

Aquabis(2-methylquinolin-8-olato- κ^2N,O)zinc(II)

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Key indicators

Single-crystal X-ray study
 $T = 299\text{ K}$
Mean $\sigma(\text{C}–\text{C}) = 0.012\text{ \AA}$
 R factor = 0.062
 wR factor = 0.154
Data-to-parameter ratio = 12.3

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

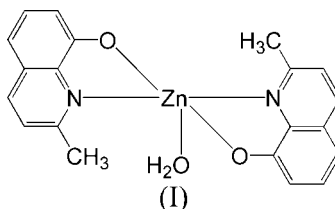
The title molecule, $[\text{Zn}(\text{C}_{10}\text{H}_8\text{NO})_2(\text{H}_2\text{O})]$, lies on a crystallographic twofold axis. The Zn^{II} atom is in a trigonal-bipyramidal coordination geometry formed by two N atoms of two quinoline groups, two O atoms and a water molecule. In the crystal structure, intermolecular hydrogen bonds link molecules into a two-dimensional network.

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Comment

Hydroxyquinolines and their derivatives are well known complexing agents in analytical chemistry (Czugler *et al.*, 2001). In addition, metal chelators based on the quinoline core have been developed as potential agents for neuroprotection in neurodegenerative diseases (Zheng *et al.*, 2005). In this context, clioquinol is a leading compound being investigated as a biomarker for β -amyloid Zn^{II} complexes in Alzheimer's disease (Opazo *et al.*, 2006). We have an interest in investigating new metal-chelator probes for neuroprotection in neurodegenerative diseases (da Silva *et al.*, 2006*a,b,c,d,e*). The crystal structure of the title compound, (I), is reported here.



The molecular structure of (I) is shown in Fig. 1. The Zn atom is in a trigonal-bipyramidal coordination geometry formed by two N atoms of two quinoline groups, two O atoms and a water molecule. The Zn atom and the water O atom both lie on a crystallographic twofold axis. Selected bond distances and angles are given in Table 1. The hydroxyquinoline ligands are bis-chelating through the quinoline N atom and the O atom, forming a five-membered ring including the Zn atom.

In the crystal structure, two intermolecular hydrogen bonds link the molecules into a two-dimensional network (Fig. 2 and Table 2).

Experimental

The title compound was prepared according to a literature procedure (Macías *et al.*, 2003). 2-Methyl-8-quinolinol (1.5 mmol) was dissolved in methanol (75 ml) and aqueous ammonia (2 ml *M* NH_4OH) aqueous ammonia was added. During stirring of this solution, ZnCl_2 (0.75 mmol) dissolved in methanol (50 ml) was added dropwise. When the addition was complete, a green precipitate was formed

which was separated by filtration (yield 80%). Single crystals of (I) suitable for X-ray data collection appeared after 2 d from a methanol solution.

Crystal data

[Zn(C₁₀H₈NO)₂(H₂O)]
M_r = 399.73
 Orthorhombic, *Pbcn*
a = 7.284 (1) Å
b = 9.131 (2) Å
c = 25.199 (5) Å
V = 1676.0 (5) Å³

Z = 4
D_x = 1.584 Mg m⁻³
 Cu *K*α radiation
 μ = 2.23 mm⁻¹
T = 299 (2) K
 Block, light red
 0.20 × 0.10 × 0.08 mm

Data collection

Enraf–Nonius CAD-4 diffractometer
 ω/2θ scans
 Absorption correction: ψ scan (North *et al.*, 1968)
T_{min} = 0.615, *T_{max}* = 0.841
 2878 measured reflections

1499 independent reflections
 655 reflections with *I* > 2σ(*I*)
R_{int} = 0.198
 θ_{max} = 67.0°
 3 standard reflections
 frequency: 120 min
 intensity decay: 1%

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.062
wR(*F*²) = 0.154
S = 0.93
 1499 reflections
 122 parameters

H atoms treated by a mixture of independent and constrained refinement
w = 1/[σ²(*F_o*²) + (0.0499*P*)²]
 where *P* = (*F_o*² + 2*F_c*²)/3
 (Δσ)_{max} = 0.010
 Δρ_{max} = 0.77 e Å⁻³
 Δρ_{min} = -0.59 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

