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Key indicators

Single-crystal X-ray study

T = 298 K

Mean $\sigma(\text{C}-\text{C}) = 0.005 \text{ \AA}$

Disorder in main residue

R factor = 0.072

wR factor = 0.215

Data-to-parameter ratio = 13.3

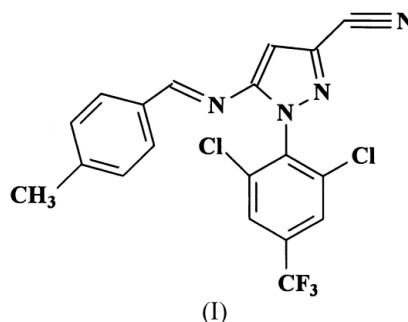
For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.1-[2,6-Dichloro-4-(trifluoromethyl)phenyl]-
5-[(4-methylbenzylidene)amino]-1H-pyrazole-
3-carbonitrileThe title compound, $\text{C}_{19}\text{H}_{11}\text{Cl}_2\text{F}_3\text{N}_4$, is a tricyclic amide with an overall U-shape, each of the three rings being planar. These include a benzene ring with two chloro and one trifluoromethyl substituent, a central pyrazole ring with a cyano substituent, and a benzene ring with one methyl substituent.

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Comment

The title compound, (I), is an important starting material for the synthesis of 5-amino-3-cyano-1-[2,6-dichloro-4-(trifluoromethyl)phenyl]-4-(trifluoromethyl)thiopyrazole, 5-amino-3-cyano-1-[2,6-dichloro-4-(trifluoromethyl)phenyl]-4-(trifluoromethyl)sulfonylpyrazole and 5-amino-3-cyano-1-[2,6-dichloro-4-(trifluoromethyl)phenyl]-4-(trifluoromethyl)sulfonylpyrazole, all of which are good insecticides (Hatton *et al.*, 1993).The structure of (I) is shown in Fig. 1, with the atom-numbering scheme. The molecule contains three planar groups, forming an overall U-shape, *viz.* a 2,6-dichloro-4-(trifluoromethyl)phenyl, a pyrazole and a 4-methylphenyl group. The dihedral angles between the pyrazole and the C1–C6 and C12–C17 benzene rings are 3.6 (3) and 78.7 (1)°, respectively.

Experimental

Following the method of Hatton *et al.* (1993), reaction of 2,6-dichloro-4-(trifluoromethyl)phenylamine with a suspension of nitrosyl sulfuric acid followed by reaction with a solution of ethyl 2,3-dicyanopropionate in acetic acid gave 5-amino-3-cyano-1-[2,6-dichloro-4-(trifluoromethyl)phenyl]pyrazole, which was then reacted with 4-methylbenzaldehyde and hydrochloric acid in anhydrous ethanol to give the title compound. Single crystals suitable for X-ray analysis were obtained by slow evaporation of an ethyl acetate/petroleum ether (1:2) solution (m.p. 460–462 K). IR (KBr, $\nu \text{ cm}^{-1}$): 2242, 1603, 1568, 1523, 1507, 1356, 1307, 881, 826; ^1H NMR (CDCl_3): δ 9.02 (s, 1H), 8.12 (s, 2H), 7.71 (d, $J = 8.0$ Hz, 2H), 7.30 (d, $J = 8.0$ Hz, 2H), 7.25 (s, 1H), 2.39 (s, 3H); ^{13}C NMR (CDCl_3): δ 166.6 (1C), 154.0 (1C),

145.0 (1C), 138.1 (1C), 134.1 (1C), 133.4 (1C), 130.5 (2C), 130.4 (2C), 128.8 (1C), 127.2 (2C), 127.1 (2C), 119.7 (1C), 114.3 (1C), 98.5 (1C), 21.7 (1C).

