

Gang Xie,^a Ming-Hua Zeng,^b
San-Ping Chen^a and Sheng-Li
Gao^{a*}^aDepartment of Chemistry, Northwest
University, Xi'an 710069, Shaanxi, People's
Republic of China, and ^bDepartment of
Chemistry, Guangxi Normal University, Guilin
541000, Guangxi, People's Republic of China

Correspondence e-mail: gaoshli@nwu.edu.cn

Key indicators

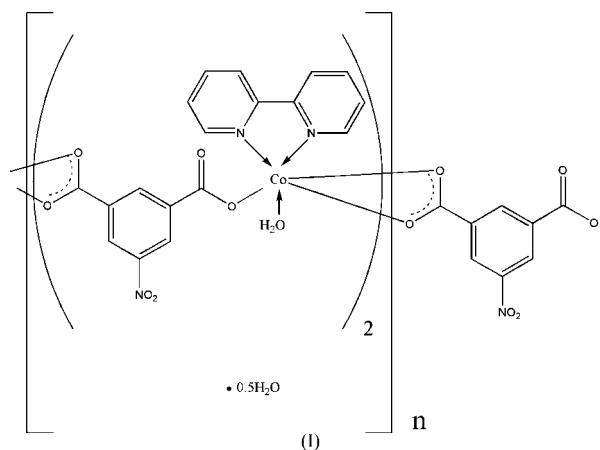
Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
Disorder in solvent or counterion
 R factor = 0.038
 wR factor = 0.112
Data-to-parameter ratio = 12.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**catena-Poly[[[aqua(2,2'-bipyridine)cobalt(II)]-
 μ -5-nitrobenzene-1,3-dicarboxylato- $\kappa^3\text{O}:\text{O}',\text{O}''$]
0.25-hydrate]**

In the title compound, $\{[\text{Co}(\text{C}_8\text{H}_3\text{NO}_6)(\text{C}_{10}\text{H}_8\text{N}_2)(\text{H}_2\text{O})] \cdot 0.25\text{H}_2\text{O}\}_n$, the coordination polyhedron of the Co^{II} ion is an octahedron. Each pair of adjacent Co^{II} ions is bridged by a dianion of 5-nitro-1,3-benzenedicarboxylic acid (H_2nmbdc) to form a chain running along the a axis. These chains are linked by π - π stacking interactions and $\text{O}(\text{water})-\text{H}\cdots\text{O}$ hydrogen bonds into a supramolecular structure.

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Comment

The dianion of 5-nitro-1,3-benzenedicarboxylic acid (H_2nmbdc) can act as a bridging ligand in a bis-monodentate coordination mode (Xiao *et al.*, 2005) or a bis-bridging coordination mode (He *et al.*, 2004). In this paper, two carboxylate groups of the nmbdc ligand coordinate in a different mode than previously reported (Xie *et al.*, 2005).



In the title compound, (I), there are two Co^{II} atoms, two 2,2'-bipyridine molecules, two nmbdc ligands, two coordinated water molecules and a solvent water half-molecule in the asymmetric unit (Fig. 1). Each pair of adjacent Co^{II} atoms is bridged by an nmbdc ligand to form a chain running along the a axis. The mode of coordination of the two carboxylate groups on each nmbdc ligand differs: one coordinates in a bidentate fashion and the other coordinates in a monodentate fashion. Each pair of adjacent chains is linked by π - π stacking interactions between the benzene rings and also $\text{O}(\text{water})-\text{H}\cdots\text{O}$ hydrogen bonds into double chains (Fig. 2 and Tables 2 and 3). The solvent water molecules are linked to the chains by $\text{O}(\text{water})-\text{H}\cdots\text{O}$ hydrogen bonds. In the crystal structure, there are many π - π stacking interactions involving the 2,2'-bipyridine ligands (Fig. 3). Geometric parameters for the π - π stacking interactions are listed in Table 3. The double chains are linked by these π - π stacking interactions into a supramolecular structure (Fig. 4).

