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4-(4-Chlorophenyl)-1-methyl-3-trifluoromethyl-1*H*-pyrazol-5-amineXiao-Ping Jiang^a and Seik Weng Ng^{b*}^aDepartment of Biology and Chemistry, Hunan University of Science and Engineering, Yongzhou, Hunan 425100, People's Republic of China, and^bDepartment of Chemistry, University of Malaya, 50603 Kuala Lumpur, Malaysia

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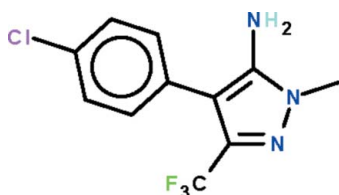
Received 12 August 2010; accepted 12 August 2010

Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.042; wR factor = 0.119; data-to-parameter ratio = 15.0.

The five-membered ring of the title compound, $\text{C}_{11}\text{H}_9\text{ClF}_3\text{N}_3$, is almost planar (r.m.s. deviation = 0.002 Å) and the phenylene ring is aligned at 44.8 (1)°. The N atom of the amino substituent shows a pyramidal geometry and is a hydrogen-bond donor to a Cl atom and to a ring N atom, which together generate a layer motif.

Related literature

For the synthesis of the title compound, see: Coispeau (1977); Nishiwaki *et al.* (1995).



Experimental

Crystal data

 $\text{C}_{11}\text{H}_9\text{ClF}_3\text{N}_3$ $M_r = 275.66$ Monoclinic, $P2_1/c$ $a = 5.8958$ (5) Å $b = 16.8618$ (13) Å $c = 12.1087$ (10) Å $\beta = 98.459$ (1)° $V = 1190.68$ (17) Å³ $Z = 4$ Mo $K\alpha$ radiation $\mu = 0.34$ mm⁻¹ $T = 293$ K

0.40 × 0.40 × 0.20 mm

Data collection

Bruker SMART area-detector

diffractometer

Absorption correction: multi-scan

(SADABS; Sheldrick, 1996)

 $T_{\min} = 0.864$, $T_{\max} = 1.000$

5716 measured reflections

2581 independent reflections

1769 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.029$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.042$ $wR(F^2) = 0.119$ $S = 1.02$

2581 reflections

172 parameters

2 restraints

H atoms treated by a mixture of independent and constrained refinement

 $\Delta\rho_{\max} = 0.32$ e Å⁻³ $\Delta\rho_{\min} = -0.26$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N3}-\text{H1}\cdots\text{Cl1}^i$	0.88 (1)	2.65 (2)	3.413 (2)	146 (2)
$\text{N3}-\text{H2}\cdots\text{N2}^{ii}$	0.88 (1)	2.54 (2)	3.180 (3)	130 (2)

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

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HG2700).

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supporting information

Acta Cryst. (2010). E66, o2336 [https://doi.org/10.1107/S1600536810032435]

4-(4-Chlorophenyl)-1-methyl-3-trifluoromethyl-1*H*-pyrazol-5-amine**Xiao-Ping Jiang and Seik Weng Ng****S1. Comment**

The title compound (Scheme I) is first mentioned in a synthesis by the cyclocondensation of hydrazines with 4,4,4-trifluoro-2-arylacetoacetonitriles in the context of colorants for polyacrylonitriles (Coispeau, 1977). The structure has been elucidated by carbon-13 NMR spectroscopy (Nishiwaki *et al.*, 1995). We have used a modification of the published procedure to synthesize the compound, which is intended for further study on its pharmaceutical activity. There are no other reports on this compound aside from these studies.

S2. Experimental

Sodium metal (0.35 g, 15 mmol) was dissolved in absolute ethanol (50 ml). To this solution was added 2-(4-chlorophenyl)acetonitrile (1.52 g, 10 mmol) followed by ethyl trifluoroacetate (1.42 g, 10 mmol). The solution was heated for 3 h. The solution was concentrated under vacuum. To the residue was added acetic acid (20 ml) followed by methylhydrazine (0.55 g, 12 mmol). The mixture was stirred for 12 h. The solution was again concentrated under vacuum. The residue was treated with water (30 ml) and the organic compound was extracted with ethyl acetate. The organic phase was washed with saturated sodium bicarbonate (230 ml) and then dried over sodium sulfate. The solvent was removed and the residue was chromatographed on a silica gel column with ethyl acetate:petroleum ether (1:10) as eluant. This gave 2 g (70%) of product as a yellow solid, which was recrystallized from ethyl acetate.

