

Zhong-Ke Hou,^{a*} Ye-Guo Ren,^b
Ming-Zhi Huang,^b Jian Song^a and
Li-Gong Chen^a

^aCollege of Pharmaceuticals and Biotechnology,
Tianjin University, Tianjin 300072, People's
Republic of China, and ^bHunan Research
Institute of Chemical Industry, Changsha
410007, People's Republic of China

Correspondence e-mail: houzk@yahoo.com

Key indicators

Single-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$
R factor = 0.030
wR factor = 0.060
Data-to-parameter ratio = 15.4

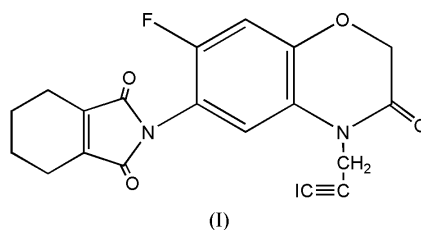
For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

2-[7-Fluoro-3,4-dihydro-4-(3-iodoprop-2-ynyl)- 3-oxo-2H-1,4-benzoxazin-6-yl]-4,5,6,7-tetrahydro- 2H-isoindole-1,3-dione

The title compound, $\text{C}_{19}\text{H}_{14}\text{FIN}_2\text{O}_4$, known as a protox
inhibitor, was synthesized from flumioxazin. The cyclohexene
ring in the molecule is in a distorted chair conformation. The
molecules are connected into two-dimensional layers by
 $\text{C}=\text{O} \cdots \text{I}$ interactions.

Comment

Flumioxazin is a selective post-emergence herbicide which
inhibits protoporphyrinogen oxidase (Yoshida *et al.*, 1991).
The title compound, (I), is a new flumioxazin analogue. It
exhibits excellent control for broadleaf weeds at low dosage
and no damage at high dosage for monocotyledon plants, such
as wheat and corn (Hou *et al.*, 2003).



The molecular structure of (I) is illustrated in Fig. 1. The
cyclohexene ring in the molecule has a distorted chair
conformation. In the crystal structure of (I), screw-related
molecules are linked by $\text{C}=\text{O} \cdots \text{I}$ interactions to form layers
parallel to the *bc* plane (Fig. 2). Such interactions have been
reported previously (Novoselov *et al.*, 1994; de Faria *et al.*,
1999).

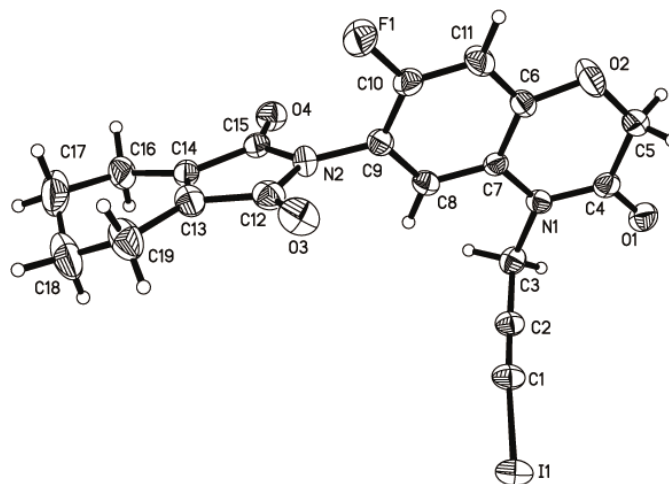


Figure 1

The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.

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Experimental

Flumioxazin (10.0 mmol) was mixed with *N*-iodosuccinimide (15.0 mmol) in acetone (10.0 ml) in the presence of an AgNO_3 catalyst (0.1 g) at ambient temperature for 24 h (Nagano *et al.*, 1987) (yield 78%). The reaction mixture was poured into iced water. The resulting precipitate was filtered off, washed with water and dried. The crude product was purified by column chromatography (silica gel, petroleum ether–acetone 5:1) to give the title compound, (I) (m.p. 555–556 K). $^1\text{H NMR}$ (CDCl_3 , δ , p.p.m): 1.66 (4H, *m*), 1.99 (4H, *m*), 4.62 (1H, *s*), 4.96 (2H, *s*), 6.76 (1H, *s*), 7.42 (1H, *s*). Compound (I) (20 mg) was dissolved in ethyl acetate (20 ml) for recrystallization. Single crystals of (I), suitable for X-ray analysis, were grown by natural evaporation of the solvent.

Crystal data

$\text{C}_{19}\text{H}_{14}\text{FIN}_2\text{O}_4$	Mo $K\alpha$ radiation
$M_r = 480.22$	Cell parameters from 908 reflections
Orthorhombic, <i>Pca2</i>	$\theta = 3.7\text{--}24.4^\circ$
$a = 9.613$ (4) Å	$\mu = 1.75 \text{ mm}^{-1}$
$b = 11.759$ (5) Å	$T = 293$ (2) K
$c = 16.571$ (7) Å	Block, colourless
$V = 1873.1$ (14) Å ³	$0.42 \times 0.38 \times 0.24 \text{ mm}$
$Z = 4$	
$D_x = 1.703 \text{ Mg m}^{-3}$	

Data collection

Bruker SMART CCD area-detector diffractometer	3752 independent reflections
φ and ω scans	2828 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 1997)	$R_{\text{int}} = 0.033$
$T_{\text{min}} = 0.472$, $T_{\text{max}} = 0.658$	$\theta_{\text{max}} = 26.4^\circ$
10 073 measured reflections	$h = -11 \rightarrow 12$
	$k = -14 \rightarrow 10$
	$l = -20 \rightarrow 19$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0247P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.030$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.060$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 0.99$	$\Delta\rho_{\text{max}} = 0.36 \text{ e \AA}^{-3}$
3752 reflections	$\Delta\rho_{\text{min}} = -0.41 \text{ e \AA}^{-3}$
244 parameters	Absolute structure: Flack (1983),
H-atom parameters constrained	1769 Friedel pairs
	Flack parameter = 0.01 (2)

H atoms were positioned geometrically, with C–H = 0.93–0.97 Å, and were refined in a riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

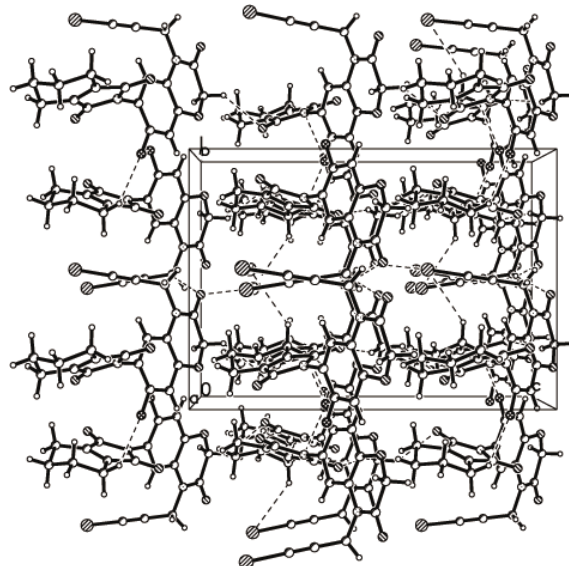


Figure 2
The crystal structure of (I), viewed along the *a* axis. The dashed lines indicate C=O...I interactions.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 1997); 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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