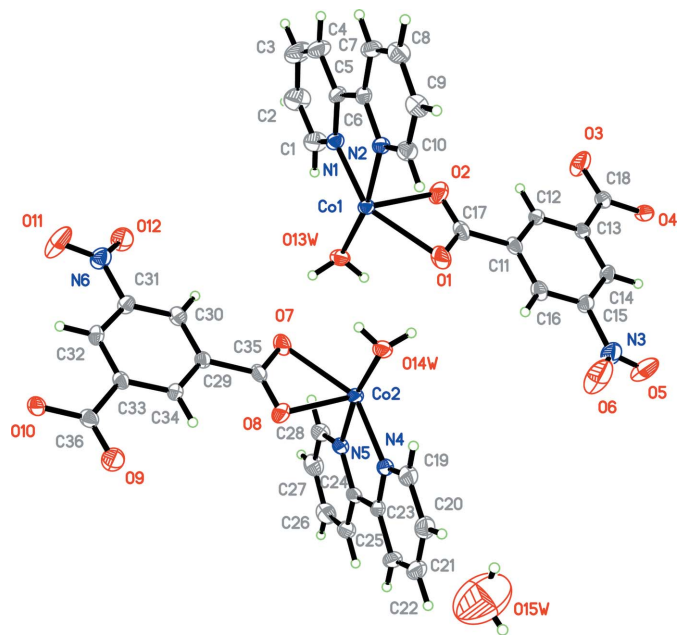


Figure 1
The asymmetric unit of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as spheres of arbitrary radii.

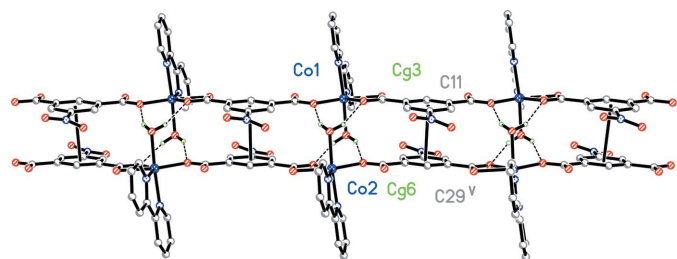


Figure 2
The double chains linked by π - π stacking interactions and hydrogen bonds along the c axis. For the sake of clarity, H atoms not involved in hydrogen bonding and solvent water molecules have been omitted [symmetry code: (v) $1 + x, y, z$].

Experimental

A mixture of cobalt nitrate hexahydrate (0.073 g, 0.25 mmol), 5-nitroisophthalic acid (0.053 g, 0.25 mmol), 2,2'-bipyridine (0.039 g, 0.25 mmol), sodium hydroxide (0.02 g, 0.5 mmol) and water (10 ml) was stirred in air for 5 min, then transferred to and sealed in a 23 ml Teflon-lined stainless steel Parr bomb, which was heated at 433 K for 120 h and then cooled to room temperature. Red block-shaped crystals were obtained after washing with deionized water (yield 31%, based on Co).

Crystal data

[Co(C₈H₃NO₆)(C₁₀H₈N₂)(H₂O)]·0.25H₂O
M_r = 446.75
 Monoclinic, $P2_1/n$
 $a = 10.0561(2) \text{ \AA}$
 $b = 23.3986(5) \text{ \AA}$
 $c = 15.3811(3) \text{ \AA}$
 $\beta = 96.317(1)^\circ$
 $V = 3597.18(13) \text{ \AA}^3$
 $Z = 8$
 $D_x = 1.650 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation
 Cell parameters from 3310 reflections
 $\theta = 2.5\text{--}23.3^\circ$
 $\mu = 1.00 \text{ mm}^{-1}$
 $T = 293(2) \text{ K}$
 Block, red
 $0.26 \times 0.14 \times 0.12 \text{ mm}$

