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Key indicators

Single-crystal X-ray study T = 290 KMean σ (C–C) = 0.004 Å R factor = 0.082 wR factor = 0.137 Data-to-parameter ratio = 11.2

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

5-Benzyl-1-(4-fluorophenyl)-2-phenyl-4,5,6,7tetrahydro-1*H*-pyrrolo[3,2-c]pyridine

In the title compound, $C_{26}H_{23}FN_2$, the dihedral angle between the 4-fluorophenyl ring and the adjacent phenyl ring is $62.3 (1)^\circ$. The crystal structure is stabilized by $C-H\cdots\pi$ interactions. Received 10 February 2006 Accepted 22 March 2006.

Comment

Molecular organization and molecular interactions are the features that are responsible for molecules exhibiting different functional properties. An understanding of non-covalent interactions becomes essential for interpreting and predicting relationships between chemical structure and function (Hunter et al., 2001). Crystal engineering via the manipulation of hydrogen bonding has attracted much attention in the recent literature (Aakeröy, 1997; Desiraju, 2000). Weak intermolecular forces of the type $C-H \cdots \pi$ play an important role in various systems of biological and chemical interest (Nishio *et al.*, 2004). Intramolecular C–H··· π interactions are responsible for a molecule adopting a particular conformation in the solid state (Jennings et al., 2001). 4-Keto-4,5,6,7-tetrahydroindoles have been tested for anti-implantation agents in rats and have been found to exhibit potent biological activity. Against this background, we have studied the crystal structure of the title compound (I).



A molecular view of (I), with the atom-labelling scheme, is shown in Fig. 1. Relevant bond distances and torsion angles are given in Table 1. Atom N1 deviates by 0.604 (2) Å from

© 2006 International Union of Crystallography All rights reserved the least-squares plane passing through atoms C5/C3/C4/C7/ C6. The crystal structure is stabilized by $C-H\cdots\pi$ intermolecular interactions (Table 2), forming molecular dimers (Fig. 2).

Experimental

The compound (I) was synthesized according to the procedure reported in the literature (Nagarajan et al., 1985), and crystallized by slow evaporation of a chloroform/hexane (2:1 v/v) solution at 278 K.

 $D_x = 1.264 \text{ Mg m}^{-3}$

Cell parameters from 650

Mo $K\alpha$ radiation

reflections

 $\theta = 1.4 - 25.8^{\circ}$ $\mu = 0.08 \text{ mm}^{-1}$

T = 290 (2) K

Block, colourless

 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.17 \text{ e} \text{ Å}^{-3}$ $\Delta \rho_{\rm min}$ = -0.15 e Å⁻³

 $w = 1/[\sigma^2(F_0^2) + (0.0366P)^2]$ + 0.563P]

where $P = (F_0^2 + 2F_c^2)/3$

 $0.25 \times 0.20 \times 0.15 \text{ mm}$

Crystal data

C26H23FN2 $M_{\rm r} = 382.46$ Monoclinic, $P2_1/c$ a = 13.220 (8) Å b = 18.431 (11) Åc = 8.475 (5) Å $\beta = 103.223 (11)^{\circ}$ $V = 2010 (2) \text{ Å}^3$ Z = 4

Data collection

Bruker SMART APEX CCD area	3975 independent reflections
detector diffractometer	2935 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.043$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.4^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -16 \rightarrow 16$
$T_{\min} = 0.939, T_{\max} = 0.988$	$k = -22 \rightarrow 22$
15238 measured reflections	$l = -10 \rightarrow 10$

Refinement

Refinement on F^2	
$R[F^2 > 2\sigma(F^2)] = 0.082$	
$wR(F^2) = 0.137$	
S = 1.23	
3975 reflections	
354 parameters	
All H-atom parameters refined	

Table 1

Selected geometric parameters (Å, °).

N2-C4	1.392 (3)	N1-C20	1.461 (3)
N2-C1	1.393 (3)	N1-C5	1.467 (3)
N2-C8	1.431 (3)	N1-C6	1.473 (3)
F1-C11	1.356 (3)		
C5-N1-C20-C26	60.4 (3)	C20-N1-C6-C7	169.2 (2)
C6-N1-C20-C26	-176.4(3)	C5-N1-C6-C7	-67.5(3)
C19-C14-C1-C2	-144.7(3)		

Table 2

Hvarogen-pond geometry (A.	metrv (Å. °	geome	1-bond	vdrogen	H١
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$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C9-H9\cdots Cg1^{i}$	0.95 (3)	2.79 (3)	3.648 (4)	150 (2)

Symmetry code: (i) -x, -y + 1, -z + 1. Cg1 is the centroid of the five-membered ring C1-C4/N2

All H atoms were located in difference Fourier maps and refined isotropically. The C–H bond distances are in the range 0.92 (3) Å to 1.02 (2) Å.







Figure 2

A molecular dimer. Dashed lines indicate $C-H \cdot \cdot \pi$ interactions. Atoms labelled with an asterisk (*) are at the symmetry position (-x, 1 - y, 1 - y)1 - z).

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SIR92 (Altomare et al., 1993); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1999) and CAMERON (Watkin et al., 1993); software used to prepare material for publication: PLATON (Spek, 2003).

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