S3. Refinement

Carbon bound H-atoms were positioned geometrically and refined using the riding model, and with C–H = 0.93 to 0.96 Å and $U(H)$ set to 1.2–1.5 $U_{eq}(C)$. The amino H-atoms were located in a difference Fourier map, and were refined with a distance restraint of N–H 0.88±0.01 Å; their temperature factors were refined.

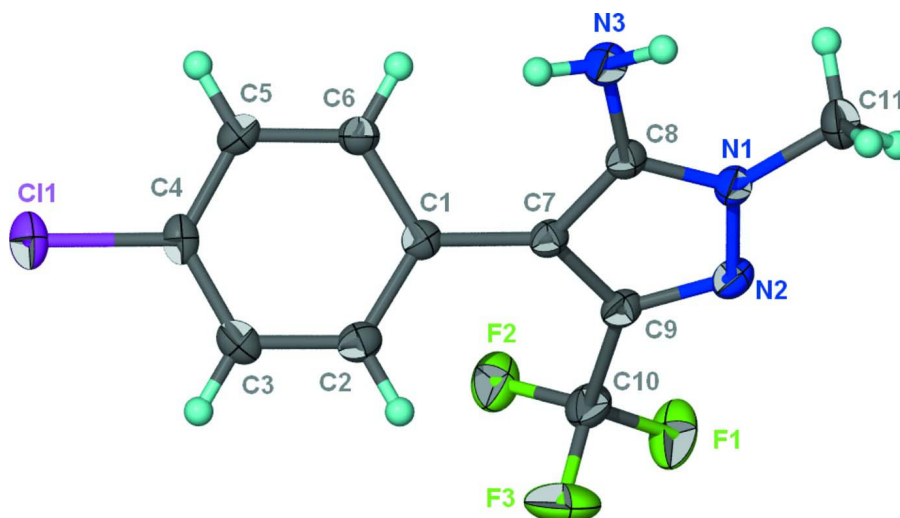


Figure 1

Thermal ellipsoid plot (Barbour, 2001) of $C_{11}H_9ClF_3N_3$ showing displacement ellipsoids at the 50% probability level. H-atoms are drawn as spheres of arbitrary radius.

4-(4-Chlorophenyl)-1-methyl-3-trifluoromethyl-1H-pyrazol-5-amine

Crystal data

$C_{11}H_9ClF_3N_3$

$M_r = 275.66$

Monoclinic, $P2_1/c$

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

$a = 5.8958$ (5) Å

$b = 16.8618$ (13) Å

$c = 12.1087$ (10) Å

$\beta = 98.459$ (1)°

$V = 1190.68$ (17) Å³

$Z = 4$

$F(000) = 560$

$D_x = 1.538$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 2152 reflections

$\theta = 2.5$ – 26.6 °

$\mu = 0.34$ mm⁻¹

$T = 293$ K

Block, yellow

$0.40 \times 0.40 \times 0.20$ mm

Data collection

Bruker SMART area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.864$, $T_{\max} = 1.000$

5716 measured reflections

2581 independent reflections

1769 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.029$

$\theta_{\max} = 27.0$ °, $\theta_{\min} = 2.1$ °

$h = -7 \rightarrow 7$

$k = -21 \rightarrow 10$

$l = -15 \rightarrow 12$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.042$

$wR(F^2) = 0.119$

$S = 1.02$

2581 reflections

172 parameters

2 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

$$w = 1/[\sigma^2(F_o^2) + (0.0629P)^2 + 0.1525P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$

$$\Delta\rho_{\max} = 0.32 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.26 \text{ e } \text{\AA}^{-3}$$