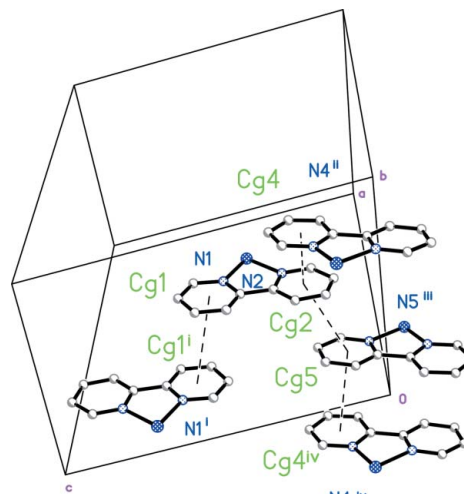


Figure 3
Part of the crystal structure of (I), showing π - π stacking interactions. For the sake of clarity, nmbdc ligands, water molecules and H atoms have been omitted [symmetry code: (i) $1 - x, -y, 1 - z$; (ii) $-\frac{1}{2} + x, \frac{1}{2} - y, -\frac{1}{2} + z$; (iii) $\frac{3}{2} - x, -\frac{1}{2} + y, \frac{1}{2} - z$; (iv) $\frac{1}{2} - x, -\frac{1}{2} + y, \frac{1}{2} - z$]

Data collection

Bruker APEX area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 2002)
 $T_{\min} = 0.780, T_{\max} = 0.889$
 28059 measured reflections
 7054 independent reflections
 4747 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.053$
 $\theta_{\max} = 26.0^\circ$
 $h = -12 \rightarrow 12$
 $k = -25 \rightarrow 28$
 $l = -18 \rightarrow 18$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.112$
 $S = 0.95$
 7054 reflections
 548 parameters
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0675P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.66 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.61 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters ($\text{\AA}, ^\circ$).

Co1—N1	2.069 (2)	Co2—N4	2.077 (2)
Co1—N2	2.108 (2)	Co2—N5	2.114 (2)
Co1—O1	2.276 (2)	Co2—O7	2.307 (2)
Co1—O2	2.199 (2)	Co2—O8	2.176 (2)
Co1—O4 ⁱ	2.048 (2)	Co2—O10 ⁱⁱ	2.040 (2)
Co1—O13W	2.135 (2)	Co2—O14W	2.115 (2)
N1—Co1—N2	77.82 (9)	N4—Co2—N5	77.77 (9)
N1—Co1—O1	146.07 (9)	N4—Co2—O8	88.41 (9)
N1—Co1—O2	88.38 (9)	N4—Co2—O14W	95.01 (9)
N1—Co1—O13W	93.44 (8)	N4—Co2—O7	145.72 (9)
N2—Co1—O1	99.16 (9)	N5—Co2—O7	97.17 (9)
N2—Co1—O2	95.02 (9)	N5—Co2—O8	94.80 (9)
N2—Co1—O13W	169.57 (9)	N5—Co2—O14W	170.88 (9)
O2—Co1—O1	57.98 (8)	O8—Co2—O7	57.94 (8)
O4 ⁱ —Co1—N1	126.38 (9)	O10 ⁱⁱ —Co2—N4	125.63 (9)
O4 ⁱ —Co1—N2	91.83 (8)	O10 ⁱⁱ —Co2—N5	91.14 (9)
O4 ⁱ —Co1—O1	87.27 (8)	O10 ⁱⁱ —Co2—O7	88.01 (8)
O4 ⁱ —Co1—O2	145.22 (9)	O10 ⁱⁱ —Co2—O8	145.90 (8)
O4 ⁱ —Co1—O13W	88.80 (8)	O10 ⁱⁱ —Co2—O14W	88.58 (8)
O13W—Co1—O1	91.28 (9)	O14W—Co2—O7	91.93 (8)
O13W—Co1—O2	90.36 (9)	O14W—Co2—O8	90.52 (9)

Symmetry codes: (i) $x - 1, y, z$; (ii) $x + 1, y, z$.

