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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.043 wR factor = 0.120 Data-to-parameter ratio = 15.1

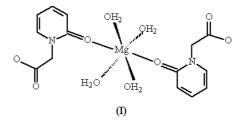
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

© 2004 International Union of Crystallography Printed in Great Britain – all rights reserved The title complex, tetraaquabis[$(2-\infty -1,2-dihydropyridin-1-yl)$ acetato]magnesium(II), [Mg(C₇H₆NO₃)₂(H₂O)₄], is a neutral mononuclear molecule consisting of an Mg^{II} ion, two (2-0x0-4*H*-pyridin-1-yl)acetate ligands and four coordinated water molecules. The Mg^{II} atom, located on a symmetry center, has octahedral coordination involving two carbonyl O atoms of different (2-0x0-4*H*-pyridin-1-yl)acetate ligands and four water molecules. O—H···O intermolecular hydrogen bonds form a layer structure.

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Comment

(4-Oxo-4*H*-pyridin-1-yl)acetic acid, known as an important medical intermediate (Edwards *et al.*, 1977), is a potential multidentate ligand with a versatile binding mode. However, there is little information on the structure of metal complexes of this acid. Recently, we used the 4-oxo-1(4*H*)-(carboxy-methyl)pyridinium hydroxide inner salt to prepare metal complexes (Gao *et al.*, 2004; Zhang *et al.*, 2004). In order to explore the structural properties of related complexes, we have synthesized the title magnesium(II) complex, (I), by the reaction of (2-oxo-4*H*-pyridin-1-yl)acetic acid with magnesium(II) nitrate hexahydrate under basic conditions. The crystal structure of (I) is described here.



As illustrated in Fig. 1, the title complex has a mononuclear structure, in which the (2-oxo-4H-pyridin-1-yl)acetate groups are bonded to the Mg^{II} atom through the carbonyl O atoms in a monodentate fashion. The Mg^{II} atom is located on an inversion center and is coordinated by two carbonyl O atoms and four water molecules, forming an octahedral coordination geometry. The Mg $-O_{carbonyl}$ bond distance is 2.027 (1) Å, and the average Mg-O_{water} distance is 2.080 (1) Å. The carboxy group and pyridine ring in the (2-oxo-4H-pyridin-1-yl)acetate anion are not coplanar; they form a dihedral angle of $80.8 (3)^{\circ}$. The C2-C3, C4-C5 and C1-O3 bond lengths are 1.360 (3), 1.351 (3) and 1.246 (2) Å, respectively. $O-H \cdots O$ intermolecular hydrogen bonds are formed between water molecules and the uncoordinated O atoms in the carboxylate groups of adjacent molecules, with O···O distances in the range 2.717 (2)–2.933 (2) Å and O–H···O bond angles of

metal-organic papers

175 (3)–179 (2)°, resulting in a layered structure (Table 2 and Fig. 2).

Experimental

The title complex was prepared by the addition of Mg(NO₃)₂·6H₂O (5.13 g, 20 mmol) to an aqueous solution of (2-oxo-4*H*-pyridin-1-yl)acetic acid (5.84 g, 40 mmol). The resulting solution was stirred and the pH was adjusted to 7 with 0.2 *M* NaOH solution. After evaporation at room temperature for a week, colorless single crystals were obtained from the filtered solution. Analysis calculated for C₁₄H₂₀MgN₂O₁₀: C 43.12, H 4.63, N 7.01%; found: C 41.97, H 5.03, N 6.99%.

