

Rerefinement of *catena*-poly[[tetraaquacobalt(II)- μ -pyrazine] phthalate] in a lower-symmetry space group

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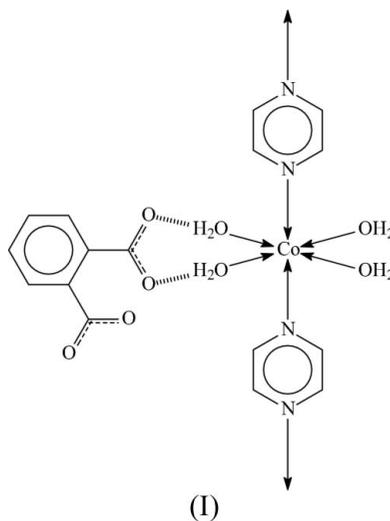
Key indicators

Single-crystal X-ray study
 $T = 298$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.023
 wR factor = 0.074
Data-to-parameter ratio = 11.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

The polymeric title compound, $[\{\text{Co}(\text{C}_4\text{H}_4\text{N}_2)(\text{H}_2\text{O})_4\}(\text{C}_8\text{H}_4\text{O}_4)]_n$, when refined in *Imm2* from the diffraction intensities originally used for refinement in *Immm* [Yang, Li, Cao & Yao (2003). *Acta Cryst. E* **59**, m961–m963], is an ordered structure whose Co^{II} atom, pyrazine ligand and phthalate counter-ion lie on special positions. The metal atom and *N*-heterocycle have *mm2* site symmetry and the dianion *m* site symmetry.

Comment

For diffraction measurements that cannot be unambiguously assigned to a particular space group, the structure should preferably be refined in a centrosymmetric space group, even though the structure is disordered in this higher-symmetry setting (Marsh, 1986). As such, the structure of $[\text{Co}(\text{H}_2\text{O})_4(\text{C}_4\text{H}_4\text{N}_2)](\text{C}_8\text{H}_4\text{O}_4)$, (I), was originally refined with disorder in *Immm*; a number of restraints were applied to the model, and the high residual index ($R_1 = 0.096$) was attributed to disorder and not to low quality of the measurements (Yang *et al.*, 2003).



The zinc analogue crystallizes with similar cell dimensions; the authors of that study refined the structure in *Imm2* because the compound exhibited second-harmonic generation (SHG) (Zhang *et al.*, 2005); the SHG test is one which is able to distinguish between centrosymmetric and non-centrosymmetric structures. Refinement of the cobalt compound in this space group led to a significantly improved *R* index ($R_1 = 0.023$). In this setting, the Co and pyrazine lie on special positions of *mm2* site symmetry and the phthalate on a special position of *m* site symmetry (Fig. 1). The phthalate is ordered; the anions surround the linear polycationic chain and interact with the chains through hydrogen bonds (Fig. 2 and Table 2).

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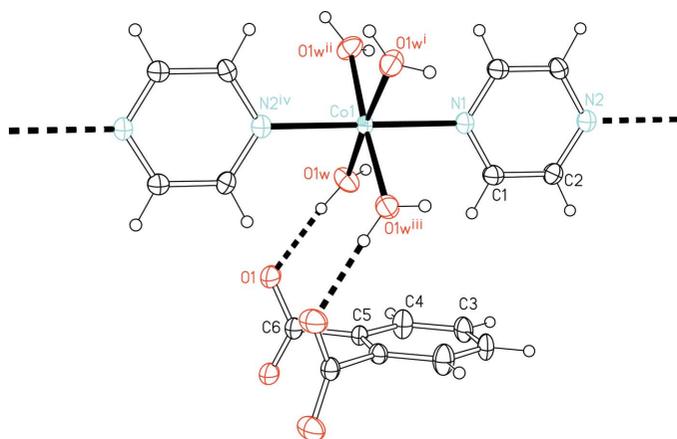


Figure 1
A plot of (I). Displacement ellipsoids are drawn at the 50% probability level. H atoms are drawn as spheres of arbitrary radii. [Symmetry codes: (i) $-x, 1 - y, z$; (ii) $-x, y, z$; (iii) $x, 1 - y, z$; (iv) $x, y, 1 + z$.]

