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5-Fluoroindoline-2,3-dione

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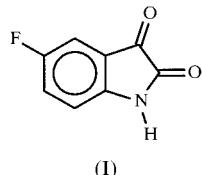
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In the title compound (5-fluoroisatin), $C_8H_4FNO_2$, the N atom of one molecule is linked to the amido-O atom of an adjacent molecule across a center of symmetry by a cyclic hydrogen bond [$N \cdots O = 2.888 (4) \text{ \AA}$].

Comment

Substituted isatin (2,3-indolinedione) compounds display biological (Anastasova, 1997) and cytostatic (Anastasova *et al.*, 2000; Juranic *et al.*, 1990) activity. QSAR studies have correlated the cytostatic activity of the monosubstituted isatins (including 5-fluoroisatin) to the nature and position of the substituent in the molecule. The fluorine substituent at the 5-position enhances antiproliferative activity; for the related uracil systems, the 5-fluoro compounds are well known anti-tumor agents (Bretner *et al.*, 1993). The crystal structure of isatin has been well investigated (Frolova *et al.*, 1988; Goldschmidt & Llewellyn, 1950; Palenik *et al.*, 1990; Palmer *et al.*, 1987); the crystal structures of monosubstituted isatins have not been reported, other than for the *N*-acetyl derivative (Zukerman-Schpector *et al.*, 1992); the molecular structures of the title compound, (I), and other 5- and *N*-substituted isatins have been studied by spectroscopic methods (Morales-Rios & Joseph-Nathan, 1991; Naumov & Anastasova, 2000).



The 5-fluoroisatin molecule is planar and all bond distances and angles are normal, except for the elongated $C(=O) - C(=O)$ single bond of $1.554 (4) \text{ \AA}$. Such long bonds are, aside from a number of other reasons (Palmer *et al.*, 1987), attributed to non-bonded lone-pair/lone-pair repulsions (Palenik *et al.*, 1990). A shorter bond [$1.538 (2) \text{ \AA}$] is found in the *N*-ethyl

derivative (Zukerman-Schpector *et al.*, 1992). In the F-substituted derivative, the amido-N atom of one molecule is linked to the amido-O atom of an adjacent molecule across a center of symmetry [$N \cdots O 2.888 (4) \text{ \AA}$].

The *ab initio* [HF/6-31G(d,p)] theoretical structure of 5-fluoroisatin reveals a slightly longer N—C(aromatic) bond; otherwise, the molecular structure is insignificantly different from that of isatin itself. Intermolecular (*e.g.* hydrogen bonding) rather than intrinsic factors probably account for the slightly longer 3-carbonyl bond in isatin [2-carbonyl = $1.220 (3) \text{ \AA}$ and 3-carbonyl = $1.211 (3) \text{ \AA}$ (Frolova *et al.*, 1988); 2-carbonyl = $1.220 (3) \text{ \AA}$ and 3-carbonyl = $1.213 (3) \text{ \AA}$ (Palenik *et al.*, 1990); 2-carbonyl = $1.220 (2) \text{ \AA}$ and 3-carbonyl = $1.214 (2) \text{ \AA}$ (Palmer *et al.*, 1987)] and 5-fluoroisatin [3-carbonyl = $1.222 (3) \text{ \AA}$ and 2-carbonyl = $1.211 (3) \text{ \AA}$]. These are indistinguishable in the optimized structures (isatin: 1.184 and 1.183 \AA ; 5-fluoroisatin: 1.183 and 1.182 \AA). The acetyl group in *N*-acetylisatin (Zukerman-Schpector *et al.*, 1992) hinders the formation of isatin-like cetrosymmetric dimers, which explains the reverse order of bond lengths [2-carbonyl = $1.196 (2) \text{ \AA}$ and 3-carbonyl = $1.207 (2) \text{ \AA}$]. The long $Csp^2 - Csp^2$ single bond in isatin and its derivatives allows for the ready cleavage of the five-membered ring in acidic or alkaline media, the reaction can explain the biological activity. *Ab initio* treatment (isatin: 1.555 \AA ; 5-F isatin: 1.553 \AA) supports the insensitivity of this bond to substitution of the hydrogen by fluorine.

Experimental

5-Fluoroisatin was obtained by cyclization of the fluorinated iso-nitroso precursor; the synthesis will be detailed elsewhere (Anastasova *et al.*, 2000). Theoretical calculations: the molecular geometry of 5-fluoroisatin was first CNDO-optimized with HYPERCHEM (Hyperecube Inc., 1996) and then further optimized at the HF SCF level within the standard basis set 6-31G(d,p) by the analytical gradient method in the GAUSSIAN-94w package (Frisch *et al.*, 1995). Harmonic vibrational analyses at the same level were performed on the resulting stationary points to ascertain energy minima.

Crystal data

$C_8H_4FNO_2$	$D_x = 1.587 \text{ Mg m}^{-3}$
$M_r = 165.12$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 2310 reflections
$a = 3.789 (7) \text{ \AA}$	$\theta = 2.15 - 25.97^\circ$
$b = 12.20 (1) \text{ \AA}$	$\mu = 0.134 \text{ mm}^{-1}$
$c = 14.99 (2) \text{ \AA}$	$T = 298 (2) \text{ K}$
$\beta = 94.41 (1)^\circ$	Parallelepiped, brown
$V = 691 (2) \text{ \AA}^3$	$0.30 \times 0.25 \times 0.25 \text{ mm}$
$Z = 4$	

Data collection

MarResearch Image Plate diffractometer	$R_{\text{int}} = 0.034$
Method: 95 frames at 2° intervals,	$\theta_{\text{max}} = 25.97^\circ$
counting time 2 min.	$h = 0 \rightarrow 4$
2310 measured reflections	$k = -15 \rightarrow 14$
1265 independent reflections	$l = -18 \rightarrow 18$
851 reflections with $I > 2\sigma(I)$	Intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.166$
 $S = 1.074$
1265 reflections
110 parameters
H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0891P)^2 + 0.1412P]$$

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

$$(\Delta/\sigma)_{\max} < 0.001$$

$$\Delta\rho_{\max} = 0.20 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$$

Extinction correction: *SHELXL97*
Extinction coefficient: 0.04 (1)

Table 1

Hydrogen-bonding geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1 ⁱ —O2 ⁱ	0.86	2.06	2.888 (4)	163

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

Data collection: *XDS* (Kabsch, 1988); cell refinement: *XDS*; data reduction: *XDS*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); software used to prepare material for publication: *SHELXL97*.

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