

## 6-(4-Fluorophenyl)-5,6-dihydrobenzimidazo[1,2-c]quinazoline

John Nicolson Low,<sup>a\*†</sup> Braulio Insuasty,<sup>b</sup> Harlen Torres<sup>b</sup> and Justo Cobo<sup>c</sup>

<sup>a</sup>Department of Chemistry, University of Aberdeen, Meston Walk, Old Aberdeen AB24 3UE, Scotland, <sup>b</sup>Grupo de Investigación de Compuestos Heterocíclicos, Departamento de Química, Universidad de Valle, AA 25360, Cali, Colombia, and <sup>c</sup>Departamento de Química Inorgánica y Orgánica, Universidad de Jaén, 23071 Jaén, Spain

† Postal address: Department of Electrical Engineering and Physics, University of Dundee, Dundee DD1 4HN, Scotland

Correspondence e-mail: j.n.low@abdn.ac.uk

## Key indicators

Single-crystal X-ray study

$T = 120\text{ K}$

Mean  $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$

$R$  factor = 0.057

w $R$  factor = 0.162

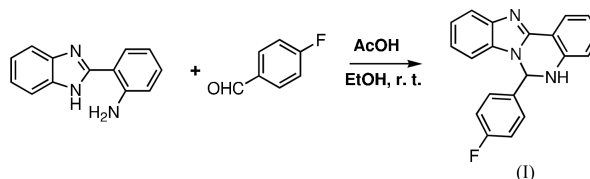
Data-to-parameter ratio = 17.3

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

The title compound,  $\text{C}_{20}\text{H}_{14}\text{FN}_3$ , contains one conventional  $\text{N}-\text{H}\cdots\text{N}$  hydrogen bond which links the molecules into spiral chains running parallel to the  $c$  axis; the conformation of this chain appears to be stabilized by an antiparallel  $\text{C}-\text{H}\cdots\text{F}$  contact. The structure also contains non-localized solvent in isolated cavities which lie along the crystallographic threefold axis.

## Comment

Due to the increasing number of immunocompromized individuals, fungal infections have increased steadily in the last two decades, affecting millions of people worldwide. Since compounds containing the quinazoline skeleton have shown interesting antifungal properties (Chan *et al.*, 1995; Castaldo *et al.*, 1979; Bartroli *et al.*, 1998) this current project is aimed at developing synthetic strategies for preparing heterocyclic compounds containing the quinazoline ring system. The title compound was prepared from 2-(1*H*-benzimidazol-2-yl)-aniline and 4-fluorobenzaldehyde.



The title compound, (I), contains no unusual bond lengths or angles (Fig. 1). The supramolecular structure is determined by two direction-specific interactions which link the molecules together into spirals running parallel to the  $c$  axis. These interactions are a conventional  $\text{N}-\text{H}\cdots\text{N}$  hydrogen bond and a short  $\text{C}-\text{H}\cdots\text{F}$  contact (Table 1). Atom N1 in the molecule at  $(x, y, z)$  forms, *via* H1, a hydrogen bond to N7 in the molecule at  $(\frac{2}{3} - x + y, \frac{1}{3} - x, \frac{1}{3} + z)$ , so forming a  $C(6)$  (Bernstein *et al.*, 1995) spiral chain which runs parallel to the  $c$  axis. Atom C8 in this latter molecule forms a contact *via* H8 to atom F24 in the molecule at  $(\frac{1}{3} - y, x - y - \frac{1}{3}, z - \frac{1}{3})$ , forming a  $C(11)$  chain. Atom N1 in this molecule, *via* H1, links to N7 in the molecule at  $(x, y, z)$ , forming an  $R_3^3(19)$  ring. Fig. 2 shows a stereoview of the ring structure running along the  $c$  axis.

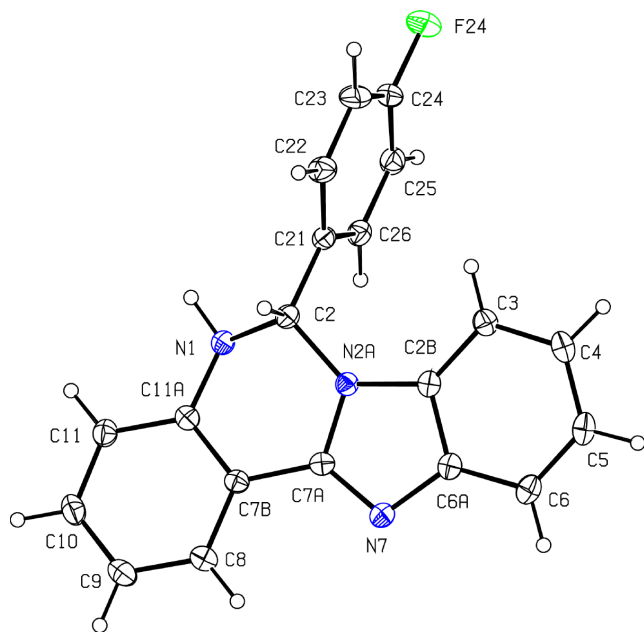
## Experimental

A solution of 2-(1*H*-benzimidazol-2-yl)aniline and 4-fluorobenzaldehyde (16 mmol) in ethanol (10 ml) was treated with acetic acid (1 ml). The mixture was stirred at room temperature for 10 min and then allowed to crystallize in a refrigerator overnight. The resulting solid was collected and recrystallized from ethanol, affording yellow crystals suitable for X-ray analysis (m.p. 477 K, yield 85%). Analysis

