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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.004 Å R factor = 0.050 wR factor = 0.135 Data-to-parameter ratio = 16.9

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

5-{[(4-Chlorophenyl)methylene]amino}-1-[2,6-dichloro-4-(trifluoromethyl)phenyl]-1*H*-pyrazole-3-carbonitrile

The title compound, $C_{18}H_8Cl_3F_3N_4$, is a tricyclic imide with an overall U-shape, each of the three rings being planar. The packing is stabilized by π - π interactions.

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Comment

The molecular structure of the title compound, (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-chlorophenyl ring. Bond lengths and angles (Table 1) are in agreement with those observed in similar compounds (Zhong, Yang & Shi, 2005; Zhong, Yang, Shi *et al.*, 2005; Chen *et al.*, 2005). The dihedral angles between the pyrazole and the C1–C6 and C12–C17 benzene rings are 19.02 (16) and 86.98 (8)°, respectively. There are π - π interactions between the pyrazole ring and the C1–C6 benzene ring. In the crystal structure, the molecules stack along the *a* axis, as shown in Fig. 2.



Experimental

Following the method of Hatton *et al.* (1993), reaction of 2,6-dichloro-4-(trifluoromethyl)aniline with a suspension of nitrosylsulfuric 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-chlorobenzaldehyde and hydrochloric acid in anhydrous ethanol to give the title compound, (I). Single crystals suitable for X-ray analysis were obtained by slow evaporation of an ethyl acetate/petroleum ether (1:2) solution (m.p. 452–454 K). IR (KBr, ν cm⁻¹): 3081, 2238, 2234, 1610, 1507, 1310, 880, 825; ¹H NMR (CDCl₃, δ): 9.08 (*s*, 1H), 8.11 (*s*, 2H), 7.84 (*d*, 2H), 7.50 (*d*, 2H), 7.29 (*s*, 1H); ¹³C NMR (CDCl₃, δ): 166.0 (1C), 153.9 (1C), 140.0 (1C), 137.0 (1C), 135.1 (1C), 134.7 (1C), 132.2 (2C), 130.6 (2C), 128.8 (1C), 127.5 (2C), 127.4 (2C), 123.8 (1C), 114.6 (1C), 99.5 (1C).

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Figure 1

The structure of (I), showing the atomic numbering scheme and displacement ellipsoids at the 50% probability level.



Figure 2

Packing diagram for (I), viewed down the c axis. H atoms have been omitted for clarity.

Crystal data

$C_{18}H_8Cl_3F_3N_4$	Z = 2	
$M_r = 443.64$	$D_x = 1.574 \text{ Mg m}^{-3}$	
Triclinic, $P\overline{1}$	Mo $K\alpha$ radiation	
a = 9.7553 (9) Å	Cell parameters from 3541	
b = 9.7554 (9) Å	reflections	
c = 11.7319 (11) Å	$\theta = 1.9-28.2^{\circ}$	
$\alpha = 68.582 \ (1)^{\circ}$	$\mu = 0.53 \text{ mm}^{-1}$	
$\beta = 68.580 \ (1)^{\circ}$	T = 298 (2) K	
$\gamma = 69.340 \ (1)^{\circ}$	Block, colourless	
$V = 936.06 (15) \text{ Å}^3$	$0.22 \times 0.18 \times 0.15 \text{ mm}$	

Data collection

 $R[F^2 > 2\sigma(F^2)] = 0.050$ wR(F²) = 0.135

S = 1.03

4267 reflections

253 parameters

Bruker SMART APEX area-
detector diffractometer
φ and ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 2002)
$T_{\min} = 0.892, \ T_{\max} = 0.924$
5873 measured reflections
Refinement
Refinement on F^2

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0645P)^{2} + 0.354P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 0.33 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.41 \text{ e} \text{ Å}^{-3}$

 $R_{\rm int} = 0.012$

 $\begin{array}{l} \theta_{\max} = 28.2^{\circ} \\ h = -12 \rightarrow 12 \end{array}$

 $\begin{array}{l} k = -12 \rightarrow 12 \\ l = -15 \rightarrow 9 \end{array}$

4627 independent reflections

3320 reflections with $I > 2\sigma(I)$

Table 1 Selected geometric parameters (Å, °).

H-atom parameters constrained

Cl1-C1	1.740 (2)	N3-C10	1.332 (3)
F1-C18	1.311 (3)	N4-C11	1.133 (3)
N1-C7	1.268 (3)	C4-C7	1.453 (3)
N1-C8	1.387 (3)	C8-C9	1.368 (3)
N2-N3	1.345 (2)	C9-C10	1.392 (3)
N2-C8	1.366 (3)	C10-C11	1.437 (3)
N2-C12	1.424 (2)	C15-C18	1.507 (3)
C7-N1-C8	118.03 (18)	C8-C9-C10	104.57 (19)
N3-N2-C8	113.56 (16)	N3-C10-C9	113.45 (19)
N3-N2-C12	122.26 (17)	N3-C10-C11	120.3 (2)
C8-N2-C12	123.83 (17)	C9-C10-C11	126.2 (2)
C10-N3-N2	102.62 (17)	N4-C11-C10	176.5 (3)
N1-C7-C4	123.2 (2)	C17-C12-N2	119.97 (18)
N2-C8-C9	105.78 (18)	F1-C18-F2	107.3 (3)
N2-C8-N1	118.46 (18)	F1-C18-C15	112.6 (2)
C9-C8-N1	135.8 (2)		

All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H distances set at 0.93 Å and $U_{iso}(H)$ values set at $1.2U_{eq}(C)$.

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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