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	0.24293 (10)	0.63048 (3)	0.86611 (5)	0.03743 (19)
F1	0.9033 (3)	0.36727 (9)	0.43732 (12)	0.0619 (5)
F2	0.7674 (3)	0.46344 (8)	0.52371 (12)	0.0503 (4)
F3	0.5700 (3)	0.35880 (11)	0.48636 (14)	0.0685 (5)
N1	1.1139 (3)	0.27925 (10)	0.73828 (14)	0.0272 (4)
N2	1.0499 (3)	0.29559 (10)	0.62838 (14)	0.0292 (4)
N3	1.0445 (3)	0.31883 (12)	0.91929 (16)	0.0314 (4)
H1	0.929 (3)	0.3386 (15)	0.948 (2)	0.054 (8)*
H2	1.087 (4)	0.2714 (8)	0.944 (2)	0.044 (7)*
C1	0.6890 (4)	0.43079 (11)	0.77159 (17)	0.0238 (5)
C2	0.4645 (4)	0.43995 (12)	0.71925 (18)	0.0283 (5)
H2A	0.4060	0.4049	0.6629	0.034*
C3	0.3258 (4)	0.50032 (13)	0.74945 (19)	0.0293 (5)
H3	0.1767	0.5063	0.7129	0.035*
C4	0.4114 (4)	0.55085 (12)	0.83380 (19)	0.0271 (5)
C5	0.6307 (4)	0.54242 (12)	0.89124 (18)	0.0282 (5)
H5	0.6846	0.5763	0.9499	0.034*
C6	0.7681 (4)	0.48252 (12)	0.85957 (18)	0.0266 (5)
H6	0.9161	0.4764	0.8975	0.032*
C7	0.8443 (4)	0.37063 (11)	0.73603 (18)	0.0239 (4)
C8	0.9973 (3)	0.32257 (12)	0.80477 (18)	0.0253 (5)
C9	0.8878 (4)	0.35046 (12)	0.62864 (18)	0.0267 (5)
C10	0.7831 (4)	0.38467 (14)	0.5202 (2)	0.0377 (6)
C11	1.2992 (4)	0.22382 (14)	0.7729 (2)	0.0398 (6)
H11A	1.4002	0.2456	0.8349	0.060*
H11B	1.3831	0.2146	0.7119	0.060*
H11C	1.2369	0.1746	0.7948	0.060*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0397 (3)	0.0332 (3)	0.0428 (4)	0.0094 (2)	0.0172 (3)	0.0028 (2)
F1	0.0999 (14)	0.0632 (11)	0.0261 (8)	0.0360 (9)	0.0205 (8)	0.0060 (7)
F2	0.0765 (11)	0.0380 (8)	0.0359 (9)	0.0191 (7)	0.0067 (7)	0.0071 (6)
F3	0.0630 (11)	0.0885 (13)	0.0445 (10)	-0.0080 (9)	-0.0238 (9)	-0.0024 (9)
N1	0.0322 (10)	0.0243 (9)	0.0245 (10)	0.0041 (8)	0.0025 (8)	-0.0016 (7)
N2	0.0370 (11)	0.0260 (9)	0.0242 (10)	-0.0007 (8)	0.0037 (8)	-0.0031 (7)
N3	0.0414 (12)	0.0300 (11)	0.0223 (10)	0.0046 (9)	0.0023 (9)	0.0029 (8)
C1	0.0276 (11)	0.0209 (10)	0.0230 (11)	-0.0014 (8)	0.0040 (9)	0.0029 (8)
C2	0.0305 (12)	0.0269 (11)	0.0273 (12)	-0.0045 (9)	0.0036 (9)	-0.0009 (9)
C3	0.0244 (11)	0.0300 (11)	0.0334 (13)	-0.0007 (9)	0.0039 (9)	0.0074 (9)

C4	0.0304 (12)	0.0221 (10)	0.0316 (12)	0.0037 (9)	0.0139 (10)	0.0055 (9)
C5	0.0333 (12)	0.0274 (11)	0.0243 (11)	-0.0033 (9)	0.0057 (9)	-0.0041 (9)
C6	0.0272 (11)	0.0258 (11)	0.0259 (12)	-0.0005 (9)	0.0007 (9)	-0.0002 (9)
C7	0.0272 (11)	0.0212 (10)	0.0232 (11)	-0.0017 (8)	0.0030 (9)	-0.0003 (8)
C8	0.0283 (11)	0.0205 (10)	0.0271 (12)	-0.0035 (8)	0.0034 (9)	-0.0004 (9)
C9	0.0312 (11)	0.0228 (10)	0.0257 (12)	-0.0028 (9)	0.0026 (9)	-0.0028 (9)
C10	0.0496 (15)	0.0362 (13)	0.0260 (13)	0.0060 (11)	0.0014 (11)	-0.0025 (10)
C11	0.0408 (14)	0.0353 (13)	0.0425 (15)	0.0127 (11)	0.0038 (11)	-0.0030 (11)

Geometric parameters (Å, °)