Table 2
Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O13W—H13A \cdots O7	0.853 (10)	1.99 (2)	2.741 (3)	147 (3)
O13W—H13B \cdots O10 ⁱⁱ	0.850 (10)	1.98 (2)	2.753 (3)	151 (3)
O14W—H14B \cdots O1	0.842 (10)	2.06 (2)	2.769 (3)	142 (3)
O14W—H14C \cdots O4 ⁱ	0.849 (10)	1.923 (18)	2.715 (3)	155 (3)
O15W—H15A \cdots O2 ⁱⁱⁱ	0.86	2.28	2.833 (12)	123

Symmetry codes: (i) $x - 1, y, z$; (ii) $x + 1, y, z$; (iii) $-x + \frac{3}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$.**Table 3**
Geometric parameters of the π - π stacking interactions (Å, °).

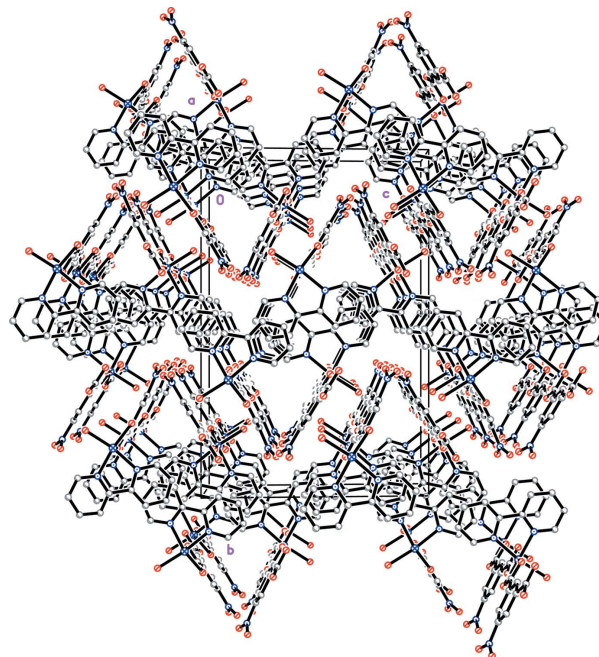
$Cg1$ is the centroid of the ring containing N1, $Cg1^i$ is the centroid of the ring containing N1ⁱ, $Cg2$ is the centroid of the ring containing N2, $Cg3$ is the centroid of the ring containing C11, $Cg4$ is the centroid of the ring containing N4ⁱⁱ, $Cg4^{iv}$ is the centroid of the ring containing N4^{iv}, $Cg5$ is the centroid of the ring containing N5ⁱⁱⁱ and $Cg6$ is the centroid of the ring containing C29^v.

Rings	Distance	Dihedral angle
$Cg1\cdots Cg1^i$	3.977 (2)	0
$Cg2\cdots Cg4$	3.583 (2)	3.57 (5)
$Cg2\cdots Cg5$	3.889 (2)	6.65 (5)
$Cg3\cdots Cg6$	3.662 (2)	4.9 (1)
$Cg5\cdots Cg4^{iv}$	3.609 (2)	3.08 (5)

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

H atoms on C atoms were positioned geometrically and refined using a riding model, with $C-H = 0.93$ Å and $U_{iso}(H) = 1.2U_{eq}(C)$. The coordinated water H atoms were located in difference Fourier maps, and were refined with distance restraints of $O-H = 0.85$ (2) Å and $H\cdots H = 1.39$ (2) Å. The uncoordinated water H atoms were located in difference Fourier maps and constrained to $O-H = 0.86$ Å, with $U_{iso}(H) = 1.2U_{eq}(O)$. The 0.5 occupancy factor results from satisfactory elemental analyses.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2002); software used to prepare material for publication: *SHELXTL*.

**Figure 4**
A packing diagram of (I). H atoms and solvent water molecules have been omitted.

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