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Aquabis(2-methyl-4-oxopyrido[1,2-*a*]-pyrimidin-9-olato)zinc(II) monohydrate

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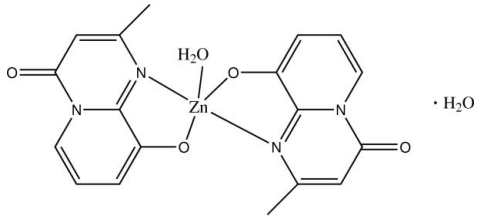
Key indicators: single-crystal X-ray study; T = 293 K; mean σ(C–C) = 0.005 Å; R factor = 0.049; wR factor = 0.128; data-to-parameter ratio = 14.2.

The crystal structure of the title compound, $[Zn(C_9H_7N_2O_2)_2(H_2O)]\cdot H_2O$, involves discrete mononuclear complex molecules. The special positions on the rotation twofold axis are occupied by Zn^{II} and O atoms of the coordinated and uncoordinated water molecules. The coordination around the Zn^{II} atom can be described as transitional from trigonal-bipyramidal to square-pyramidal. The two chelating 2-methyl-4-oxopyrido[1,2-*a*]pyrimidin-9-olate ligands and the coordinated water molecule form the Zn coordination. O—H···O hydrogen bonds between the coordinated water molecule and the ligand and between the uncoordinated water molecule and the ligand dominate the crystal packing.

Related literature

For the design and synthesis of self-assembling systems with organic ligands containing N and O donors, see: Bayot *et al.* (2006); Chen *et al.* (2007). For the structures of quinolin-8-ol complexes, see: Wu *et al.* (2006).



Experimental

Crystal data

$[Zn(C_9H_7N_2O_2)_2(H_2O)]\cdot H_2O$
M_r = 451.73
Orthorhombic, *Pbcn*
a = 7.7670 (16) Å
b = 16.045 (3) Å
c = 14.006 (3) Å
V = 1745.4 (6) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 1.46 mm^{-1}
T = 293 (2) K
0.25 × 0.15 × 0.12 mm

Data collection

Rigaku Scxmini 1K CCD area-detector diffractometer
Absorption correction: multi-scan (*CrystalClear*; Rigaku, 2005)
T_{min} = 0.752, T_{max} = 0.831
16899 measured reflections
2005 independent reflections
1470 reflections with $I > 2\sigma(I)$
R_{int} = 0.070

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.049
$wR(F^2)$ = 0.128
S = 1.07
2005 reflections
141 parameters
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.50 e Å^{-3}
$\Delta\rho_{min}$ = −0.56 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
O3—H3B···O2^{i}	0.72 (4)	2.11 (4)	2.823 (3)	170 (5)
O4—H4B···O1^{ii}	0.78 (5)	2.23 (5)	3.008 (4)	176 (6)

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

Data collection: *CrystalClear* (Rigaku, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: KP2191).

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supporting information

Acta Cryst. (2009). E**65**, m91 [doi:10.1107/S1600536808039615]

Aquabis(2-methyl-4-oxopyrido[1,2-*a*]pyrimidin-9-olato)zinc(II) monohydrate

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S1. Comment

Considerable attention has been paid to the design and synthesis of self-assembling systems with organic ligands containing N and O donors (Bayot *et al.*, 2006; Chen, *et al.*, 2007). Quinolin-8-ol is one such ligand and several crystal structures of complexes containing it have been reported (Wu *et al.*, 2006). We report here the synthesis and crystal structure of the title complex, (I) (Fig. 1). In (I), the Zn atom is penta-coordinated by two pyridine nitrogen atoms and two oxygen atoms from the hydroxy groups and water molecule (Fig. 1 and Table 1). Intermolecular O—H···O hydrogen bonds (Table 2 and Fig. 2) connect the molecules of (I) define the crystal packing.

S2. Experimental

All chemicals used (reagent grade) were commercially available. An aqueous solution (5 ml) of $ZnCl_2$ (13.6 mg, 0.1 mmol) was added by constant stirring to an ethanol solution (10 ml) containing 2-methyl-9-hydroxylpyrido [1,2-*a*]pyrimidin-4-one (17.6 mg, 0.1 mmol) then filtered off. After a few days, colourless, well shaped single crystals in the form of prisms deposited in the mother-liquid. They were separated off, washed with cold ethanol and dried in air at room temperature.

