

# Ethyl 2-amino-4-(4-fluorophenyl)-7,7-dimethyl-5-oxo-5,6,7,8-tetrahydro-4*H*-benzo[*b*]pyran-3-carboxylate

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## Key indicators

Single-crystal X-ray study

$T = 296\text{ K}$

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

$R$  factor = 0.040

$wR$  factor = 0.114

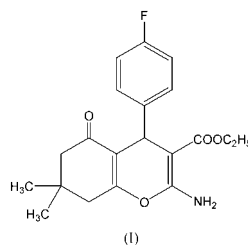
Data-to-parameter ratio = 13.4

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

The title compound,  $\text{C}_{20}\text{H}_{22}\text{FNO}_4$ , was synthesized by the reaction of 4-fluorobenzaldehyde with 5,5-dimethyl-1,3-cyclohexanedione and ethyl cyanoacetate. The pyran ring adopts a boat conformation, while the fused six-membered ring adopts a distorted boat conformation.

## Comment

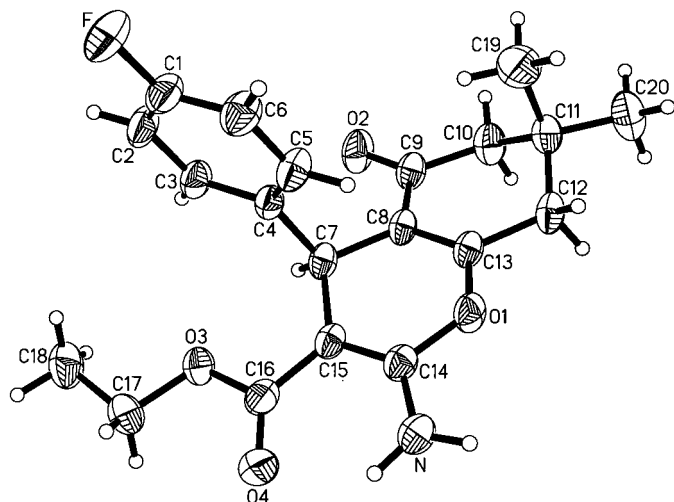
Benzopyrans possess a wide range of pharmacological activities (Bloxham *et al.*, 1994; Nawwar *et al.*, 1991; Zamocka *et al.*, 1991). The use of water as a solvent in organic chemistry was rediscovered in the 1980s (Breslow & Rideout, 1980). As part of our program aimed at developing new selective and environmentally friendly methodologies for the preparation of fine chemicals (Shi *et al.*, 2003), we carried out a reaction employing water as the reaction medium. We report here the X-ray crystal structure of the title compound, (I).



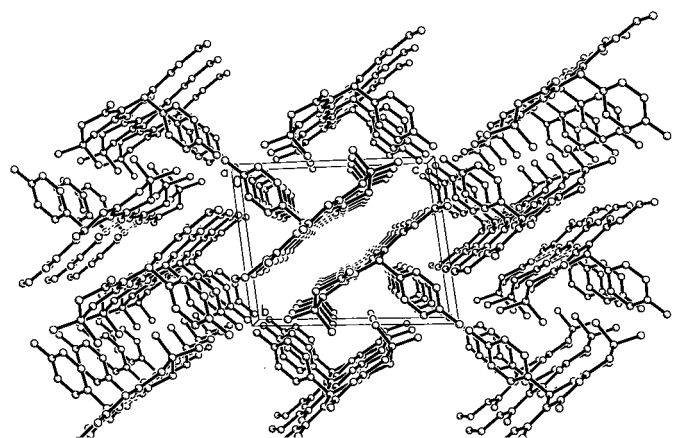
The pyran ring adopts a boat conformation (Fig. 1 and Table 1); atoms C8, C13, C14 and C15 are coplanar, with atoms O1 and C7 deviating from the plane by 0.120 (1) and 0.258 (1) Å, respectively. A similar distortion was observed in the structure of 2-amino-7,7-dimethyl-4-(3-fluorophenyl)-5-oxo-3-ethoxycarbonyl-5,6,7,8-tetrahydro-4*H*-benzo(*b*)pyran (Sharanina *et al.*, 1986). In addition, the N—C14 bond length of 1.330 (2) Å is shorter than the typical  $\text{Csp}^2\text{—N}$  distance (Lorente *et al.*, 1995). The fused six-membered ring adopts a distorted boat conformation; atoms C8, C9, C10 and C13 are coplanar, while atoms C11 and C12 deviate from the plane by 0.761 (1) and 0.212 (1) Å, respectively. Molecules are linked by N—H...O hydrogen bonds (Table 2 and Fig. 2).

