

Acta Crystallographica Section E

# **Structure Reports Online**

ISSN 1600-5368

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#### **Key indicators**

Single-crystal X-ray study T = 173 KMean  $\sigma(\text{C-C}) = 0.003 \text{ Å}$  R factor = 0.039 wR factor = 0.097Data-to-parameter ratio = 11.9

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

# 1-(4-Fluorophenyl)-1,3-dihydroisobenzofuran-5-carbonitrile

The title compound,  $C_{15}H_{10}FNO$ , crystallizes with two molecules in the asymmetric unit, which differ only in the orientation of the fluorophenyl ring with respect to the isobenzofuran system.

Received 15 October 2004 Accepted 27 October 2004 Online 6 November 2004

# Comment

1,3-Dihydroisobenzofuran (or phthalan) is used for the preparation of 1,2-di(lithiomethyl)benzene (Almena *et al.*, 1995). Electron-transfer-induced reductive cleavage of phthalans has been reported (Azzena *et al.*, 1996). The title compound, (I), 4-fluorophenyl-5-phthalan carbonitrile, is a key intermediate in the synthesis of citalopram, which is a versatile antidepressant (Liechti *et al.*, 2000). In order to examine the conformation of the phthalan moiety and to study the influence of fluorophenyl and cyano groups, the crystal structure determination of (I) has been carried out and the results are presented here.

Compound (I) crystallizes with two molecules in the asymmetric unit, and perspective views of these are shown in Figs. 1 and 2. Bond lengths and angles can be regarded as normal (Cambridge Structural Database, Version 1.6 plus three updates; *MOGUL* Version 1.0; Allen, 2002). The isobenzofuran moiety is essentially planar (r.m.s. deviations 0.020 and 0.022 Å for the two molecules). Bond lengths and angles in the two different molecules are essentially equal. The two molecules differ only in the orientation of the fluorophenyl ring with respect to the isobenzofuran system. This difference can be expressed by the corresponding torsion angles (Table 1), which differ by approximately 50°. A least-squares fit of the isobenzofuran moieties including the cyano group (Fig. 3) gives an r.m.s. deviation of 0.015 Å.

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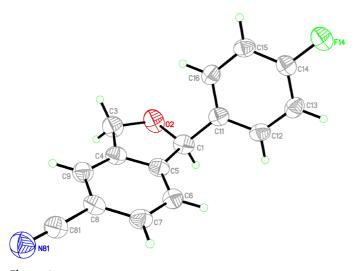


Figure 1

A perspective view of molecule 1 of the two molecules in the asymmetric unit of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

There are no classical hydrogen bonds in the crystal structure of (I), but several short  $C-H\cdots N$  and  $C-H\cdots F$  contacts are observed (Table 2).

# **Experimental**

5-Bromo-3H-isobenzofuran-1-one (1.0 g ,4.7 mmol) was subjected to a Grignard reaction with 4-fluorophenyl magnesium bromide (1.12 g, 5.6 mmol) in tetrahydrofuran (20 ml) at 273–278 K. The resulting product was treated with sodium borohydride (0.19 g ,5.2 mmol) in methanol (5 ml) to obtain the diol, which was cyclized with p-toluene sulfonic acid monohydrate (0.1 g, 0.5 mmol) in toluene (10 ml) to obtain the cyclized product. The cyclized product was refluxed with CuCN (0.5 g, 5.6 mmol) in dimethylformamide (5 ml) to obtain the title compound (Bigler et al., 1977). X-ray quality crystals of (I) were obtained after recrystallization from solution in acetonitrile.  $Crystal\ data$ 

$C_{15}H_{10}FNO$	$D_x = 1.404 \text{ Mg m}^{-3}$
$M_r = 239.24$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 8495
a = 14.606 (2)  Å	reflections
b = 5.8056 (6) Å	$\theta = 2.8 – 25.1^{\circ}$
c = 27.037 (4)  Å	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 99.032 (11)^{\circ}$	T = 173 (2)  K
$V = 2264.2 (5) \text{ Å}^3$	Block, light yellow
Z = 8	$0.47 \times 0.42 \times 0.36 \text{ mm}$

