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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.032 wR factor = 0.083 Data-to-parameter ratio = 8.4

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

N-(4-Methoxyphenyl)-2-(quinolin-8-yloxy)-acetamide monohydrate

In the title compound, $C_{18}H_{16}N_2O_3\cdot H_2O$, all bond lengths and angles are within normal ranges. The dihedral angle formed by the benzene ring with the quinoline moiety is 67.06 (7)°. Molecules are linked into chains along the *a* axis by intermolecular N-H···O, O-H···O and O-H···N hydrogen bonds involving the solvent water molecule. The packing is further stabilized by π - π interactions.

Comment

Recently, we have reported the structure of an amide-type acyclic compound with 8-hydroxyquinolinate as the skeleton, namely *N*-phenyl-2-(quinolin-8-yloxy)acetamide hemihydrate, (II) (Li *et al.*, 2005). We have synthesized and carried out the structure determination of the title compound, (I), reported here.



All bond lengths and angles in (I) (Table 1) are within normal ranges (Allen et al., 1987) and comparable with those in the related compound (II). The dihedral angle formed by the benzene ring with the quinoline moiety is $67.06 (7)^\circ$. There is one intramolecular hydrogen bond, viz. $C13-H13A\cdots O2$, forming a six-membered ring. In the crystal structure, molecules are linked into chains along the *a* axis by intermolecular N2-H2A···O1 W^{i} , $O1W - H2W1 \cdots O2^{ii}$ and O1W-H1W1···N1ⁱⁱ hydrogen bonds (Table 2 and Fig. 2) involving the solvent water molecule. The packing is further stabilized by π - π interactions involving the benzene rings (Cg2 is the centroid of the C12–C17 benzene ring): $Cg3 \cdot \cdot \cdot Cg3(-\frac{1}{2} + x, \frac{1}{2} - x)$ y, -z) = 3.737 Å.

Experimental

To a solution of 8-hydroxyquinoline (2.9 g, 20 mmol) in acetone (40 ml) were added 2-chloro-*N*-(4-methoxyphenyl)acetamide (3.99 g, 20 mmol), K_2CO_3 (3.04 g, 22 mmol) and KI (0.5 g), and the resulting mixture was stirred at 333 K for 5 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 yellow powder at room temperature for 48 h. Yellow single crystals suitable for X-ray diffraction study were obtained by slow evaporation of a petroleum ether–ethyl acetate (1:3 ν/ν) solution over a period of 6 d.

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Crystal data

 $C_{18}H_{16}N_2O_3 \cdot H_2O$ $M_r = 326.34$ Orthorhombic, $P2_12_12_1$ a = 6.8814 (13) Å b = 11.627 (2) Å c = 20.468 (4) Å V = 1637.6 (5) Å³ Z = 4 $D_x = 1.324$ Mg m⁻³

Data collection

Siemens SMART 1000 CCD areadetector diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{min} = 0.961, T_{max} = 0.989$ 9310 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.032$ $wR(F^2) = 0.084$ S = 1.121901 reflections 225 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Selected bond lengths (Å).

01 - C8	1 371 (2)	N2-C11	1340(2)
O1-C10	1.418 (2)	N2-C12	1.426 (2)
O2-C11	1.223 (2)		

Mo $K\alpha$ radiation

reflections

 $\begin{array}{l} \theta = 2.6 - 25.8^{\circ} \\ \mu = 0.10 \ \mathrm{mm}^{-1} \end{array}$

T = 293 (2) K

Plate, yellow

 $R_{\rm int} = 0.018$

 $\theta_{\rm max} = 26.1^{\circ}$

 $h = -8 \rightarrow 8$

 $k = -13 \rightarrow 14$

 $l = -25 \rightarrow 24$

Cell parameters from 4023

 $0.42 \times 0.42 \times 0.12 \text{ mm}$

1901 independent reflections

1737 reflections with $I > 2\sigma(I)$

 $w = 1/[\sigma^2(F_0^2) + (0.0392P)^2]$

+ 0.1826P] where $P = (F_0^2 + 2F_c^2)/3$

 $\Delta \rho_{\rm max} = 0.13 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.14 \ {\rm e} \ {\rm \AA}^{-3}$

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

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots $
$\begin{array}{c} N2 - H2A \cdots O1W^{i} \\ O1W - H1W1 \cdots N1^{ii} \\ O1W - H2W1 \cdots O2^{ii} \\ C13 - H13A \cdots O2^{ii} \end{array}$	0.86	2.01	2.853 (2)	166
	0.87 (3)	2.02 (3)	2.886 (3)	176 (3)
	0.84 (3)	1.93 (3)	2.762 (2)	173 (2)
	0.93	2.41	2.889 (3)	112

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

All H atoms were located in difference Fourier maps and all except water H atoms, which were refined freely, were constrained to ride on their parent atoms, with C–H distances in the range 0.93–0.97 Å, and with $U_{\rm iso}(\rm H) = 1.2 U_{eq}(\rm C)$ or $1.5 U_{eq}(\rm C)$ for methyl H atoms. The Friedel reflections were merged 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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Figure 1

The structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme.



Figure 2

Packing diagram of (I), showing the intermolecular hydrogen bonds (dashed lines), viewed down the b axis.

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