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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.049 wR factor = 0.126 Data-to-parameter ratio = 16.3

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

N,*N*'-[(2,3,5,6-Tetrafluoro-1,4-phenylene)dimethylene]bis(pyridine-4-carboxamide)

In the centrosymmetric molecule of the title compound, $C_{20}H_{14}F_4N_4O_2$, the pyridine ring is twisted with respect to the benzene ring with a dihedral angle of 66.85 (10)°. In the crystal structure, molecules are linked by intermolecular hydrogenbonding and $C-H\cdots\pi$ interactions.

Comment

Pyridinecarboxamide derivatives have attracted much attention because of their biological activity. Some derivatives of pyridinecarboxamide possess anticancer (Liu *et al.*, 1996) and anti-allergic activities (Tsuzurahara *et al.*, 1987), and are efficient as IKK inhibitors (Li & Prakash, 2005), monoamine oxidase B inhibitors (Blauenstein *et al.*, 1998) and anticonvulsant drugs (Palmer *et al.*, 1993). As part of a search for new pyridinecarboxamide compounds with better biological activity, the title compound, (I), has been synthesized. Here we report the crystal its structure.



The molecular structure of (I) is shown in Fig. 1. The molecule of (I) is centrosymmetric. The pyridine ring is twisted with respect to the benzene ring, with a dihedral angle of $66.85 (10)^{\circ}$.

Weak C-H··· π interactions are observed in the crystal structure (Fig. 2); C3-H3 = 0.93, H3···Cg^{iv} = 2.97, C3···Cg^{iv} = 3.726 (3) Å and C3-H3···Cg^{iv} = 139°; C5-H5 = 0.93, H5···Cg^v = 3.05, C5···Cg^v = 3.756 (3) Å and C5-H5···Cg^v = 133° [Cg is the centroid of the pyridine ring; symmetry codes: (iv) $\frac{5}{2} - x$, $\frac{1}{2} + y$, $\frac{3}{2} - z$; (v) $\frac{3}{2} - x$, $-\frac{1}{2} + y$, $\frac{3}{2} - z$]. Classical N-H···O and weak C-H···O and C-H···F hydrogen bonding occurs in the crystal structure (Table 1), which helps to stabilize the crystal structure (Fig. 3).

Experimental

To a stirred suspension of isonicotinyl chloride (3.92 g, 22 mmol) and 1,4-bis(aminomethyl)-2,3,5,6-tetrafluorobenzene (2.08 g, 10 mmol) in dry tetrahydrofuran (25 ml) was added dropwise triethylamine (3 ml) at 273 K. The mixture was maintained at room temperature for 48 h

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 $03716 \quad \text{He et al.} \cdot C_{20}H_{14}F_4N_4O_2$

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Figure 1

The molecular structure of (I), drawn with 30% probability ellipsoids (arbitrary spheres for H atoms) [symmetry code: (A) 1 - x, 1 - y, 2 - z].



Figure 2 The C-H··· π interactions (dashed lines) in (I).



Figure 3 A packing diagram for (I). Dashed lines indicate hydrogen bonds.

with stirring under a nitrogen atmosphere, then evaporated to dryness. The resulting residue was washed with iced water (25 ml), saturated sodium carbonate solution (12 ml), and again with iced water (15 ml). After work-up, the crude product was dried under vacuum and purified by recrystallization from methanol to yield (I) (3.81 g, 91% yield) as a white solid. Single crystals of (I) were obtained by slow evaporation of a methanol solution.

Z = 2

 $D_x = 1.567 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

 $\mu = 0.13 \text{ mm}^{-1}$

T = 298 (2) K

 $R_{\rm int} = 0.044$

 $\theta_{\rm max} = 28.5^{\circ}$

Block, colorless

 $0.30 \times 0.20 \times 0.13~\text{mm}$

2213 independent reflections

1302 reflections with $I > 2\sigma(I)$

Crystal data

 $C_{20}H_{14}F_4N_4O_2$ $M_r = 418.35$ Monoclinic, $P2_1/n$ a = 8.3488 (17) Åb = 5.1431 (11) Å c = 20.665 (4) Å $\beta = 91.843 \ (2)^{\circ}$ V = 886.9 (3) Å³

Data collection

Bruker SMART APEX CCD areadetector diffractometer φ and φ scans Absorption correction: none 7535 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0524P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.049$	+ 0.1461P]
$wR(F^2) = 0.126$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.03	$(\Delta/\sigma)_{\rm max} < 0.001$
2213 reflections	$\Delta \rho_{\rm max} = 0.18 \text{ e} \text{ \AA}^{-3}$
136 parameters	$\Delta \rho_{\rm min} = -0.21 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdots A$
$N2-H2A\cdots O1^{i}$	0.88	2.10	2.927 (2)	155
$C4 - H4 \cdots F2^{ii}$ $C7 - H7B \cdots O1^{iii}$	0.93 0.97	2.53 2.50	3.242 (3) 3.393 (3)	134 153
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Symmetry codes: (i) x, y - 1, z; (ii) $x + \frac{1}{2}, -y + \frac{1}{2}, z - \frac{1}{2}$; (iii) -x + 2, -y + 2, -z + 2.

The imino H atoms were located in a difference Fourier map and refined as riding in their as-found relative positions, with $U_{iso}(H) =$ $1.2U_{eq}(N)$. Other H atoms were placed in calculated positions, with C-H = 0.97 (methylene) or 0.93 Å (aromatic), and refined in riding mode, with $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Bruker, 2000); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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