 $D_x = 1.486 \text{ Mg m}^{-3}$

Cell parameters from 5735

2054 independent reflections

 $w = 1/[\sigma^2(F_o^2) + (0.0672P)^2]$

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

+ 0.2876P]

 $\Delta \rho_{\rm min} = -0.27 \ {\rm e} \ {\rm \AA}^{-3}$

 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.35 \text{ e } \text{\AA}^{-3}$

1839 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

reflections

 $\theta = 3.3 - 27.4^{\circ}$

 $\mu = 0.16 \text{ mm}^{-1}$

T = 293 (2) K

 $R_{\rm int} = 0.029$

 $\theta_{\rm max} = 27.5^{\circ}$

 $h = -13 \rightarrow 13$

 $\begin{array}{l} k=-8\rightarrow9\\ l=-16\rightarrow16 \end{array}$

Prism, colorless $0.38 \times 0.24 \times 0.20$ mm

Crystal data

 $\begin{bmatrix} Mg(C_7H_6NO_3)_2(H_2O)_4 \end{bmatrix} \\ M_r = 400.63 \\ Monoclinic, P_{2_1}/c \\ a = 10.550 (2) Å \\ b = 7.099 (1) Å \\ c = 13.056 (3) Å \\ \beta = 113.70 (3)^{\circ} \\ V = 895.3 (4) Å^3 \\ Z = 2 \end{bmatrix}$

Data collection

Rigaku R-AXIS RAPID diffractometer ω scans Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995) $T_{min} = 0.943, T_{max} = 0.969$ 8176 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.043$ $wR(F^2) = 0.120$ S = 1.062054 reflections 136 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Selected geometric parameters (Å, °).

Mg1 - O1W	2.060(1)	O3-C1	1.246 (2)
Mg1-O2W	2.099(1)	C2-C3	1.360 (3)
Mg1-O3	2.027(1)	C4-C5	1.351 (3)
O3-Mg1-O1W	90.97 (7)	O1W-Mg1-O2W	91.08 (5)
$O3 - Mg1 - O1W^{i}$	89.03 (7)	$O1W - Mg1 - O2W^{i}$	88.92 (5)
$O3 - Mg1 - O2W^{i}$	88.51 (5)	N1-C6-C7	111.3 (1)
O3-Mg1-O2W	91.49 (5)		

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

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$ \begin{array}{c} \hline O1W-H1W2\cdots O1^{ii} \\ O1W-H1W1\cdots O1^{iii} \\ O2W-H2W2\cdots O2^{iv} \\ O2W-H2W1\cdots O1^{i} \end{array} $	0.86 (2) 0.86 (2) 0.86 (2) 0.86 (2)	1.88 (1) 1.93 (1) 1.86 (1) 2.09 (2)	2.735 (2) 2.779 (2) 2.717 (2) 2.933 (2)	175 (3) 175 (3) 177 (2) 179 (2)
Symmetry codes: (i) -	x, -y, 1-z; (ii) $x, \frac{1}{2} - y, z - y$	$-\frac{1}{2}$; (iii) $-x, 1$	-y, 1-z; (iv)

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

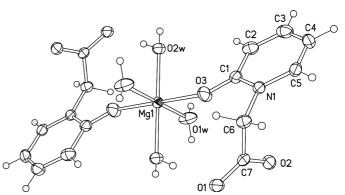


Figure 1

Structure of the title compound, showing 30% probability displacement ellipsoids for the non-H atoms.

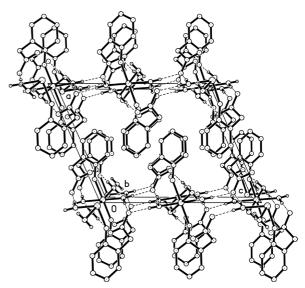


Figure 2

Packing diagram of the complex, viewed along the b axis. Hydrogen bonds are shown as dashed lines.

H atoms of the water molecules were located in difference Fourier maps and refined isotropically, with the O–H and H···H distances restrained to 0.85 (1) and 1.39 (1) Å, respectively, and with $U_{iso}(H) = 1.5U_{eq}(O)$. The other H atoms were placed in calculated positions [C-H = 0.93 and 0.97 Å] and were included in the refinement in the riding model approximation, with $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSC & Rigaku, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP*II (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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