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## 2-(1,3-Dioxo-4,5,6,7-tetrahydro-1Hisoindol-2-yl)-N-[7-fluoro-3-oxo-4-(prop-2-ynyl)-3,4-dihydro-2H-benzoxazin-6-yl]acetamide monohydrate

In the title compound,  $C_{21}H_{18}FN_3O_5 H_2O$ , the cyclohexene ring exhibits a distorted chair conformation. The crystal packing is stabilized by intra- and intermolecular hydrogen bonds.

# Comment

Herbicides inhibiting protoporphyrinogen oxidase (protox) have been sold commercially for nearly 40 years (Dayan & Duke, 1997). The title compound, (I), may belong to this family of protox-inhibiting herbicides and we present its crystal structure here.





© 2006 International Union of Crystallography The molecular structure of (I), showing displacement ellipsoids drawn at the 50% probability level. All disorder components are shown.



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#### Kev indicators

Single-crystal X-ray study T = 292 K Mean  $\sigma$ (C–C) = 0.004 Å Disorder in main residue R factor = 0.062 wR factor = 0.167 Data-to-parameter ratio = 11.3

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



#### Figure 2

The packing of (I), with hydrogen bonds shown as dashed lines. Only one component of each disordered group is shown.

Therefore, atoms N1, N2 and N3 are  $sp^2$  hybridized. The molecules of (I) form two-dimensional layers through hydrogen bonds in the ac plane (Table 2 and Fig. 2).

#### **Experimental**

2-[1,3-Dioxo-4,5,6,7-tetrahydro-1*H*-isoindol-2-yl]acetyl choride (1.2 mmol) in dry toluene (10 ml) was added dropwise to a solution of 6-amino-7-fluoro-4-(prop-2-ynyl)-2H-benzoxazin-3(4H)-one (1 mmol) and triethylamine (1.2 mmol) in dry toluene (10 ml) under  $N_2$  at room temperature, and the resulting mixture was stirred for 2 h. After filtration, the solid was washed with water and recrystallized from petroleum ether and methanol (4:1 v/v). Colorless plate-shaped crystals of (I) were obtained by evaporation of the solvent over a period of two weeks.

#### Crystal data

$C_{21}H_{18}FN_3O_5 \cdot H_2O$	$D_x = 1.339 \text{ Mg m}^{-3}$
$M_r = 429.40$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 2160
a = 19.2192 (19)  Å	reflections
b = 4.7354 (5) Å	$\theta = 2.7 – 21.4^{\circ}$
c = 23.421 (2) Å	$\mu = 0.11 \text{ mm}^{-1}$
$\beta = 92.091 \ (2)^{\circ}$	T = 292 (2) K
V = 2130.2 (4) Å <sup>3</sup>	Plate, colorless
Z = 4	$0.40 \times 0.10 \times 0.02 \text{ mm}$

#### Data collection

Bruker SMART CCD area-detector	37
diffractometer	24
$\varphi$ and $\omega$ scans	$R_{\rm ir}$
Absorption correction: multi-scan	$\theta_{\rm m}$
(SADABS; Bruker, 2000)	h =
$T_{\min} = 0.959, \ T_{\max} = 0.998$	<i>k</i> =
14145 measured reflections	<i>l</i> =

#### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.062$  $wR(F^2) = 0.167$ S = 1.043719 reflections 328 parameters H atoms treated by a mixture of independent and constrained refinement

19 independent reflections 11 reflections with  $I > 2\sigma(I)$  $n_{\rm nt} = 0.050$  $_{\rm max} = 25.0^{\circ}$  $= -21 \rightarrow 22$  $= -5 \rightarrow 5$  $-27 \rightarrow 27$ 

 $w = 1/[\sigma^2(F_0^2) + (0.0734P)^2]$ + 0.5971P] where  $P = (F_0^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\rm max} < 0.001$  $\Delta \rho_{\rm max} = 0.26 \text{ e } \text{\AA}^{-3}$  $\Delta \rho_{\rm min} = -0.21 \text{ e } \text{\AA}^{-3}$ 

#### Table 1

Selected geometric parameters (Å, °).

C2-C3	1.560 (7)	C10-N2	1.318 (4)
C8-N1-C7 C8-N1-C9 C7-N1-C9 C10-N2-C11	109.7 (2) 126.7 (3) 123.5 (3) 126.1 (2)	109.7 (2) C17-N3-C15 126.7 (3) C17-N3-C19 123.5 (3) C15-N3-C19 126.1 (2)	
C1 - C2 - C3 - C4	31.5 (10)	C2-C3-C4-C5	-52.1 (12)

Table 2		
Hydrogen-bond geometry	(Å,	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C21 - H21 \cdots O2^{i}$	0.93	2.33	3.244 (5)	168
C19−H19B····O4	0.97	2.30	2.735 (4)	106
C13−H13···O1 <sup>ii</sup>	0.93	2.44	3.354 (4)	170
$C9 - H9A \cdots O3^{iii}$	0.97	2.48	3.189 (8)	130
$C4 - H4B \cdots O4^{iv}$	0.97	2.54	3.454 (8)	157
$N2-H2\cdots O3^{iii}$	0.851 (10)	2.04 (2)	2.847 (9)	158 (3)

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

The amide and water H atoms were located in a difference map and were refined with the restraints N-H = 0.86 (1) Å and O-H =0.82 Å, and with  $U_{iso}(H) = 1.2U_{eq}(\text{carrier})$  for H1 and  $1.5U_{eq}(\text{carrier})$ for water H atoms. Other H atoms were positioned geometrically, with C-H = 0.93 or 0.97 Å, and refined in a riding model, with  $U_{iso}(H) = 1.2U_{eq}(C)$  or  $1.5U_{eq}(methyl C)$ . Two of the C atoms in the cyclohexene ring were disordered over two positions, and the occupancy factors for disordered positions C3/C3' and C4/C4' were refined to 0.709 (12) and 0.291 (12). Atom O3/O3' was disordered over two positions, with occupancies of 0.77 (9) and 0.23 (9). Water atoms O6 and O6', with partial occupancies of 0.50 [initially refined to 0.504 (1)], were assigned tentatively, based only on the crystallographic evidence; the water probably derives from the methanol solvent used for recrystallization.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

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