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Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$
 R factor = 0.042
 wR factor = 0.122
Data-to-parameter ratio = 13.7For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.***N'*-(2-Fluorobenzylidene)-2-(quinolin-8-yloxy)-
acetohydrazide methanol solvate**

In the title compound, $\text{C}_{18}\text{H}_{14}\text{FN}_3\text{O}_2 \cdot \text{CH}_4\text{O}$, all bond lengths and angles are within normal ranges. The $\text{C}_{18}\text{H}_{14}\text{FN}_3\text{O}_2$ molecule is essentially planar, with a dihedral angle of $7.83(7)^\circ$ between the planes of the benzene ring and the quinoline group. Each methanol molecule is linked to one $\text{C}_{18}\text{H}_{14}\text{FN}_3\text{O}_2$ molecule *via* intermolecular $\text{N}-\text{H} \cdots \text{O}$, $\text{O}-\text{H} \cdots \text{O}$ and $\text{O}-\text{H} \cdots \text{N}$ hydrogen bonds.

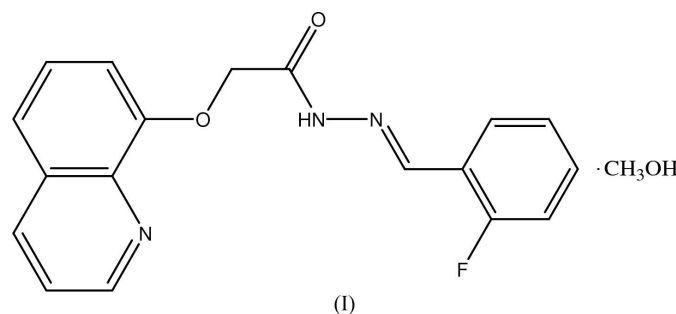
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Comment

As part of our ongoing search for good extractants of metal ions or a biologically active material, the title compound, (I), was obtained in the reaction of quinolin-8-yloxyacetic acid hydrazide and 2-fluorobenzaldehyde. We report here the crystal structure of (I).



The bond lengths and angles in (I) (Table 1) are within normal ranges (Allen *et al.*, 1987), and are comparable to those in the related compound *N,N*-diphenyl-2-(quinolin-8-yloxy)acetamide monohydrate (Wen *et al.*, 2005). The $\text{C}_{18}\text{H}_{14}\text{FN}_3\text{O}_2$ molecule is essentially planar, with a dihedral angle of $7.83(7)^\circ$ between the planes of the benzene ring and the quinoline group. The intramolecular $\text{N}2-\text{H}2\text{A} \cdots \text{O}1$ hydrogen bond, forming a five-membered ring, contributes to the planarity of the title molecule. Each methanol molecule is linked to one $\text{C}_{18}\text{H}_{14}\text{FN}_3\text{O}_2$ molecule *via* intermolecular $\text{N}-\text{H} \cdots \text{O}$, $\text{O}-\text{H} \cdots \text{O}$ and $\text{O}-\text{H} \cdots \text{N}$ hydrogen bonds (Table 2 and Fig. 2).

Experimental

Quinolin-8-yloxyacetic acid hydrazide (2.18 g, 10 mmol), 2-fluorobenzaldehyde (1.30 g, 10.5 mmol), ethanol (40 ml) and some drops of acetic acid were added to a 100 ml flask, and refluxed for 3 h. After cooling to room temperature, the mixture was filtered. Colourless single crystals suitable for X-ray diffraction study were obtained by slow evaporation of a tetrahydrofuran–methanol (1:2, *v/v*) solution over a period of 5 d.

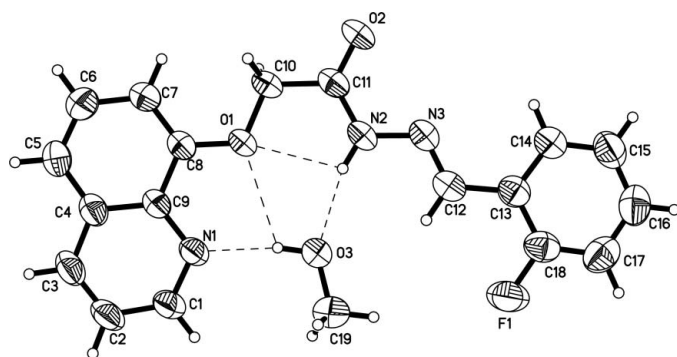


Figure 1
View of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme. The dashed lines indicate hydrogen bonds.