## Data collection

Stoe IPDS II two-circle	2571 reflections with $I > 2\sigma(I)$
diffractometer	$R_{\rm int} = 0.045$
$\omega$ scans	$\theta_{\rm max} = 25.2^{\circ}$
Absorption correction: none	$h = -17 \rightarrow 15$
10 504 measured reflections	$k = -6 \rightarrow 6$
3878 independent reflections	$l = -30 \rightarrow 32$

# Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.039$	$w = 1/[\sigma^{2}(F_{o}^{2}) + (0.059P)^{2}]$
$wR(F^2) = 0.097$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.88	$(\Delta/\sigma)_{\text{max}} = 0.001$
3878 reflections	$\Delta \rho_{\text{max}} = 0.14 \text{ e Å}^{-3}$
325 parameters	$\Delta \rho_{\min} = -0.20 \text{ e Å}^{-3}$

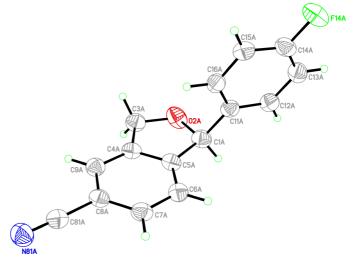


Figure 2
A perspective view of molecule 2 of the two molecules in the asymmetric unit of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

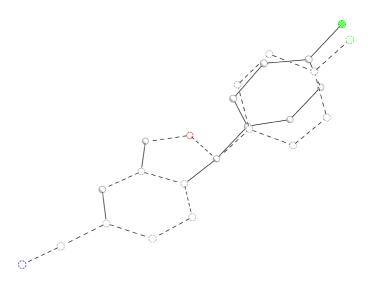


Figure 3
A least-squares fit of the isobenzofuran moieties of the asymmetric unit of (I).

Table 1 Selected geometric parameters  $(\mathring{A}, {}^{\circ})$ .

C1-O2	1.432(2)	C1A - O2A	1.441(2)
O2-C3	1.436 (2)	O2A - C3A	1.430(2)
C8-C81	1.451 (3)	C8A - C81A	1.453 (3)
C81-N81	1.131 (2)	C81A-N81A	1.129(2)
C14-F14	1.361 (2)	C14A – F14A	1.359 (2)
O2-C1-C11-C12	-149.46 (15)	O2A-C1A-C11A-	C12A-96.76 (18)
O2-C1-C11-C16	33.5 (2)	O2A - C1A - C11A -	C16A 80.41 (19)

**Table 2** Geometry of hydrogen bonds and weak C−H···F interactions (Å, °).

$D-H\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
C7−H7···N81A <sup>i</sup>	0.95	2.60	3.472 (3)	152
C7 <i>A</i> − H7 <i>A</i> ···N81 <sup>i</sup>	0.95	2.62	3.511 (3)	156
C9−H9···N81 <sup>ii</sup>	0.95	2.56	3.484 (2)	166
C9A−H9A···N81A <sup>iii</sup>	0.95	2.63	3.479 (2)	148
$C6-H6\cdots F14A^{iv}$	0.95	2.80	3.750(2)	177
C3−H3 <i>B</i> ···F14 <sup>v</sup>	0.99	2.69	3.223 (2)	114
$C6A - H6A \cdot \cdot \cdot F14^{iv}$	0.95	2.55	3.491 (2)	170
$C16A - H16A \cdot \cdot \cdot F14^{vi}$	0.95	2.80	3.611 (2)	144

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

H atoms were refined with fixed individual isotropic displacement parameters [ $U_{\rm iso}({\rm H})=1.2U_{\rm eq}({\rm C})$ ], using a riding model, with C-H = 1.00, 0.99 and 0.95 Å for tertiary CH, secondary CH and aromatic CH, respectively.

Data collection: X-AREA (Stoe & Cie, 2001); cell refinement: X-AREA; data reduction: X-AREA; program(s) used to solve

structure: *SHELXS*97 (Sheldrick, 1990); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL-Plus* (Sheldrick, 1991); software used to prepare material for publication: *SHELXL*97 and *PLATON* (Spek, 1990).

One of the authors (BN) is grateful to University of Mysore for laboratory facilities.

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