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Key indicators

Single-crystal X-ray study
 T = 295 K
 Mean $\sigma(\text{C}-\text{C}) = 0.003 \text{ \AA}$
 R factor = 0.034
 wR factor = 0.092
 Data-to-parameter ratio = 15.2

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

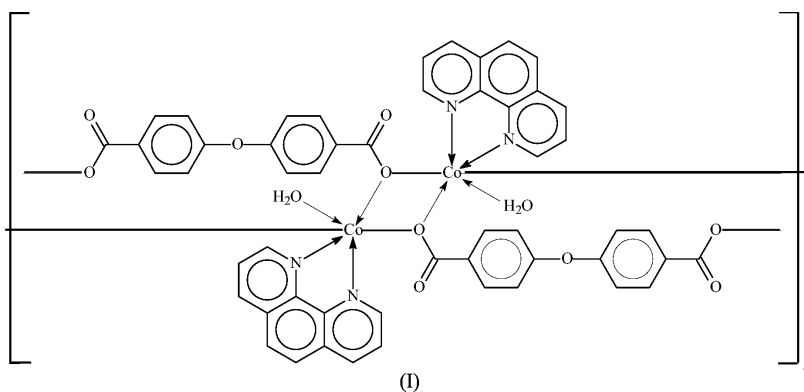
catena-Poly[bis[aqua(1,10-phenanthroline- κ^2N,N')-cobalt(II)]-di- μ -4,4'-oxydibenzoato-1:2 $\kappa^4O:O'$]

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In the crystal structure of the title compound, $[\text{Co}_2(\text{C}_{14}\text{H}_8\text{O}_5)_2(\text{C}_{10}\text{H}_8\text{N}_2)_2(\text{H}_2\text{O})_2]_n$, the dianion functions as a bridging ligand, bonding through both carboxyl $-\text{CO}_2$ end groups to link symmetry-related $[\text{Co}(\text{C}_{10}\text{H}_8\text{N}_2)(\text{H}_2\text{O})]$ entities into a ribbon structure.

Comment

In the compound $[\text{Co}(\text{C}_{14}\text{H}_8\text{O}_5)(\text{C}_{10}\text{H}_8\text{N}_2)]_n$, the bidentate bonding mode of the carboxylate dianion leads to a linear chain motif (Skakle *et al.*, 2001). The reaction of cobalt(II) acetate, 4,4'-oxybisbenzoic acid and 1,10-phenanthroline affords the related polymeric title compound, $[\text{Co}_2(\text{C}_{14}\text{H}_8\text{O}_5)_2(\text{H}_2\text{O})_2]_n$, (I), in which 1,10-phenanthroline replaces 2,2'-bipyridine. The hydrothermal synthesis yields a compound having two water molecules in the formula unit.



Although the dicarboxylate moiety of (I) links the $[\text{Co}(\text{C}_{10}\text{H}_8\text{N}_2)(\text{H}_2\text{O})]$ entities into a linear chain, as in the 2,2'-bipyridine compound, in (I) the two carboxyl $-\text{CO}_2$ fragments are only monodentate in the chain. The O atom of one repeat unit bonds to a Co atom of a symmetry-related unit, completing the distorted octahedral Co environment (Fig. 1). This gives rise to the formation of a ribbon structure (Fig. 2).

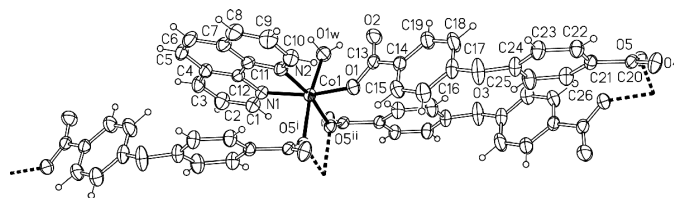


Figure 1
 A plot illustrating the octahedral geometry of the Co atom in a fragment of the ribbon structure of (I). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii. [Symmetry codes: (i) $1 - x, 2 - y, 1 - z$; (ii) $x, y, z - 1$].

