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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.005 Å R factor = 0.069 wR factor = 0.204 Data-to-parameter ratio = 12.8

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-[(2-furyl)methyleneamino]-1*H*-pyrazole-3-carbonitrile

The title compound, $C_{16}H_7Cl_2F_3N_4O$, is a tricyclic imide with an overall U-shape. There are $\pi - \pi$ interactions between the pyrazole and furyl rings.

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-3cyano-1-[2,6-dichloro-4-(trifluoromethyl)phenyl]-4-(trifluoromethylsulfenyl)pyrazole and 5-amino-3-cyano-1-[2,6dichloro-4-(trifluoromethyl)phenyl]-4-(trifluoromethylsulfonyl)pyrazole, which are all good insecticides (Hatton *et al.*, 1993).



The structure of (I) is shown in Fig. 1, with the atomnumbering scheme. The molecule contains three planar moieties, forming an overall U-shape. The dihedral angles between the pyrazole and the furyl and benzene rings are



Figure 1

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

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3013 independent reflections 2571 reflections with $I > 2\sigma(I)$

 $\begin{aligned} R_{\text{int}} &= 0.019\\ \theta_{\text{max}} &= 25.2^{\circ}\\ h &= -13 \rightarrow 14\\ k &= -8 \rightarrow 6\\ l &= -24 \rightarrow 25 \end{aligned}$



Figure 2 The packing of (I), viewed down the b axis.

19.8 (2) and 67.9 (1)°, respectively. The plane-to-plane separation of 3.8411 (1) Å between the furyl and pyrazole rings indicates the presence of a weak π - π interaction. In the crystal structure, the molecules are stacked along the *b* axis, as shown in Fig. 2.

Experimental

Following the method of Hatton *et al.* (1993), reaction of 2,6-dichloro-4-(trifluoromethyl)amine 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 2-furanal to give (I). Single crystals suitable for X-ray analysis were obtained by slow evaporation of a methanol solution (m.p. 449–451 K). IR (KBr, ν cm⁻¹): 3129, 2240, 1611, 1558, 1395, 1310, 1133, 873, 818; ¹H NMR (CDCl₃): δ 8.81 (*s*, 1H), 8.12 (*s*, 2H), 7.83 (*s*, 1H), 7.24 (*m*, 2H), 6.69 (*m*, 1H); ¹³C NMR (CDCl₃): 154.2 (1C), 153.3 (1C), 152.1 (1C), 149.2 (1C), 136.6 (1C), 134.4 (*q*, *J* = 34.3 Hz, 1C), 128.2 (1C), 127.05 (1C), 127.01 (1C), 126.95 (1C), 126.91 (1C), 123.3 (*q*, *J* = 271.6 Hz, 1C), 122.0 (1C), 114.2 (1C), 114.0 (1C), 98.4 (1C).

Crystal data

| $C_{16}H_7Cl_2F_3N_4O$ |
|--------------------------------|
| $M_r = 399.16$ |
| Monoclinic, $P2_1/n$ |
| a = 11.8828 (9) Å |
| b = 6.7072 (5) Å |
| c = 21.1191 (16) Å |
| $\beta = 92.084 \ (1)^{\circ}$ |
| $V = 1682.1 (2) \text{ Å}^3$ |
| Z = 4 |

D_x = 1.576 Mg m⁻³ Mo Kα radiation Cell parameters from 3298 reflections θ = 3.2–25.0° μ = 0.43 mm⁻¹ T = 298 (2) K Block, colorless 0.38 × 0.31 × 0.29 mm

Data collection

| Bruker APEX area-detector |
|--|
| diffractometer |
| φ and ω scans |
| Absorption correction: multi-scan |
| (SADABS; Bruker, 2002) |
| $T_{\min} = 0.854, \ T_{\max} = 0.885$ |
| 8571 measured reflections |
| Refinement |
| Refinement on F^2 |

| Refinement on F^2 | $w = 1/[\sigma^2(F_0^2) + (0.1164P)^2]$ |
|---------------------------------|--|
| $R[F^2 > 2\sigma(F^2)] = 0.069$ | + 2.0094P] |
| $wR(F^2) = 0.204$ | where $P = (F_0^2 + 2F_c^2)/3$ |
| S = 1.05 | $(\Delta/\sigma)_{\rm max} < 0.001$ |
| 3013 reflections | $\Delta \rho_{\rm max} = 1.12 \text{ e } \text{\AA}^{-3}$ |
| 235 parameters | $\Delta \rho_{\rm min} = -0.64 \text{ e } \text{\AA}^{-3}$ |
| H-atom parameters constrained | |

| Table 1 | | | | |
|----------|-----------|------------|-----|-----|
| Selected | geometric | parameters | (Å, | °). |

| Cl1-C4 | 1.724 (4) | N4-C12 | 1.273 (4) |
|------------|-----------|-------------|-----------|
| F1-C1 | 1.271 (9) | N4-C11 | 1.382 (4) |
| O1-C16 | 1.352 (4) | C9-C10 | 1.389 (5) |
| O1-C13 | 1.362 (4) | C10-C11 | 1.374 (5) |
| N1-N2 | 1.348 (4) | C13-C14 | 1.350 (5) |
| N1-C11 | 1.374 (4) | C14-C15 | 1.403 (5) |
| N2-C9 | 1.337 (5) | C15-C16 | 1.340 (6) |
| N3-C8 | 1.147 (5) | | |
| C16-O1-C13 | 106.2 (3) | C11-C10-C9 | 104.9 (3) |
| N2-N1-C11 | 113.0 (3) | C10-C11-N1 | 105.7 (3) |
| C9-N2-N1 | 103.2 (3) | C14-C13-O1 | 109.3 (3) |
| F3-C1-F2 | 111.0 (7) | C13-C14-C15 | 107.6 (3) |
| N3-C8-C9 | 179.1 (5) | C16-C15-C14 | 105.6 (3) |
| N2-C9-C10 | 113.1 (3) | C15-C16-O1 | 111.4 (3) |
| | | | |

All H atom were initially observed in a difference Fourier map and were then placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H = 0.93 Å and $U_{iso}(H) = 1.22_{eq}(C)$. The low U_{eq} value of atom C1 compared with its neighbours may be attributed to the three possibly disordered F atoms. The highest peak is located 1.27 Å from atoms C1 and F.

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: *SHELXL97*.

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