## organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.005 Å R factor = 0.047 wR factor = 0.097 Data-to-parameter ratio = 7.6

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

# *N*-(3-Methoxyphenyl)-2-(quinolin-8-yloxy)-acetamide monohydrate

In the title compound,  $C_{18}H_{16}N_2O_3 \cdot H_2O$ , the organic molecules and water molecules are linked by intermolecular N- $H \cdot \cdot \cdot O$ ,  $O-H \cdot \cdot \cdot O$  and  $O-H \cdot \cdot \cdot N$  hydrogen bonds. The structure is also stabilized by  $C-H \cdot \cdot \cdot O$  hydrogen bonds.

#### Comment

In our ongoing studies of amide-type acyclic compounds with 8-hydroxyquinolinate as the skeleton (Wen *et al.*, 2005), the title compound has been synthesized and its structure is reported here.



All bond lengths and angles in (I) are normal (Allen *et al.*, 1987) and comparable with those in a related compound (Wen *et al.*, 2005). The molecules are linked by intermolecular hydrogen bonds, with N and O atoms acting as donors and acceptors (Table 1, Fig. 2). There are also three weak C– $H \cdots O$  hydrogen bonds, one of them being intramolecular (Table 1). The angle between the mean planes of the benzene and quinoline rings is 56.2 (1)° (Nardelli, 1995).

#### **Experimental**

2-Chloro-*N*-(3-methoxyphenyl)acetamide (4.0 g, 20 mmol),  $K_2CO_3$ (3.04 g, 22 mmol) and KI (0.5 g, 3 mmol) were added to a solution of 8-hydroxyquinoline (2.9 g, 20 mmol) in acetone (40 ml). The mixture was stirred at 333 K for 6 h. After cooling to room temperature, the mixture was washed three times with water and then filtered. The title compound was obtained after drying the resulting white powder at room temperature for 48 h (elemental analysis found: C 66.20, H 5.62, N 8.55, O 19.66; calculated: C 66.25, H 5.56, N 8.58, O 19.61 wt. %). Colourless single crystals suitable for X-ray diffraction studies were obtained by slow evaporation of an ethyl acetate solution over a period of 5 d.

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Crystal data
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\begin{array}{l} C_{18}H_{16}N_2O_3 \cdot H_2O\\ M_r = 326.34\\ \text{Monoclinic, }P2_1\\ a = 8.1881 (11) \text{ Å}\\ b = 7.0491 (10) \text{ Å}\\ c = 14.094 (2) \text{ Å}\\ \beta = 100.553 (2)^\circ\\ V = 799.75 (19) \text{ Å}^3 \end{array}
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Z = 2  $D_x = 1.355 \text{ Mg m}^{-3}$ Mo K\alpha radiation  $\mu = 0.10 \text{ mm}^{-1}$  T = 293 (2) KBlock, colourless  $0.26 \times 0.16 \times 0.12 \text{ mm}$ 

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#### Figure 1

The structure of compound (I), showing 40% probability displacement ellipsoids and the atom-numbering scheme. Hydrogen bonds are shown as dashed lines. [Symmetry code: (I) x, -1 + y, 1 + z.]

#### Data collection

Siemens SMART 1000 CCD area-	4571 measured reflections
detector diffractometer	1706 independent reflections
$\omega$ scans	1288 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan	$R_{\rm int} = 0.026$
(SADABS; Sheldrick, 1996)	$\theta_{\rm max} = 26.1^{\circ}$
T = 0.975 T = 0.989	

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0469P)^2]$		
$R[F^2 > 2\sigma(F^2)] = 0.047$	+ 0.0065P]		
$wR(F^2) = 0.098$	where $P = (F_0^2 + 2F_c^2)/3$		
S = 1.07	$(\Delta/\sigma)_{\rm max} < 0.001$		
1706 reflections	$\Delta \rho_{\rm max} = 0.14 \text{ e } \text{\AA}^{-3}$		
224 parameters	$\Delta \rho_{\rm min} = -0.15 \ {\rm e} \ {\rm \AA}^{-3}$		
H atoms treated by a mixture of			
independent and constrained			
refinement			

#### Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$O1W-H1W\cdots O2^i$	0.85 (3)	1.99 (3)	2.838 (4)	174 (4)
$N2-H2B\cdots O1W^{ii}$	0.86	2.00	2.844 (4)	167
$O1W - H2W \cdot \cdot \cdot N1^{i}$	0.86 (3)	1.98 (3)	2.830 (4)	173 (4)
C8-H8A···O2 <sup>iii</sup>	0.93	2.50	3.365 (4)	154
$C13-H13A\cdots O1W^{ii}$	0.93	2.59	3.329 (4)	136
C17−H17A···O2	0.93	2.37	2.906 (5)	116

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

All H atoms were located in a difference Fourier map. Water H atoms were refined with O1W-H1W and O1W-H2W distance restraints of 0.85 Å and with a common  $U_{iso}$  value. The remaining H atoms were placed in idealized positions and constrained to ride on their parent atoms, with  $C_{aryl}-H = 0.93$  Å,  $C_{methyl}-H = 0.96$  Å,  $C_{methylene}-H = 0.97$  Å, and N-H = 0.86 Å, and with  $U_{iso}(H) = 1.2$   $U_{eq}(C,N)$  or  $1.5U_{iso}(methyl)$  C). 1307 Friedel pairs were merged





A packing diagram for (I), viewed down the *a* axis. Only the intermolecular hydrogen bonds with O or N donor atoms are shown, as dashed lines.

before the final refinement because of the absence of any significant anomalous scattering effects.

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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