0172 (8)	0.0020 (6)	−0.0078 (6)	−0.0004 (6)
C13	0.0169 (7)	0.0152 (7)	0.0129 (7)	−0.0004 (6)	−0.0034 (6)	−0.0011 (6)
C14	0.0164 (7)	0.0161 (7)	0.0145 (8)	−0.0023 (6)	−0.0042 (6)	−0.0019 (6)
C15	0.0184 (8)	0.0136 (7)	0.0151 (8)	−0.0020 (6)	−0.0008 (6)	−0.0001 (6)
C16	0.0145 (7)	0.0177 (8)	0.0210 (8)	−0.0034 (6)	−0.0005 (6)	−0.0042 (6)
C17	0.0156 (8)	0.0205 (8)	0.0180 (8)	−0.0001 (6)	−0.0056 (6)	−0.0050 (6)
C18	0.0210 (8)	0.0169 (7)	0.0152 (8)	−0.0014 (6)	−0.0062 (6)	−0.0014 (6)
C19	0.0208 (8)	0.0168 (8)	0.0179 (8)	−0.0068 (6)	−0.0072 (6)	0.0007 (6)
C20	0.0230 (9)	0.0149 (8)	0.0304 (10)	−0.0057 (6)	−0.0090 (7)	0.0055 (7)
C21	0.0248 (9)	0.0227 (9)	0.0321 (10)	−0.0076 (7)	−0.0168 (7)	0.0028 (7)
C22	0.0303 (9)	0.0217 (9)	0.0249 (9)	−0.0067 (7)	−0.0149 (7)	0.0051 (7)
N5	0.0228 (7)	0.0146 (7)	0.0203 (7)	−0.0053 (5)	−0.0101 (6)	0.0051 (5)

Geometric parameters (Å, °)

Cl1—C2	1.7304 (17)	C11—C12	1.380 (2)
Cl2—C3	1.7266 (17)	C11—H11	0.9500
S1—C7	1.7392 (15)	C12—H12	0.9500
S1—C9	1.7492 (15)	C13—N5	1.3694 (19)
N1—C7	1.361 (2)	C14—C19	1.394 (2)
N1—C6	1.424 (2)	C14—C15	1.399 (2)
N1—C20	1.4626 (19)	C14—N5	1.407 (2)
N2—C7	1.310 (2)	C15—C16	1.382 (2)
N2—C8	1.3778 (19)	C15—H15	0.9500
N3—C12	1.330 (2)	C16—C17	1.392 (2)
N3—C13	1.3561 (19)	C16—H16	0.9500
N4—C13	1.3346 (19)	C17—C18	1.391 (2)
N4—C10	1.3530 (19)	C17—H17	0.9500
C1—C2	1.386 (2)	C18—C19	1.396 (2)
C1—C6	1.386 (2)	C18—C22	1.507 (2)
C1—H1	0.9500	C19—H19	0.9500
C2—C3	1.394 (2)	C20—H20A	0.9800
C3—C4	1.380 (2)	C20—H20B	0.9800
C4—C5	1.384 (2)	C20—H20C	0.9800
C4—H4	0.9500	C21—H21A	0.9800
C5—C6	1.385 (2)	C21—H21B	0.9800
C5—H5	0.9500	C21—H21C	0.9800
C8—C9	1.372 (2)	C22—H22A	0.9800

