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Key indicators

Single-crystal X-ray study T = 292 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ R factor = 0.065 wR factor = 0.155 Data-to-parameter ratio = 15.1

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

1-Amino-2-anilino-4-(4-fluorobenzylidene)-1*H*imidazol-5(4*H*)-one

In the title compound, $C_{16}H_{13}FN_4O$, centrosymmetric hydrogen-bonded dimers are linked into one-dimensional chains along the [100] direction. The chains are further linked to give a sheet parallel to the (101) plane by $C-H\cdots F$ hydrogen bonds. Received 18 May 2006 Accepted 20 July 2006

Comment

Imidazol-4-one derivatives are important heterocycles, which have good fungicidal and antiphlogistic activities (Ding *et al.*, 2001; Trivedi & Shah, 1993). In order to find compounds presenting both low toxicity and high biological activity, we synthesized a series of new imidazol-4-ones containing 4fluorobenzene derivatives. In this context, we have crystallized the title compound, (I), and report its crystal structure here.



The molecular conformation is illustrated in Fig. 1. In the molecular structure, two benzene rings and the imidazole ring are almost coplanar; the torsion angles C6-C7-C8-N1 and N1-C10-N4-C12 are -1.3 (5) and 1.1 (4), respectively. All other molecular geometric features are unremarkable.



© 2006 International Union of Crystallography All rights reserved **Figure 1** Molecular structure of (I), showing displacement ellipsoids drawn at the 50% probability level. In the packing, the reference molecule at (x, y, z) is linked to the centrosymmetrically related molecule at (-x, -y, 2-z), forming a hydrogen-bonded dimer centered at (0,0,1). These dimers are linked into a two-dimensional sheet parallel to the (101) plane by N-H···O (or ···N) and C-H···F hydrogen bonds (Fig. 2 and Table 1). Analysis using *PLATON* (Spek, 2003) shows that no specific interaction is found between the adjacent layers.

Experimental

Compound (I) was synthesized according to the procedure of Ding *et al.* (2001). Suitable crystals were obtained by slow evaporation of an acetone solution at room temperature.

Z = 4

 $D_r = 1.418 \text{ Mg m}^{-3}$

 $0.20 \times 0.10 \times 0.04 \text{ mm}$

Mo $K\alpha$ radiation

 $\mu = 0.10 \text{ mm}^{-1}$

T = 292 (2) K

Block, purple

Crystal data

 $\begin{array}{l} C_{16}H_{13}FN_4O\\ M_r = 296.30\\ Monoclinic, P2_1/n\\ a = 5.5218 \ (5) \ \AA\\ b = 25.529 \ (2) \ \AA\\ c = 10.0165 \ (9) \ \AA\\ \beta = 100.553 \ (2)^\circ\\ V = 1388.1 \ (2) \ \AA^3 \end{array}$

Data collection

Bruker SMART APEX CCD areadetector diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 2001) $T_{min} = 0.980, T_{max} = 0.996$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.065$ $wR(F^2) = 0.155$ S = 0.983144 reflections 208 parameters 9440 measured reflections 3144 independent reflections 1654 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.057$ $\theta_{\text{max}} = 27.5^{\circ}$

H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0624P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.18 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.16 \text{ e} \text{ Å}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C13-H13\cdots F1^{i}$ $N3-H3B\cdots O1^{ii}$ $N3-H3C\cdots N1^{iii}$	0.93 0.862 (10) 0.866 (10)	2.49 2.52 (3) 2.413 (11)	3.348 (3) 2.973 (3) 3.279 (3)	154 114 (2) 178 (3)
$N4-H4A\cdots N3$	0.862 (10)	2.28 (2)	2.762 (3)	115 (2)

Symmetry codes: (i) x - 1, y, z + 1; (ii) -x, -y, -z + 2; (iii) x - 1, y, z.

All C-bound H atoms were placed in idealized locations and were refined using a riding model, with C–H = 0.93 Å and $U_{\rm iso}({\rm H})$ =



Figure 2

Packing of the molecules, with hydrogen bonds shown as dashed lines. Cbound H atoms have been omitted.

 $1.2U_{eq}(C)$. Atoms H3B, H3C and H4A were located in difference maps and refined with the restraints N-H = 0.86 (1) Å and $U_{iso}(H) = 1.2U_{eq}(N)$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

References

- Bruker (2001). SAINT (Version 6.45) and SMART (Version 5.628). Bruker AXS Inc., Madison, Wisconsin, USA.
- Ding, M.-W., Xu, Z.-F., Liu, Z.-J. & Wu, T.-J. (2001). Synth. Commun. 31, 1053– 1057.
- Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.
- Sheldrick, G. M. (2001). SADABS. Version 2.10. Bruker AXS Inc., Madison, Wisconsin, USA.
- Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.
- Trivedi, B. & Shah, V. H. (1993). J. Indian Chem. Soc. 70, 645-648.