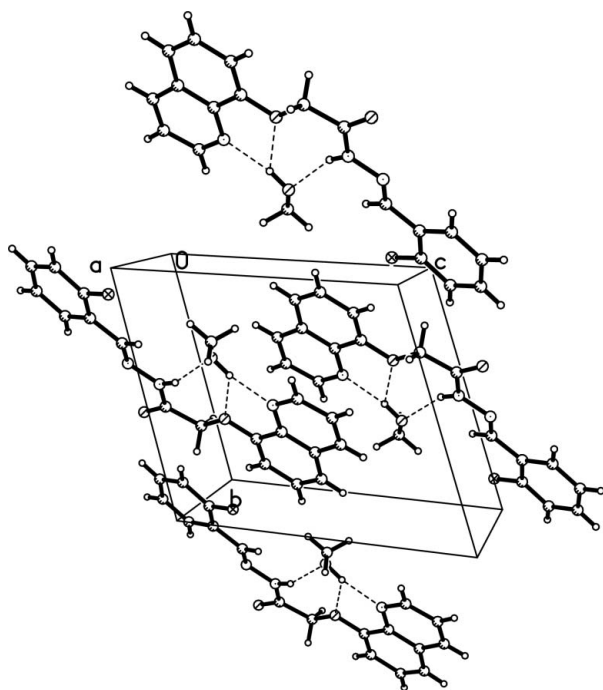


Figure 2
Packing diagram of the title compound, showing intermolecular hydrogen bonds (dashed lines).

Crystal data

$C_{18}H_{14}FN_3O_2 \cdot CH_4O$
 $M_r = 355.36$
 Triclinic, $P\bar{1}$
 $a = 9.071$ (2) Å
 $b = 9.546$ (2) Å
 $c = 11.506$ (3) Å
 $\alpha = 66.450$ (4)°
 $\beta = 71.709$ (4)°
 $\gamma = 74.490$ (4)°
 $V = 855.6$ (3) Å³

$Z = 2$
 $D_x = 1.379$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 2286 reflections
 $\theta = 2.4$ – 26.0 °
 $\mu = 0.10$ mm⁻¹
 $T = 293$ (2) K
 Column, colourless
 $0.42 \times 0.27 \times 0.13$ mm

Data collection

Siemens SMART 1000 CCD area-detector diffractometer
 ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{min} = 0.958$, $T_{max} = 0.987$
 4778 measured reflections

3286 independent reflections
 2742 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.010$
 $\theta_{max} = 26.1$ °
 $h = -11 \rightarrow 11$
 $k = -6 \rightarrow 11$
 $l = -14 \rightarrow 14$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.122$
 $S = 1.06$
 3286 reflections
 239 parameters
 H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0668P)^2 + 0.0967P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.14$ e Å⁻³
 $\Delta\rho_{min} = -0.22$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

F1—C18	1.3588 (18)	N2—C11	1.3451 (19)
O1—C8	1.3723 (16)	N2—N3	1.3819 (17)
O1—C10	1.4241 (16)	N3—C12	1.271 (2)
O2—C11	1.2167 (16)		
C8—O1—C10	115.37 (11)	C12—N3—N2	115.58 (12)
C11—N2—N3	118.17 (11)	O1—C10—C11	113.46 (12)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N2—H2A \cdots O1	0.86	2.33	2.698 (2)	106
N2—H2A \cdots O3	0.86	2.06	2.879 (2)	158
O3—H1O3 \cdots O1	0.90 (3)	2.50 (3)	3.011 (2)	116 (2)
O3—H1O3 \cdots N1	0.90 (3)	1.88 (3)	2.771 (2)	169 (2)

The solvent H atoms were located in a difference Fourier map, while the other H atoms were positioned geometrically. The hydroxy atom H1O3 was refined freely. All other H atoms were allowed to ride on their parent atoms, with C—H = 0.93–0.97 Å, N—H = 0.86 Å and $U_{iso}(H) = 1.2$ – $1.5U_{eq}(\text{parent atom})$.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

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