S3. Refinement

In general, H atoms bound to carbon were placed in geometrical positions and refined using a riding model, with C—H = 0.94Å and U_{iso}(H) = 1.2U_{eq}(C). The H of water were located from the difference map and refined freely.

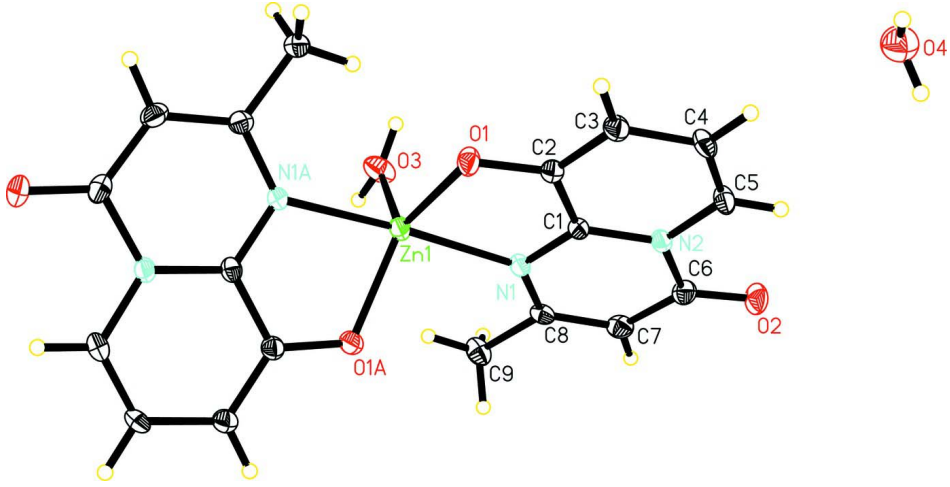

Figure 1

The molecular structure of the title molecule and the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. [Symmetry code A: -*x*, *y*, 0.5 - *z*.]