C8—C21	1.497 (2)	C22—H22B	0.9800
C9—C10	1.447 (2)	C22—H22C	0.9800
C10—C11	1.392 (2)	N5—H5N	0.8800
C7—S1—C9	88.31 (7)	N4—C13—N3	126.28 (14)
C7—N1—C6	120.09 (13)	N4—C13—N5	119.49 (14)
C7—N1—C20	118.53 (13)	N3—C13—N5	114.23 (13)
C6—N1—C20	119.69 (12)	C19—C14—C15	118.38 (14)
C7—N2—C8	110.18 (13)	C19—C14—N5	124.83 (14)
C12—N3—C13	114.27 (13)	C15—C14—N5	116.80 (13)
C13—N4—C10	117.34 (13)	C16—C15—C14	120.89 (14)
C2—C1—C6	119.86 (15)	C16—C15—H15	119.6
C2—C1—H1	120.1	C14—C15—H15	119.6
C6—C1—H1	120.1	C15—C16—C17	120.25 (14)
C1—C2—C3	120.02 (15)	C15—C16—H16	119.9
C1—C2—Cl1	119.41 (12)	C17—C16—H16	119.9
C3—C2—Cl1	120.53 (13)	C18—C17—C16	119.82 (15)
C4—C3—C2	119.66 (15)	C18—C17—H17	120.1
C4—C3—Cl2	119.29 (13)	C16—C17—H17	120.1
C2—C3—Cl2	121.04 (13)	C17—C18—C19	119.58 (14)
C3—C4—C5	120.44 (16)	C17—C18—C22	121.23 (15)
C3—C4—H4	119.8	C19—C18—C22	119.20 (14)
C5—C4—H4	119.8	C14—C19—C18	121.06 (14)
C4—C5—C6	119.86 (16)	C14—C19—H19	119.5
C4—C5—H5	120.1	C18—C19—H19	119.5
C6—C5—H5	120.1	N1—C20—H20A	109.5
C5—C6—C1	120.11 (15)	N1—C20—H20B	109.5
C5—C6—N1	119.71 (15)	H20A—C20—H20B	109.5
C1—C6—N1	120.18 (14)	N1—C20—H20C	109.5
N2—C7—N1	122.48 (14)	H20A—C20—H20C	109.5
N2—C7—S1	115.90 (11)	H20B—C20—H20C	109.5
N1—C7—S1	121.59 (12)	C8—C21—H21A	109.5
C9—C8—N2	115.95 (14)	C8—C21—H21B	109.5
C9—C8—C21	127.26 (14)	H21A—C21—H21B	109.5
N2—C8—C21	116.74 (13)	C8—C21—H21C	109.5
C8—C9—C10	132.96 (14)	H21A—C21—H21C	109.5
C8—C9—S1	109.62 (11)	H21B—C21—H21C	109.5
C10—C9—S1	117.41 (11)	C18—C22—H22A	109.5
N4—C10—C11	120.73 (14)	C18—C22—H22B	109.5
N4—C10—C9	114.52 (13)	H22A—C22—H22B	109.5
C11—C10—C9	124.75 (14)	C18—C22—H22C	109.5
C12—C11—C10	116.32 (14)	H22A—C22—H22C	109.5
C12—C11—H11	121.8	H22B—C22—H22C	109.5
C10—C11—H11	121.8	C13—N5—C14	129.52 (13)
N3—C12—C11	124.83 (14)	C13—N5—H5N	115.2
N3—C12—H12	117.6	C14—N5—H5N	115.2
C11—C12—H12	117.6		
C6—C1—C2—C3	0.3 (2)	C7—S1—C9—C8	1.59 (12)
C6—C1—C2—Cl1	−177.54 (11)	C7—S1—C9—C10	−177.37 (13)

C1—C2—C3—C4	−1.4 (2)	C13—N4—C10—C11	1.2 (2)
Cl1—C2—C3—C4	176.42 (13)	C13—N4—C10—C9	−179.55 (13)
C1—C2—C3—Cl2	179.72 (12)	C8—C9—C10—N4	174.45 (17)
Cl1—C2—C3—Cl2	−2.50 (19)	S1—C9—C10—N4	−6.89 (19)
C2—C3—C4—C5	0.8 (3)	C8—C9—C10—C11	−6.3 (3)
Cl2—C3—C4—C5	179.69 (14)	S1—C9—C10—C11	172.32 (13)
C3—C4—C5—C6	1.0 (3)	N4—C10—C11—C12	−3.6 (2)
C4—C5—C6—C1	−2.1 (2)	C9—C10—C11—C12	177.22 (15)
C4—C5—C6—N1	177.66 (15)	C13—N3—C12—C11	2.2 (2)
C2—C1—C6—C5	1.5 (2)	C10—C11—C12—N3	1.8 (2)
C2—C1—C6—N1	−178.28 (13)	C10—N4—C13—N3	3.5 (2)
C7—N1—C6—C5	−108.63 (18)	C10—N4—C13—N5	−176.52 (14)
C20—N1—C6—C5	56.3 (2)	C12—N3—C13—N4	−5.1 (2)
C7—N1—C6—C1	71.1 (2)	C12—N3—C13—N5	174.89 (14)
C20—N1—C6—C1	−123.93 (16)	C19—C14—C15—C16	−1.2 (2)
C8—N2—C7—N1	−177.07 (14)	N5—C14—C15—C16	178.77 (14)
C8—N2—C7—S1	0.87 (18)	C14—C15—C16—C17	1.7 (2)
C6—N1—C7—N2	178.64 (14)	C15—C16—C17—C18	−0.9 (2)
C20—N1—C7—N2	13.5 (2)	C16—C17—C18—C19	−0.5 (2)
C6—N1—C7—S1	0.8 (2)	C16—C17—C18—C22	179.02 (15)
C20—N1—C7—S1	−164.32 (12)	C15—C14—C19—C18	−0.2 (2)
C9—S1—C7—N2	−1.46 (13)	N5—C14—C19—C18	179.85 (15)
C9—S1—C7—N1	176.49 (14)	C17—C18—C19—C14	1.0 (2)
C7—N2—C8—C9	0.5 (2)	C22—C18—C19—C14	−178.47 (16)
C7—N2—C8—C21	−177.22 (14)	N4—C13—N5—C14	14.1 (2)
N2—C8—C9—C10	177.21 (16)	N3—C13—N5—C14	−165.88 (15)
C21—C8—C9—C10	−5.4 (3)	C19—C14—N5—C13	−21.4 (3)
N2—C8—C9—S1	−1.52 (18)	C15—C14—N5—C13	158.68 (16)
C21—C8—C9—S1	175.88 (14)		

