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Key indicators

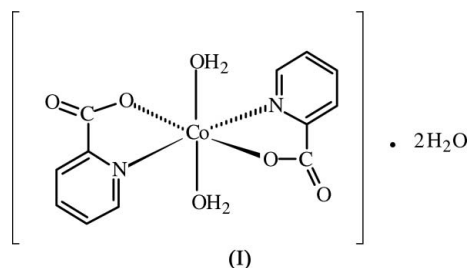
Single-crystal X-ray study
 $T = 296$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.024
 wR factor = 0.063
Data-to-parameter ratio = 14.9For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.Redetermination of *trans*-diaquabis(picolinato- $\kappa^2\text{N},\text{O}$)cobalt(II) dihydrate

The crystal structure of the title compound, $[\text{Co}(\text{C}_6\text{H}_4\text{NO}_2)_2(\text{H}_2\text{O})_2] \cdot 2\text{H}_2\text{O}$, has been reinvestigated with improved precision [previous reports: Chang *et al.* (1972). *J. Coord. Chem.* **2**, 31–34; Lumme *et al.* (1969). *Suom. Kemistil. B*, **42**, 270]. In the title compound, the Co atom is located on an inversion center and its coordination can be described as slightly distorted octahedral, equatorially *trans*-coordinated by two N and O atoms of two picolinate ligands and axially coordinated by two O atoms of the water molecules. Intermolecular O—H...O and C—H...O hydrogen-bonding interactions result in the formation of an intricate three-dimensional network.

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Comment

Picolinic acid (pyridine-2-carboxylic acid) is a well known terminal tryptophan metabolite (Mahler & Cordes, 1971). It has biological activity to induce apoptosis in leukemia HL-60 cells, which is cell death characterized by DNA fragmentation (Ogata *et al.*, 2000). Pyridine-2-carboxylic acid and its analogs have been used increasingly in the synthesis of high-nuclearity transition metallacrowns with high spin values which could be used as single-molecule magnets (Sessoli *et al.*, 1993; Thomas *et al.*, 1996; Aubin *et al.*, 1996; Gatteschi *et al.*, 2000; Winpenny, 1999; Liu *et al.*, 2001). We report here a redetermination of the crystal structure of the title compound, (I), which has been determined previously by Chang *et al.* (1972) with a higher R value (0.097) and atom H8 removed from the atom list for suspected coordinate error(s) (Cambridge Structural Database, Version 5.26; Allen, 2002), and by Lumme *et al.* (1969) with no R value given.



The complex molecule of (I) lies on an inversion center and thus the asymmetric unit contains only one half-complex. The coordination of the Co atom located on the inversion center is slightly distorted octahedral (Fig. 1). The Co atom is *trans*-coordinated by two carboxylic O atoms (O1) and two pyridine N atoms (N1) from two picolinate ligands and two O atoms (O1W) of two water molecules. The equatorial plane consists of N and O atoms from the two picolinate ligands and the axial

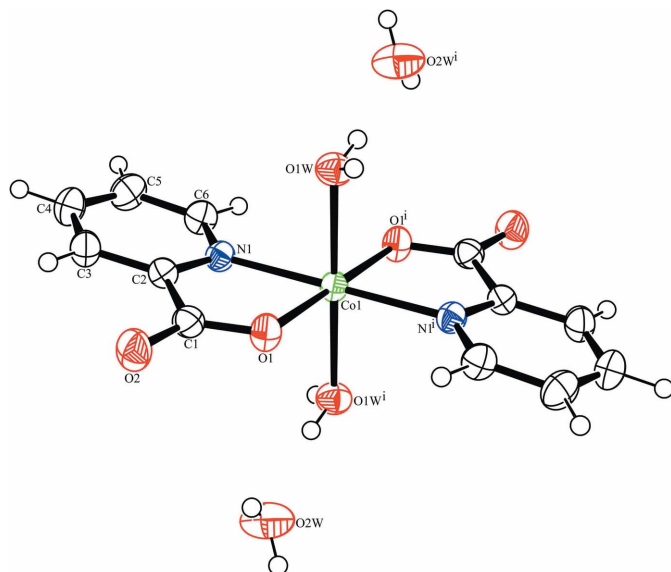


Figure 1
An ORTEP-3 (Farrugia, 1997) drawing of the structure of (I), showing the atomic numbering. Displacement ellipsoids of non-H atoms are drawn at the 50% probability level [symmetry code: (i) $1 - x, 1 - y, 1 - z$].