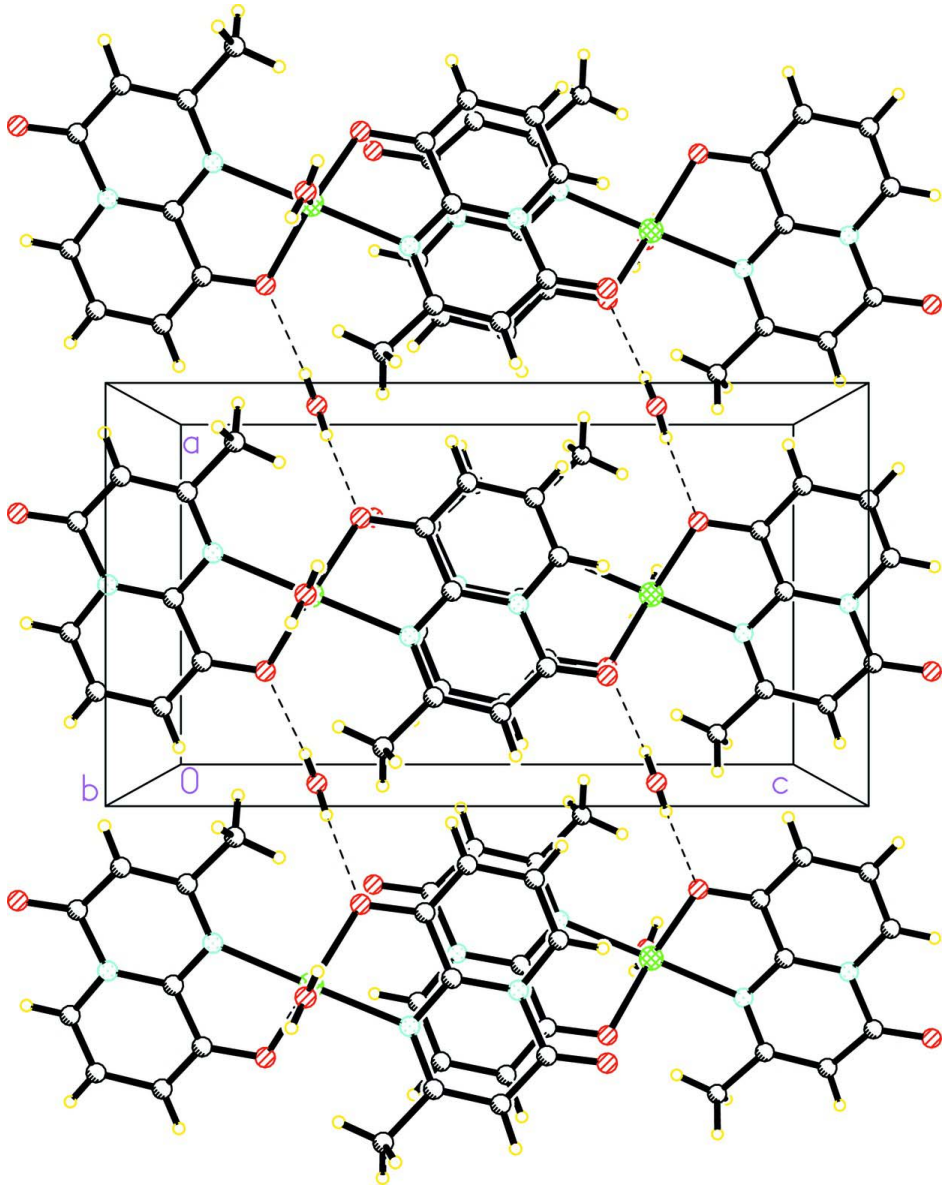

Figure 2

Crystal packing of the compound (I). Hydrogen bonds are shown as dashed lines.

Aquabis(2-methyl-4-oxopyrido[1,2-*a*]pyrimidin-9-olato)zinc(II) monohydrate

Crystal data

$[Zn(C_9H_7N_2O_2)_2(H_2O)]\cdot H_2O$
$M_r = 451.73$
Orthorhombic, *Pbcn*
Hall symbol: -P 2n 2ab
$a = 7.7670\ (16)$ Å
$b = 16.045\ (3)$ Å
$c = 14.006\ (3)$ Å
$V = 1745.4\ (6)$ Å^3
$Z = 4$
$F(000) = 928$
$D_x = 1.719$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 13380 reflections
$\theta = 3.0–27.6°$
$\mu = 1.46$ mm^{-1}
$T = 293$ K
Prism, colourless
0.25 × 0.15 × 0.12 mm

Data collection

Rigaku Scxmini 1K CCD area-detector diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 8.192 pixels mm^{-1}
Thin–slice ω scans
Absorption correction: multi-scan (*CrystalClear*; Rigaku, 2005)
T_{min} = 0.752, T_{max} = 0.831

16899 measured reflections
2005 independent reflections
1470 reflections with $I > 2\sigma(I)$
R_{int} = 0.070
θ_{max} = 27.5°, θ_{min} = 3.3°
h = −10→10
k = −20→20
l = −18→18

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.049
$wR(F^2)$ = 0.128
S = 1.07
2005 reflections
141 parameters
0 restraints
Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H atoms treated by a mixture of independent and constrained refinement
$w = 1/[\sigma^2(F_o^2) + (0.0591P)^2 + 2.2542P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max}$ = 0.50 e $Å^{-3}$
$\Delta\rho_{min}$ = −0.56 e $Å^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2, conventional R-factors R are based on F, with F set to zero for negative F^2. The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R-factors(gt) *etc*. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F, and R- factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	x	*y*	*z*	U_{iso}*/U_{eq}
C1	−0.0089 (4)	0.1222 (2)	0.4495 (2)	0.0220 (7)
C2	−0.1741 (4)	0.0991 (2)	0.4103 (2)	0.0242 (7)
C3	−0.2932 (5)	0.0659 (2)	0.4714 (3)	0.0290 (8)
H3A	−0.4012	0.0508	0.4484	0.035*
C4	−0.2541 (5)	0.0543 (2)	0.5686 (2)	0.0301 (8)
H4A	−0.3366	0.0317	0.6092	0.036*
C5	−0.0987 (5)	0.0754 (2)	0.6036 (2)	0.0308 (8)
H5A	−0.0747	0.0668	0.6680	0.037*
C6	0.1875 (5)	0.1347 (2)	0.5853 (2)	0.0287 (8)
C7	0.3026 (5)	0.1688 (2)	0.5193 (3)	0.0301 (8)
H7A	0.4105	0.1859	0.5402	0.036*
C8	0.2616 (4)	0.1779 (2)	0.4246 (2)	0.0240 (7)
C9	0.3867 (5)	0.2129 (2)	0.3541 (3)	0.0332 (9)
H9A	0.3346	0.2143	0.2920	0.050*
H9B	0.4877	0.1784	0.3523	0.050*