Hydrogen-bond geometry (Å, °)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N5—H5N···N3^{i}	0.88	2.20	3.078 (2)	177

Symmetry codes: (i) −*x*+1, −*y*+2, −*z*+1.

Fig. 1

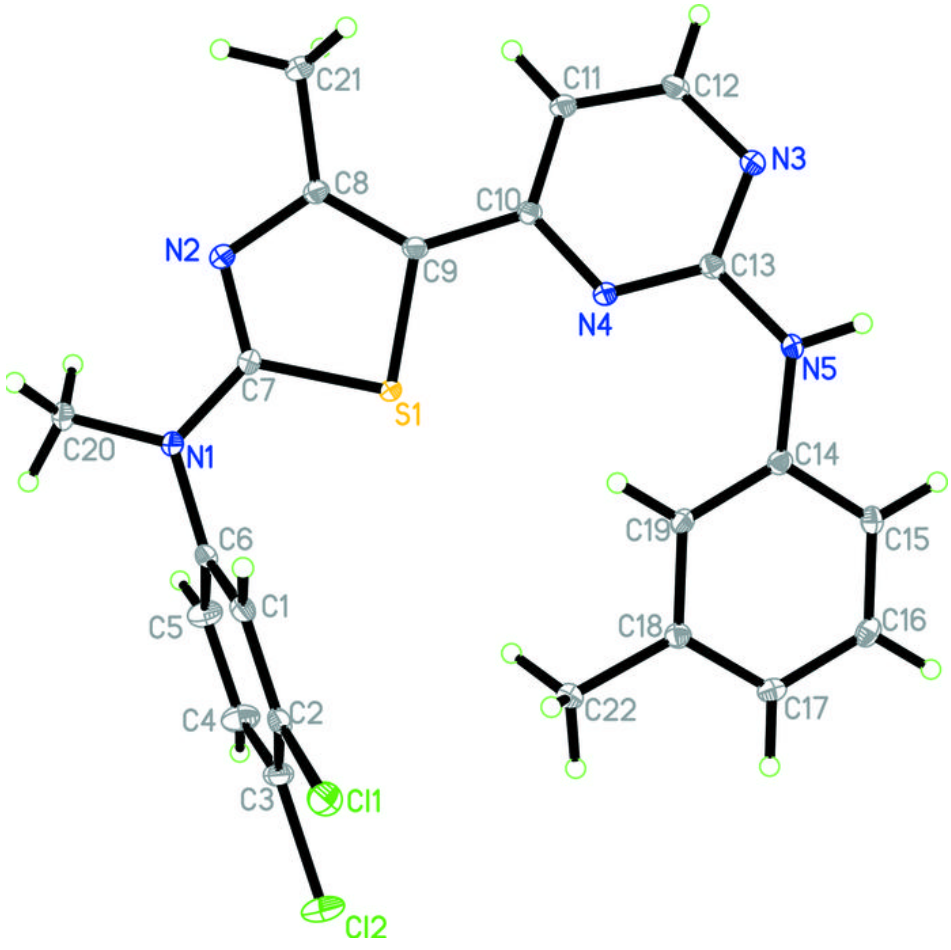

C21
C11
C12
N3
C8
C10
N2
C9
C13
N4
N5
C7
S1
C20
N1
C14
C19
C15
C6
C1
C5
C18
C16
C4
C2
C22
C17
C3
Cl1
Cl2

Fig. 2