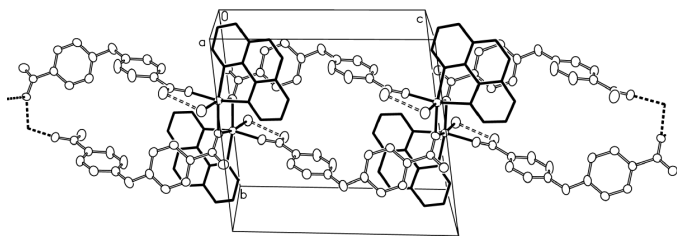


Figure 2

An illustration of the ribbon structure of (I). Dashed lines indicate hydrogen bonds.

The ribbons are linked into layers by O—H...O hydrogen bonds (Table 2).

Experimental

Cobalt(II) acetate tetrahydrate (0.125 g, 0.5 mmol), 4,4'-oxybisbenzoic acid (0.129 g, 0.5 mmol) and 1,10-phenanthroline (0.180 g, 1.0 mmol) were placed in a 30 ml Teflon-lined stainless-steel Parr bomb, together with water (20 ml). The bomb was heated at 423 K for 6 d and then cooled slowly to room temperature to furnish red crystals of (I).

Crystal data

[Co₂(C₁₄H₈O₅)₂(C₁₂H₈N₂)₂(H₂O)₂]
M_r = 1026.71
 Triclinic, *P*1
a = 7.7129 (4) Å
b = 11.5063 (6) Å
c = 13.3854 (7) Å
 α = 82.614 (1)°
 β = 83.165 (1)°
 γ = 72.724 (1)°
V = 1120.8 (1) Å³

Z = 1
D_x = 1.521 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 5331 reflections
 θ = 2.5–28.3°
 μ = 0.81 mm⁻¹
T = 295 (2) K
 Prism, red
 0.34 × 0.27 × 0.23 mm

Data collection

Bruker APEX CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 2002)
T_{min} = 0.770, *T_{max}* = 0.835
 9587 measured reflections

4919 independent reflections
 4510 reflections with *I* > 2σ(*I*)
R_{int} = 0.014
 θ_{\max} = 27.5°
h = -10 → 10
k = -14 → 14
l = -17 → 17

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.034
wR(*F*²) = 0.092
S = 1.03
 4919 reflections
 324 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0553P)^2 + 0.3018P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.40 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.30 \text{ e } \text{Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Co1—O1	2.053 (1)	Co1—O1 <i>W</i>	2.094 (1)
Co1—O5 ⁱⁱⁱ	2.108 (1)	Co1—N1	2.121 (1)
Co1—O5 ^{iv}	2.122 (1)	Co1—N2	2.161 (1)
O1—Co1—O5 ⁱⁱⁱ	91.30 (5)	O5 ⁱⁱⁱ —Co1—N2	94.16 (5)
O1—Co1—O5 ^{iv}	98.51 (5)	O5 ^{iv} —Co1—O1 <i>W</i>	88.92 (5)
O1—Co1—O1 <i>W</i>	88.10 (5)	O5 ^{iv} —Co1—N1	96.53 (5)
O1—Co1—N1	164.66 (5)	O5 ^{iv} —Co1—N2	169.23 (5)
O1—Co1—N2	88.34 (5)	O1 <i>W</i> —Co1—N1	89.18 (6)
O5 ⁱⁱⁱ —Co1—O5 ^{iv}	77.47 (5)	O1 <i>W</i> —Co1—N2	99.66 (6)
O5 ⁱⁱⁱ —Co1—O1 <i>W</i>	166.14 (5)	N1—Co1—N2	77.24 (6)
O5 ⁱⁱⁱ —Co1—N1	94.90 (5)		

Symmetry codes: (iii) *x*, *y*, *z* + 1; (iv) 1 - *x*, 1 - *y*, 1 - *z*.

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>
O1 <i>W</i> —H1 <i>W</i> 1...O2	0.85 (1)	1.86 (2)	2.675 (2)	161 (3)
O1 <i>W</i> —H1 <i>W</i> 2...O4 ^v	0.84 (1)	1.93 (1)	2.764 (2)	175 (2)

Symmetry codes: (v) *x* + 1, *y*, *z* + 1.

H atoms bonded to C atoms were included in the refinement in calculated positions in the riding-model approximation, with C—H = 0.93 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C). The water H atoms were located and refined with a distance restraint of 0.85 (1) Å and with *U*_{iso}(H) = 1.2*U*_{eq}(O).

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINTE* (Bruker, 2002); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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