H9C	0.4182	0.2684	0.3729	0.050*
N1	0.1066 (4)	0.15560 (17)	0.39001 (19)	0.0230 (6)
N2	0.0251 (3)	0.10953 (18)	0.54498 (19)	0.0235 (6)
O1	−0.1984 (3)	0.11255 (17)	0.31857 (17)	0.0322 (6)
O2	0.2095 (4)	0.12461 (18)	0.67183 (17)	0.0383 (7)
Zn1	0.0000	0.16492 (4)	0.2500	0.0259 (2)
O3	0.0000	0.2946 (3)	0.2500	0.0420 (10)
O4	−0.5000	−0.0113 (3)	0.7500	0.0529 (12)
H4B	−0.420 (7)	−0.038 (4)	0.765 (4)	0.08 (2)*
H3B	0.069 (6)	0.320 (3)	0.232 (3)	0.046 (15)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0236 (16)	0.0234 (15)	0.0189 (15)	0.0037 (15)	0.0017 (14)	−0.0002 (12)
C2	0.0215 (17)	0.0284 (18)	0.0225 (17)	0.0009 (14)	0.0003 (14)	0.0009 (14)
C3	0.0224 (17)	0.037 (2)	0.0275 (18)	−0.0033 (15)	0.0014 (15)	0.0022 (15)
C4	0.0279 (19)	0.037 (2)	0.0254 (18)	−0.0065 (16)	0.0099 (15)	0.0032 (16)
C5	0.035 (2)	0.038 (2)	0.0197 (16)	−0.0017 (17)	0.0052 (16)	0.0021 (15)
C6	0.0291 (19)	0.0329 (19)	0.0241 (17)	0.0021 (16)	−0.0055 (15)	−0.0049 (15)
C7	0.0211 (17)	0.036 (2)	0.0335 (18)	−0.0014 (15)	−0.0031 (15)	−0.0063 (16)
C8	0.0206 (16)	0.0242 (17)	0.0273 (17)	0.0008 (13)	0.0030 (14)	−0.0064 (14)
C9	0.030 (2)	0.040 (2)	0.0295 (18)	−0.0102 (17)	0.0041 (16)	−0.0066 (16)
N1	0.0222 (15)	0.0285 (15)	0.0185 (13)	−0.0024 (12)	0.0024 (11)	−0.0012 (11)
N2	0.0234 (16)	0.0295 (15)	0.0177 (13)	0.0005 (12)	0.0015 (11)	−0.0008 (11)
O1	0.0234 (13)	0.0505 (16)	0.0228 (12)	−0.0071 (12)	−0.0026 (10)	0.0072 (11)
O2	0.0387 (16)	0.0545 (17)	0.0218 (13)	−0.0045 (14)	−0.0078 (11)	0.0000 (12)
Zn1	0.0240 (3)	0.0347 (3)	0.0191 (3)	0.000	0.0024 (2)	0.000
O3	0.040 (2)	0.032 (2)	0.054 (3)	0.000	0.021 (2)	0.000
O4	0.046 (3)	0.059 (3)	0.054 (3)	0.000	0.000 (3)	0.000

Geometric parameters (Å, º)

C1—N1	1.336 (4)	C7—C8	1.372 (5)
C1—N2	1.378 (4)	C7—H7A	0.9300
C1—C2	1.444 (5)	C8—N1	1.346 (4)
C2—O1	1.316 (4)	C8—C9	1.494 (5)
C2—C3	1.368 (5)	C9—H9A	0.9600
C3—C4	1.408 (5)	C9—H9B	0.9600
C3—H3A	0.9300	C9—H9C	0.9600
C4—C5	1.346 (5)	N1—Zn1	2.134 (3)
C4—H4A	0.9300	O1—Zn1	2.001 (2)
C5—N2	1.379 (4)	Zn1—O1^{i}	2.001 (2)
C5—H5A	0.9300	Zn1—O3	2.081 (4)
C6—O2	1.234 (4)	Zn1—N1^{i}	2.134 (3)
C6—C7	1.398 (5)	O3—H3B	0.72 (4)
C6—N2	1.440 (4)	O4—H4B	0.78 (5)