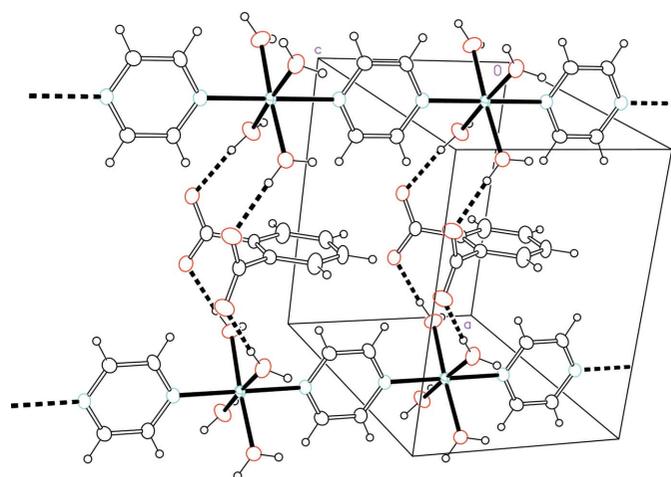


Figure 2
A plot depicting the hydrogen bonds (dashed lines).

Experimental

Crystal data

[Co(C₄H₄N₂)(H₂O)₄](C₈H₄O₄)
M_r = 375.20
 Orthorhombic, *Imm*2
a = 9.4033 (6) Å
b = 10.2886 (7) Å
c = 7.1791 (5) Å
V = 694.55 (8) Å³
Z = 2
D_x = 1.794 Mg m⁻³

Mo *K*α radiation
 Cell parameters from 2933 reflections
 θ = 2.9–28.4°
 μ = 1.28 mm⁻¹
T = 298 (2) K
 Prism, orange
 0.45 × 0.28 × 0.27 mm

Data collection

Bruker APEX area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.545, *T_{max}* = 0.723
 2928 measured reflections

871 independent reflections
 871 reflections with *I* > 2σ(*I*)
R_{int} = 0.018
 θ_{max} = 27.5°
h = -12 → 11
k = -13 → 13
l = -9 → 9

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.023
wR(*F*²) = 0.074
S = 1.25
 871 reflections
 75 parameters
 All H-atom parameters refined

$w = 1/[\sigma^2(F_o^2) + (0.0542P)^2 + 0.0753P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 (Δ/σ)_{max} = 0.001
 $\Delta\rho_{\text{max}}$ = 0.35 e Å⁻³
 $\Delta\rho_{\text{min}}$ = -0.51 e Å⁻³
 Absolute structure: Flack (1983),
 391 Friedel pairs
 Flack parameter: 0.02 (1)

Table 1

Selected geometric parameters (Å, °).

Co1—O1w	2.051 (1)	Co1—N2 ^{iv}	2.188 (4)
Co1—N1	2.194 (4)		
O1w—Co1—O1w ⁱ	176.3 (1)	O1w—Co1—N1	88.1 (1)
O1w—Co1—O1w ⁱⁱ	87.9 (1)	O1w—Co1—N2 ^{iv}	91.9 (1)
O1w—Co1—O1w ⁱⁱⁱ	92.0 (1)		

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

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>
O1w—H1w1...O1	0.84 (1)	1.87 (1)	2.709 (2)	174 (4)
O1w—H1w2...O1 ^v	0.83 (1)	1.91 (1)	2.736 (2)	178 (3)

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

The C-bound H atoms were generated geometrically (C—H = 0.93 Å) and were treated as riding, with *U*_{iso}(H) parameters set at 1.2*U*_{eq}(C). The two O-bound H atoms were refined with a distance restraint of O—H = 0.85 (1) Å; the *U*_{iso}(H) parameters were freely refined.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINTE* (Bruker, 2001); data reduction: *SAINTE*; method used to solve structure: atomic coordinates taken from the Zn analogue; 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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