C11—C4	1.749 (2)	C2—C3	1.388 (3)
F1—C10	1.344 (3)	C2—H2A	0.9300
F2—C10	1.333 (3)	C3—C4	1.368 (3)
F3—C10	1.336 (3)	C3—H3	0.9300
N1—C8	1.348 (3)	C4—C5	1.382 (3)
N1—N2	1.357 (2)	C5—C6	1.384 (3)
N1—C11	1.452 (3)	C5—H5	0.9300
N2—C9	1.331 (3)	C6—H6	0.9300
N3—C8	1.375 (3)	C7—C8	1.394 (3)
N3—H1	0.88 (1)	C7—C9	1.404 (3)
N3—H2	0.88 (1)	C9—C10	1.483 (3)
C1—C2	1.389 (3)	C11—H11A	0.9600
C1—C6	1.402 (3)	C11—H11B	0.9600
C1—C7	1.473 (3)	C11—H11C	0.9600
C8—N1—N2	112.53 (17)	C5—C6—H6	119.3
C8—N1—C11	127.18 (19)	C1—C6—H6	119.3
N2—N1—C11	120.18 (18)	C8—C7—C9	102.82 (18)
C9—N2—N1	103.59 (17)	C8—C7—C1	126.98 (19)
C8—N3—H1	109.5 (19)	C9—C7—C1	130.08 (19)
C8—N3—H2	113.0 (17)	N1—C8—N3	122.16 (19)
H1—N3—H2	114 (2)	N1—C8—C7	107.49 (19)
C2—C1—C6	117.73 (19)	N3—C8—C7	130.2 (2)
C2—C1—C7	122.32 (18)	N2—C9—C7	113.57 (19)
C6—C1—C7	119.94 (19)	N2—C9—C10	118.3 (2)
C3—C2—C1	121.3 (2)	C7—C9—C10	128.1 (2)
C3—C2—H2A	119.3	F2—C10—F3	105.6 (2)
C1—C2—H2A	119.3	F2—C10—F1	106.7 (2)
C4—C3—C2	119.1 (2)	F3—C10—F1	105.95 (19)
C4—C3—H3	120.5	F2—C10—C9	112.43 (19)
C2—C3—H3	120.5	F3—C10—C9	113.2 (2)
C3—C4—C5	121.79 (19)	F1—C10—C9	112.3 (2)
C3—C4—C11	119.06 (17)	N1—C11—H11A	109.5
C5—C4—C11	119.10 (17)	N1—C11—H11B	109.5
C4—C5—C6	118.5 (2)	H11A—C11—H11B	109.5
C4—C5—H5	120.7	N1—C11—H11C	109.5
C6—C5—H5	120.7	H11A—C11—H11C	109.5

C5—C6—C1	121.4 (2)	H11B—C11—H11C	109.5
C8—N1—N2—C9	-0.4 (2)	N2—N1—C8—C7	0.5 (2)
C11—N1—N2—C9	-176.89 (19)	C11—N1—C8—C7	176.7 (2)
C6—C1—C2—C3	2.8 (3)	C9—C7—C8—N1	-0.3 (2)
C7—C1—C2—C3	-175.81 (19)	C1—C7—C8—N1	-176.80 (19)
C1—C2—C3—C4	-1.1 (3)	C9—C7—C8—N3	175.8 (2)
C2—C3—C4—C5	-1.3 (3)	C1—C7—C8—N3	-0.6 (4)
C2—C3—C4—C11	176.15 (16)	N1—N2—C9—C7	0.2 (2)
C3—C4—C5—C6	2.0 (3)	N1—N2—C9—C10	177.86 (18)
C11—C4—C5—C6	-175.48 (16)	C8—C7—C9—N2	0.1 (2)
C4—C5—C6—C1	-0.2 (3)	C1—C7—C9—N2	176.4 (2)
C2—C1—C6—C5	-2.1 (3)	C8—C7—C9—C10	-177.3 (2)
C7—C1—C6—C5	176.54 (19)	C1—C7—C9—C10	-1.0 (4)
C2—C1—C7—C8	-138.4 (2)	N2—C9—C10—F2	-134.0 (2)
C6—C1—C7—C8	43.0 (3)	C7—C9—C10—F2	43.3 (3)
C2—C1—C7—C9	46.1 (3)	N2—C9—C10—F3	106.4 (2)
C6—C1—C7—C9	-132.4 (2)	C7—C9—C10—F3	-76.3 (3)
N2—N1—C8—N3	-176.08 (18)	N2—C9—C10—F1	-13.6 (3)
C11—N1—C8—N3	0.1 (3)	C7—C9—C10—F1	163.7 (2)

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>
N3—H1...C11 ⁱ	0.88 (1)	2.65 (2)	3.413 (2)	146 (2)
N3—H2...N2 ⁱⁱ	0.88 (1)	2.54 (2)	3.180 (3)	130 (2)

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