N1—C1—N2	122.4 (3)	C8—C9—H9A	109.5
N1—C1—C2	117.5 (3)	C8—C9—H9B	109.5
N2—C1—C2	120.1 (3)	H9A—C9—H9B	109.5
O1—C2—C3	125.2 (3)	C8—C9—H9C	109.5
O1—C2—C1	117.2 (3)	H9A—C9—H9C	109.5
C3—C2—C1	117.6 (3)	H9B—C9—H9C	109.5
C2—C3—C4	120.7 (3)	C1—N1—C8	118.9 (3)
C2—C3—H3A	119.7	C1—N1—Zn1	109.9 (2)
C4—C3—H3A	119.7	C8—N1—Zn1	131.2 (2)
C5—C4—C3	120.9 (3)	C1—N2—C5	120.2 (3)
C5—C4—H4A	119.6	C1—N2—C6	120.5 (3)
C3—C4—H4A	119.6	C5—N2—C6	119.3 (3)
C4—C5—N2	120.5 (3)	C2—O1—Zn1	115.3 (2)
C4—C5—H5A	119.7	O1—Zn1—O1^{i}	130.33 (16)
N2—C5—H5A	119.7	O1—Zn1—O3	114.83 (8)
O2—C6—C7	127.8 (4)	O1^{i}—Zn1—O3	114.83 (8)
O2—C6—N2	118.0 (3)	O1—Zn1—N1^{i}	96.48 (10)
C7—C6—N2	114.2 (3)	O1^{i}—Zn1—N1^{i}	80.11 (10)
C8—C7—C6	122.2 (3)	O3—Zn1—N1^{i}	94.02 (7)
C8—C7—H7A	118.9	O1—Zn1—N1	80.11 (10)
C6—C7—H7A	118.9	O1^{i}—Zn1—N1	96.48 (10)
N1—C8—C7	121.8 (3)	O3—Zn1—N1	94.02 (7)
N1—C8—C9	116.4 (3)	N1^{i}—Zn1—N1	171.96 (15)
C7—C8—C9	121.8 (3)	Zn1—O3—H3B	125 (4)

N1—C1—C2—O1	0.1 (5)	N1—C1—N2—C6	2.0 (5)
N2—C1—C2—O1	−179.8 (3)	C2—C1—N2—C6	−178.2 (3)
N1—C1—C2—C3	−179.2 (3)	C4—C5—N2—C1	−0.2 (5)
N2—C1—C2—C3	0.9 (5)	C4—C5—N2—C6	177.5 (3)
O1—C2—C3—C4	−179.8 (3)	O2—C6—N2—C1	177.5 (3)
C1—C2—C3—C4	−0.6 (5)	C7—C6—N2—C1	−2.0 (5)
C2—C3—C4—C5	−0.1 (6)	O2—C6—N2—C5	−0.1 (5)
C3—C4—C5—N2	0.5 (6)	C7—C6—N2—C5	−179.7 (3)
O2—C6—C7—C8	−178.9 (4)	C3—C2—O1—Zn1	178.4 (3)
N2—C6—C7—C8	0.6 (5)	C1—C2—O1—Zn1	−0.9 (4)
C6—C7—C8—N1	1.1 (5)	C2—O1—Zn1—O1^{i}	91.1 (2)
C6—C7—C8—C9	−178.7 (3)	C2—O1—Zn1—O3	−88.9 (2)
N2—C1—N1—C8	−0.2 (5)	C2—O1—Zn1—N1^{i}	173.6 (2)
C2—C1—N1—C8	179.9 (3)	C2—O1—Zn1—N1	0.9 (2)
N2—C1—N1—Zn1	−179.5 (2)	C1—N1—Zn1—O1	−0.8 (2)
C2—C1—N1—Zn1	0.7 (4)	C8—N1—Zn1—O1	−179.9 (3)
C7—C8—N1—C1	−1.3 (5)	C1—N1—Zn1—O1^{i}	−130.7 (2)
C9—C8—N1—C1	178.5 (3)	C8—N1—Zn1—O1^{i}	50.2 (3)
C7—C8—N1—Zn1	177.8 (2)	C1—N1—Zn1—O3	113.7 (2)
C9—C8—N1—Zn1	−2.4 (4)	C8—N1—Zn1—O3	−65.4 (3)

N1—C1—N2—C5	179.6 (3)	C1—N1—Zn1—N1^{i}	−66.3 (2)
C2—C1—N2—C5	−0.6 (5)	C8—N1—Zn1—N1^{i}	114.6 (3)

Symmetry code: (i) −*x*, *y*, −*z*+1/2.

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
O3—H3*B*···O2^{ii}	0.72 (4)	2.11 (4)	2.823 (3)	170 (5)
O4—H4*B*···O1^{iii}	0.78 (5)	2.23 (5)	3.008 (4)	176 (6)

Symmetry codes: (ii) −*x*+1/2, −*y*+1/2, *z*−1/2; (iii) *x*, −*y*, *z*+1/2.