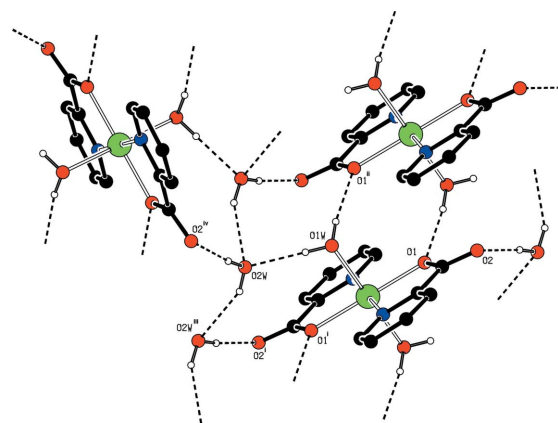


Figure 2
View showing the formation of a chain of rings through O—H...O hydrogen bonding. H atoms bonded to C atoms have been omitted. Hydrogen bonds are shown as dashed lines. [Symmetry code: (i) $1 - x, 1 - y, 1 - z$; (ii) $-x + 1, -y + 2, -z + 1$; (iii) $-x + 1, y - \frac{1}{2}, \frac{1}{2} - z$; (iv) $x, \frac{3}{2} - y, z - \frac{1}{2}$].

positions are occupied by the water molecules. The bond lengths around the metal [Co—O1 = 2.065 (1) Å, Co—N1 = 2.119 (1) Å and Co—O1W = 2.138 (1) Å] fall into normal ranges.

There is an intricate hydrogen bonding network generated by O—H...O and C—H...O interactions. Atom O1W acts as a hydrogen-bond donor to water atom O2W and to carboxylate atom O1($-x + 1, -y + 2, -z + 1$), whereas O2W acts as a hydrogen-bond donor to the symmetry-related water molecule O2W($-x + 1, y - \frac{1}{2}, \frac{1}{2} - z$) and to the second carboxylate atom O2($x, \frac{3}{2} - y, z - \frac{1}{2}$), so generating by translation a $C_3^3C(6)C_3^3C(8)C_2^2(9)R_4^3(10)R_3^3(10)$ chain of rings (Bernstein *et al.*, 1995) running parallel to the [001] direction (Table 1 and Fig. 2). In addition, atom C3 of the pyridine ring acts as a hydrogen-bond donor to the carboxylate atom

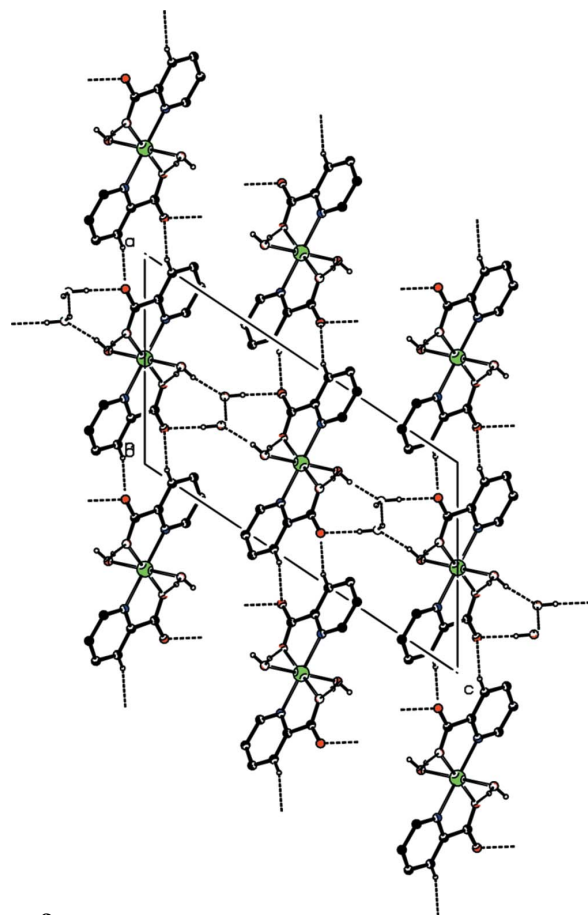


Figure 3
View down the *b* axis, showing the interconnection between the chains of rings. Hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bonding have been omitted.