## Experimental

The title compound, (I) (m.p. 433–434 K), was prepared by the reaction of 4-fluorobenzaldehyde with 5,5-dimethyl-1,3-cyclohexanedione and ethyl cyanoacetate in the presence of triethylbenzylammonium chloride in water. Single crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of an ethanol solution.



**Figure 1**  
The molecular structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme.



**Figure 2**  
The molecular packing in the crystal structure of (I).

#### Crystal data

$C_{20}H_{22}FNO_4$   
 $M_r = 359.39$   
 Triclinic,  $P\bar{1}$   
 $a = 8.266$  (1) Å  
 $b = 9.773$  (1) Å  
 $c = 12.375$  (2) Å  
 $\alpha = 81.40$  (1)°  
 $\beta = 73.41$  (1)°  
 $\gamma = 82.15$  (1)°  
 $V = 942.6$  (2) Å<sup>3</sup>

$Z = 2$   
 $D_x = 1.266$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 32 reflections  
 $\theta = 3.1$ – $15.1$ °  
 $\mu = 0.09$  mm<sup>-1</sup>  
 $T = 296$  (2) K  
 Block, colorless  
 $0.50 \times 0.46 \times 0.40$  mm

#### Data collection

Siemens P4 diffractometer  
 $\omega$  scans  
 Absorption correction:  $\psi$  scan  
 (XSCANS; Siemens, 1994)  
 $T_{\min} = 0.937$ ,  $T_{\max} = 0.964$   
 3694 measured reflections  
 3316 independent reflections  
 2258 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.008$   
 $\theta_{\text{max}} = 25.0$ °  
 $h = 0 \rightarrow 9$   
 $k = -11 \rightarrow 11$   
 $l = -14 \rightarrow 14$   
 3 standard reflections every 97 reflections  
 intensity decay: 2.3%

#### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.040$   
 $wR(F^2) = 0.114$   
 $S = 1.00$   
 3316 reflections  
 247 parameters  
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0652P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.20$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.19$  e Å<sup>-3</sup>  
 Extinction correction: SHELXTL  
 Extinction coefficient: 0.040 (4)

**Table 1**

Selected geometric parameters (Å, °).

O1—C13	1.3686 (19)	C7—C15	1.514 (2)
O1—C14	1.3770 (19)	C8—C13	1.336 (2)
N—C14	1.330 (2)	C14—C15	1.359 (2)
C7—C8	1.505 (2)		
C13—O1—C14	118.45 (12)	C15—C14—O1	122.24 (15)
N—C14—O1	109.95 (15)	C14—C15—C7	120.74 (15)
C15—C7—C8—C9	159.77 (13)	N—C14—C15—C16	-1.7 (3)
C13—C8—C9—C10	3.3 (2)	C4—C7—C15—C14	-103.32 (17)
C11—C12—C13—C8	15.1 (2)		

**Table 2**

Hydrogen-bonding geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N—H1B $\cdots$ O2 <sup>i</sup>	0.87 (2)	2.03 (2)	2.889 (2)	170.6 (19)
N—H1A $\cdots$ O4	0.90 (2)	2.04 (2)	2.688 (2)	128.1 (18)

Symmetry code: (i)  $1 + x, y, z$ .

The amino H1A and H1B atoms were refined isotropically. The positions of the other H atoms were fixed and distances to H atoms were set by SHELXTL (Sheldrick, 1997) [ $C-H = 0.93$ – $0.98$  Å and  $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$ ].

Data collection: XSCANS (Siemens, 1994); cell refinement: XSCANS; data reduction: SHELXTL (Sheldrick, 1997); program(s) used to solve structure: SHELXTL; program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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