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**Figure 1**  
A view of (I) with the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

calculated for  $C_{20}H_{14}FN_3$ : C 76.18, H 4.47, N 13.33%; found: C 76.31, H 4.41, N 13.21%.

#### Crystal data

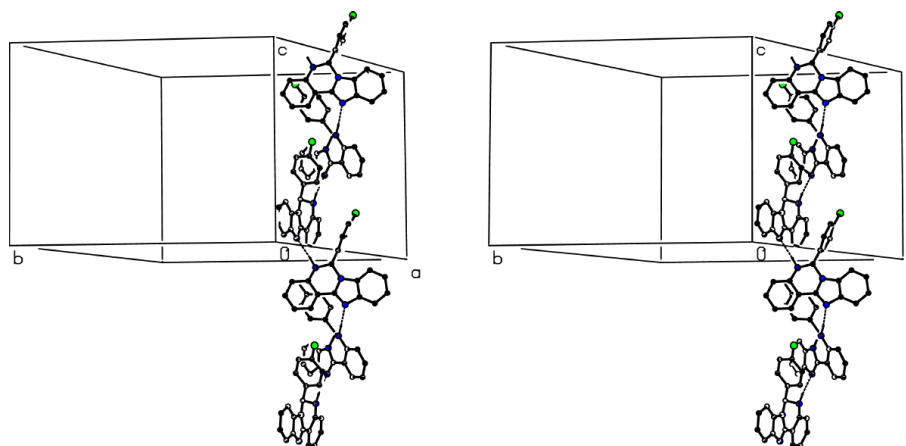
$C_{20}H_{14}FN_3$   
 $M_r = 315.34$   
 Trigonal (hexagonal axes),  $R\bar{3}$   
 $a = 22.3211$  (8) Å  
 $c = 17.2220$  (9) Å  
 $V = 7431.0$  (5) Å<sup>3</sup>  
 $Z = 18$   
 $D_x = 1.268$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation  
 Cell parameters from 3780  
 reflections  
 $\theta = 3.0$ – $27.5^\circ$   
 $\mu = 0.09$  mm<sup>-1</sup>  
 $T = 120.0$  (2) K  
 Plate, colourless  
 $0.26 \times 0.20 \times 0.06$  mm

#### Data collection

Nonius KappaCCD diffractometer  
 $\varphi$  scans and  $\omega$  scans with  $\kappa$  offsets  
 Absorption correction: multi-scan  
 (DENZO-SMN; Otwinowski &  
 Minor, 1997)  
 $T_{\min} = 0.978$ ,  $T_{\max} = 0.995$   
 28120 measured reflections

3780 independent reflections  
 2552 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.105$   
 $\theta_{\text{max}} = 27.5^\circ$   
 $h = -28 \rightarrow 28$   
 $k = -28 \rightarrow 28$   
 $l = -19 \rightarrow 22$



**Figure 2**  
A stereoview of the  $R_3^3(19)$  ring formed by the combination of the C(6) and C(11) chains.

#### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.057$   
 $wR(F^2) = 0.162$   
 $S = 1.04$   
 3780 reflections  
 218 parameters  
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0894P)^2 + 1.7404P]$$

where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.27$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.29$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bonding geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1\cdots N7^i$	0.93	2.09	3.016 (3)	173
$C8-H8\cdots F24^{ii}$	0.95	2.49	3.274 (2)	140

Symmetry codes: (i)  $\frac{2}{3} - x + y, \frac{1}{3} - x, \frac{1}{3} + z$ ; (ii)  $\frac{2}{3} - x + y, \frac{1}{3} - x, z - \frac{2}{3}$ .

The structure was refined using hexagonal axes. H atoms were treated as riding atoms, with C–H = 0.95 Å (aromatic), 0.98 Å (methyl) and 0.99 Å (CH<sub>2</sub>). The H atom attached to N1 was allowed to ride at a position based on its location in a difference map. The difference map also revealed several peaks lying in voids on and around the threefold axis. These could not be resolved in terms of sensible molecules and the SQUEEZE option in PLATON (Spek, 2003) was used to remove the contribution of electron density from these solvent-accessible voids.

Data collection: KappaCCD Server Software (Nonius, 1997); cell refinement: DENZO-SMN (Otwinowski & Minor, 1997); data reduction: DENZO-SMN; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976) and PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97 and WordPerfect macro PRPKAPPA (Ferguson, 1999).

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