O2($-x, 2 - y, 1 - z$), generating an $R_2^2(10)$ ring linking the chains into a three-dimensional network (Fig. 3).

Experimental

A solution of picolinic acid (2.4 mmol, 0.29 g) in absolute ethanol (20 ml) was added dropwise with stirring to a solution of cobalt(II) acetate tetrahydrate (1.2 mmol, 0.30 g) in absolute ethanol (20 ml). The mixture was heated to 333 K in a temperature-controlled bath and stirred for 4 h. The reaction mixture was then cooled to room temperature. The orange crystals which formed were filtered off and washed with 10 ml of water and dried in air (yield 75%, based on Co). Chemical analyses found for $C_{12}H_{16}N_2NiO_8$: C 38.41, H 4.30, N 7.47%; calculated: C 37.92, H 4.17, N 7.35%.

Crystal data

[Co(C₆H₄NO₂)₂(H₂O)₂] \cdot 2H₂O
 $M_r = 375.20$
 Monoclinic, $P2_1/c$
 $a = 9.8093$ (9) Å
 $b = 5.1846$ (3) Å
 $c = 17.5233$ (16) Å
 $\beta = 123.780$ (6) $^\circ$
 $V = 740.74$ (11) Å³
 $Z = 2$

$D_x = 1.682$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 14007
 data
 $\theta = 2.5$ – 27.9°
 $\mu = 1.20$ mm⁻¹
 $T = 296$ (2) K
 Plate, orange
 $0.37 \times 0.19 \times 0.02$ mm

Data collection

Stoe IPDS-II diffractometer	1441 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.044$
Absorption correction: integration (<i>X-RED32</i> ; Stoe & Cie, 2002)	$\theta_{\text{max}} = 27.9^\circ$
$T_{\text{min}} = 0.879$, $T_{\text{max}} = 0.961$	$h = -12 \rightarrow 12$
12035 measured reflections	$k = -6 \rightarrow 6$
1760 independent reflections	$l = -23 \rightarrow 22$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0389P)^2 + 0.0458P]$
$R[F^2 > 2\sigma(F^2)] = 0.024$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.063$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 1.03$	$\Delta\rho_{\text{max}} = 0.26 \text{ e } \text{\AA}^{-3}$
1760 reflections	$\Delta\rho_{\text{min}} = -0.25 \text{ e } \text{\AA}^{-3}$
118 parameters	
H atoms treated by a mixture of independent and constrained refinement	

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O1W-H1A \cdots O1^i$	0.847 (9)	1.892 (10)	2.7300 (17)	169.9 (19)
$O1W-H1B \cdots O2W$	0.84 (2)	1.956 (10)	2.7900 (19)	171 (2)
$O2W-H2A \cdots O2W^{ii}$	0.832 (10)	2.150 (15)	2.9264 (18)	155 (3)
$O2W-H2B \cdots O2^{iii}$	0.841 (10)	1.853 (10)	2.6891 (19)	173 (3)
$C3-H3 \cdots O2^{iv}$	0.93	2.54	3.447 (2)	166

Symmetry codes: (i) $-x+1, -y+2, -z+1$; (ii) $-x+1, y-\frac{1}{2}, -z+\frac{1}{2}$; (iii) $x, -y+\frac{3}{2}, z-\frac{1}{2}$; (iv) $-x, -y+2, -z+1$.

C-bound H atoms were positioned geometrically and were treated as riding on their parent atoms, with $C-H = 0.93 \text{ \AA}$ and $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$. The water H atoms were refined using bond restraints: $O-H = 0.85 (1) \text{ \AA}$, $H \cdots H = 1.39 \text{ \AA}$ and $U_{\text{iso}}(H) = 1.5U_{\text{eq}}(O)$.

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s)

used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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