N1–Zn1	2.114 (5)	O2–Zn1	2.103 (6)
O1–Zn1	2.022 (5)		
O1 ⁱ –Zn1–O1	125.0 (3)	O1–Zn1–N1	80.4 (2)
O1–Zn1–O2	117.48 (14)	O2–Zn1–N1	89.97 (18)
O1 ⁱ –Zn1–N1	99.6 (2)	N1–Zn1–N1 ⁱ	179.9 (4)

Symmetry code: (i) $-x, y, -z + \frac{1}{2}$.

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> –H... <i>A</i>	<i>D</i> –H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> –H... <i>A</i>
O2–H ₂ O...O1 ⁱⁱ	0.92 (7)	1.82 (7)	2.736 (6)	176 (7)
C10–H10A...N1 ⁱⁱⁱ	0.96	2.61	3.555 (10)	169

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

The symmetry-independent O-bound H atom was located in a difference map and its positional parameters were refined. The other H atoms were positioned with idealized geometry using a riding model, with C–H = 0.93 Å (aromatic) or 0.96 Å (methyl). For all H atoms, *U*_{iso}(H) = 1.2*U*_{eq}(parent atom). The ratio of observed to unique reflections is lower than normal and this has resulted in a higher than normal value for *R*_{int} (0.198). This in turn can lower the precision of the results.

Data collection: *CAD-4-PC* Software (Enraf–Nonius, 1996); cell refinement: *CAD-4-PC* Software; data reduction: *REDU4* (Stoe & Cie, 1987); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

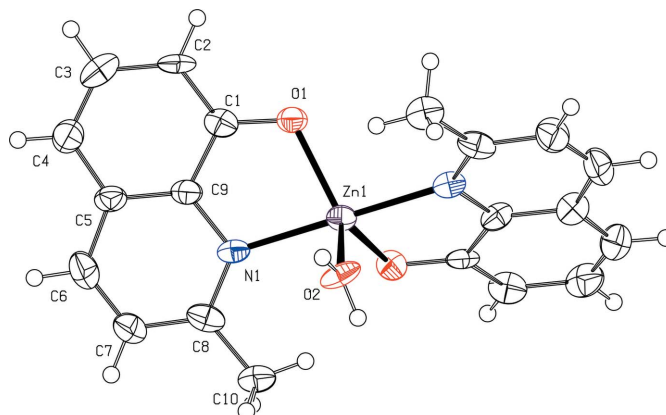


Figure 1

The molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. Unlabelled atoms are related to the corresponding labelled atoms by the symmetry operation $(-x, y, -z + \frac{1}{2})$.

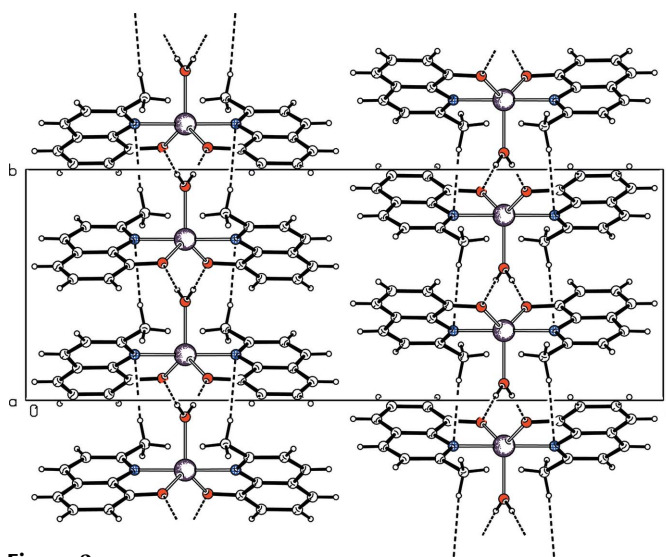


Figure 2

Part of the crystal structure of (I), with hydrogen bonds shown as dashed lines.

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