Crystal data

C₁₉H₁₁Cl₂F₃N₄
M_r = 423.22
 Triclinic, *P*1̄
a = 8.3045 (7) Å
b = 10.5134 (10) Å
c = 11.8019 (11) Å
 α = 76.106 (1)°
 β = 87.116 (2)°
 γ = 72.072 (2)°
V = 951.38 (15) Å³

Z = 2
D_x = 1.477 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 2515 reflections
 θ = 1.8–25.2°
 μ = 0.38 mm⁻¹
T = 298 (2) K
 Block, colourless
 0.18 × 0.15 × 0.13 mm

Data collection

Bruker SMART APEX area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 2002)
T_{min} = 0.932, *T_{max}* = 0.952
 5082 measured reflections

3428 independent reflections
 2829 reflections with *I* > 2σ(*I*)
R_{int} = 0.010
 θ_{max} = 25.2°
h = -9 → 9
k = -10 → 12
l = -14 → 14

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.072
wR (*F*²) = 0.215
S = 1.06
 3359 reflections
 253 parameters
 H-atom parameters constrained

w = 1/[σ²(*F_o*²) + (0.1186*P*)² + 0.8905*P*]
 where *P* = (*F_o*² + 2*F_c*²)/3
 (Δ/*σ*)_{max} < 0.001
 Δρ_{max} = 1.09 e Å⁻³
 Δρ_{min} = -0.52 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

N1—C8	1.266 (4)	N4—C12	1.140 (5)
N1—C9	1.390 (4)	C5—C8	1.455 (5)
N2—C11	1.335 (5)	C9—C10	1.364 (5)
N2—N3	1.344 (4)	C10—C11	1.391 (5)
N3—C9	1.367 (4)	C11—C12	1.437 (5)
N3—C13	1.421 (4)		
C8—N1—C9	118.5 (3)	C9—C10—C11	104.8 (3)
C11—N2—N3	102.9 (3)	N2—C11—C10	113.1 (3)
N2—N3—C9	113.2 (3)	N2—C11—C12	120.6 (3)
N2—N3—C13	121.3 (3)	C10—C11—C12	126.4 (3)
C9—N3—C13	125.4 (3)	N4—C12—C11	176.3 (5)
N1—C8—C5	122.7 (3)	C14—C13—N3	121.4 (3)
C10—C9—N3	106.0 (3)	C18—C13—N3	119.9 (3)
C10—C9—N1	136.3 (3)	F2—C19—F1	106.1 (6)
N3—C9—N1	117.7 (3)	F2—C19—C16	115.8 (4)

All H atoms were initially located in a difference Fourier map, and were then placed in geometrically idealized positions and constrained to ride on their parent atoms, with C—H distances in the range 0.93–

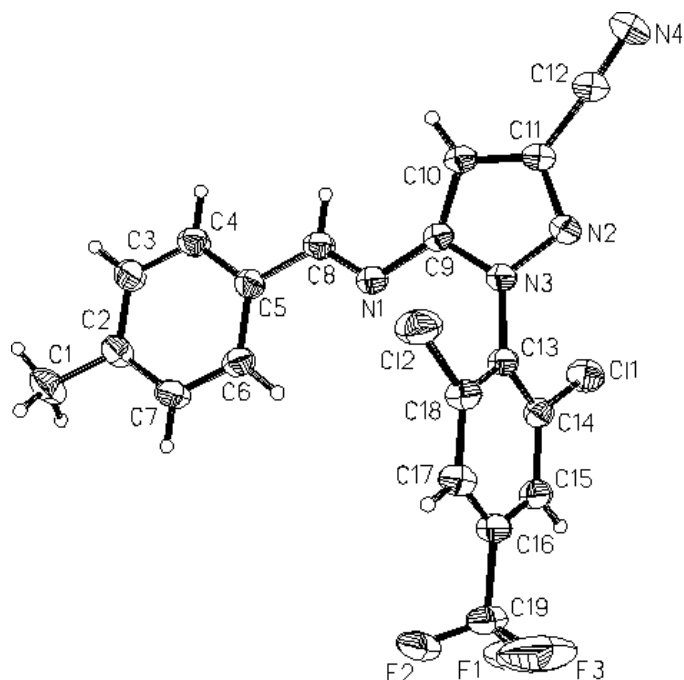


Figure 1

The structure (I), showing the atom-numbering scheme and with displacement ellipsoids at the 50% probability level. Only one disorder component is shown.

0.96 Å and with *U*_{iso}(H) = 1.22*U*_{eq}(C). The low *U*_{eq} value of C19 compared with its neighbours may be attributed to the three disordered F atoms.

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: XP (Bruker, 2002); software used to prepare material for publication: SHELXTL (Bruker, 2002).

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