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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=298 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.050$
$w R$ 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.
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## 5-\{[(4-Chlorophenyl)methylene]amino\}-1-[2,6-dichloro-4-(trifluoromethyl)phenyl]-1H-pyrazole-3-carbonitrile

The title compound, $\mathrm{C}_{18} \mathrm{H}_{8} \mathrm{Cl}_{3} \mathrm{~F}_{3} \mathrm{~N}_{4}$, is a tricyclic imide with an overall U-shape, each of the three rings being planar. The packing is stabilized by $\pi-\pi$ interactions.

## 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 C1C 6 and $\mathrm{C} 12-\mathrm{C} 17$ benzene rings are 19.02 (16) and 86.98 (8) ${ }^{\circ}$, respectively. There are $\pi-\pi$ 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.

(I)

## 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 ( $\mathrm{KBr}, v \mathrm{~cm}^{-1}$ ): 3081, 2238, 2234, $1610,1507,1310,880,825 ;{ }^{1} \mathrm{H}$ NMR ( $\left.\mathrm{CDCl}_{3}, \delta\right): 9.08(s, 1 \mathrm{H}), 8.11(s$, $2 \mathrm{H}), 7.84(d, 2 \mathrm{H}), 7.50(d, 2 \mathrm{H}), 7.29(s, 1 \mathrm{H}) ;{ }^{13} \mathrm{C}$ NMR $\left(\mathrm{CDCl}_{3}, \delta\right)$ : 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).


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

| $\mathrm{C}_{18} \mathrm{H}_{8} \mathrm{Cl}_{3} \mathrm{~F}_{3} \mathrm{~N}_{4}$ | $Z=2$ |
| :--- | :--- |
| $M_{r}=443.64$ | $D_{x}=1.574 \mathrm{Mg} \mathrm{m}^{-3}$ |
| Triclinic, $P \overline{1}$ | Mo $K \alpha$ radiation |
| $a=9.7553(9) \AA$ | Cell parameters from 3541 |
| $b=9.7554(9) \AA$ | reflections |
| $c=11.7319(11) \AA$ | $\theta=1.9-28.2^{\circ}$ |
| $\alpha=68.582(1)^{\circ}$ | $\mu=0.53 \mathrm{~mm}^{-1}$ |
| $\beta=68.580(1)^{\circ}$ | $T=298(2) \mathrm{K}$ |
| $\gamma=69.340(1)^{\circ}$ | Block, colourless |
| $V=936.06(15) \AA^{\circ}$ | $0.22 \times 0.18 \times 0.15 \mathrm{~mm}$ |

## Data collection

Bruker SMART APEX areadetector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 2002)
$T_{\text {min }}=0.892, T_{\text {max }}=0.924$
5873 measured reflections

> 4627 independent reflections
> 3320 reflections with $I>2 \sigma(I)$
> $R_{\text {int }}=0.012$
> $\theta_{\max }=28.2^{\circ}$
> $h=-12 \rightarrow 12$
> $k=-12 \rightarrow 12$
> $l=-15 \rightarrow 9$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.050$
$w R\left(F^{2}\right)=0.135$
$S=1.03$
4267 reflections
253 parameters
H -atom parameters constrained

$$
\begin{gathered}
w=1 /\left[\sigma^{2}\left(F_{o}^{2}\right)+(0.0645 P)^{2}\right. \\
\quad+0.354 P] \\
\text { where } P=\left(F_{o}^{2}+2 F_{c}^{2}\right) / 3 \\
(\Delta / \sigma)_{\max }=0.001 \\
\Delta \rho_{\max }=0.33 \mathrm{e}^{-3} \AA^{-3} \\
\Delta \rho_{\min }=-0.41 \mathrm{e}^{-3}
\end{gathered}
$$

Table 1
Selected geometric parameters $\left(\AA^{\circ},^{\circ}\right)$.

| Cl1-C1 | $1.740(2)$ | $\mathrm{N} 3-\mathrm{C} 10$ | $1.332(3)$ |
| :--- | :--- | :--- | :--- |
| $\mathrm{F} 1-\mathrm{C} 18$ | $1.311(3)$ | $\mathrm{N} 4-\mathrm{C} 11$ | $1.133(3)$ |
| $\mathrm{N} 1-\mathrm{C} 7$ | $1.268(3)$ | $\mathrm{C} 4-\mathrm{C} 7$ | $1.453(3)$ |
| $\mathrm{N} 1-\mathrm{C} 8$ | $1.387(3)$ | $\mathrm{C} 8-\mathrm{C} 9$ | $1.368(3)$ |
| $\mathrm{N} 2-\mathrm{N} 3$ | $1.345(2)$ | $\mathrm{C} 9-\mathrm{C} 10$ | $1.392(3)$ |
| $\mathrm{N} 2-\mathrm{C} 8$ | $1.366(3)$ | $\mathrm{C} 10-\mathrm{C} 11$ | $1.437(3)$ |
| $\mathrm{N} 2-\mathrm{C} 12$ | $1.424(2)$ | $\mathrm{C} 15-\mathrm{C} 18$ | $1.507(3)$ |
|  |  |  |  |
| $\mathrm{C} 7-\mathrm{N} 1-\mathrm{C} 8$ | $118.03(18)$ | $\mathrm{C} 8-\mathrm{C} 9-\mathrm{C} 10$ | $104.57(19)$ |
| N3-N2-C8 | $113.56(16)$ | $\mathrm{N} 3-\mathrm{C} 10-\mathrm{C} 9$ | $113.45(19)$ |
| N3-N2-C12 | $122.26(17)$ | $\mathrm{N} 3-\mathrm{C} 10-\mathrm{C} 11$ | $120.3(2)$ |
| $\mathrm{C} 8-\mathrm{N} 2-\mathrm{C} 12$ | $123.83(17)$ | $\mathrm{C} 9-\mathrm{C} 10-\mathrm{C} 11$ | $126.2(2)$ |
| $\mathrm{C} 10-\mathrm{N} 3-\mathrm{N} 2$ | $102.62(17)$ | $\mathrm{N} 4-\mathrm{C} 11-\mathrm{C} 10$ | $176.5(3)$ |
| $\mathrm{N} 1-\mathrm{C} 7-\mathrm{C} 4$ | $123.2(2)$ | $\mathrm{C} 17-\mathrm{C} 12-\mathrm{N} 2$ | $119.97(18)$ |
| N2-C8-C9 | $105.78(18)$ | $\mathrm{F} 1-\mathrm{C} 18-\mathrm{F} 2$ | $107.3(3)$ |
| N2-C8-N1 | $118.46(18)$ | $\mathrm{F} 1-\mathrm{C} 18-\mathrm{C} 15$ | $112.6(2)$ |
| $\mathrm{C} 9-\mathrm{C} 8-\mathrm{N} 1$ | $135.8(2)$ |  |  |

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

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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