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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and least-squares-planes data have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 71374 (29 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: SH1051]

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Structure of a Tripotassium Pyrroloquinoline Quinone Salt

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Abstract

In $3K^+ \cdot C_{14}H_3N_2O_8^{3-} \cdot 4H_2O$, all three carboxyl groups of pyrroloquinoline quinone (PQQ; 4,5-dihydro-4,5-dioxo-1*H*-pyrrolo[2,3-*f*]quinoline-2,7,9-tricarboxylic acid) are in an anion form to neutralize the three potassium cations. Each of the three potassium ions is closely coordinated to polar atoms of the PQQ^{3-} ions and waters of crystallization. Noticeable structural features within the crystal unit are the stacking of the PQQ^{3-} ions and the extensive network of hydrogen-bonding interactions.

Comment

PQQ is known as the coenzyme of a novel class of oxidoreductases (Duine, Frank Jzn & Verwiël, 1980; Duine & Frank Jzn, 1981). To understand its biological behavior, it is important to reveal the molecular structure and atomic charge distribution in the molecule. The structures of the 5-(2-oxopropyl) (Salisbury, Forrest, Cruse & Kennard, 1979; Cruse, Kennard & Salisbury, 1980) and 5-(2,4-dinitrophenylhydrazine) adducts of PQQ (van Koningsveld, Jansen, Jongejan, Frank Jzn & Duine, 1985) and the PQQ disodium salt (Ishida *et al.*, 1989) have been reported already. Ishida *et al.* (1989) demonstrated that the PQQ molecule possesses two sodium-binding sites. This is the first report on the crystal structure of the trianionic form of PQQ.

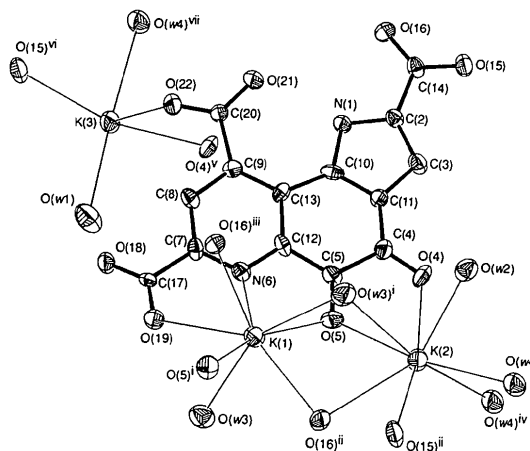


Fig. 1. Perspective view of $K_3(C_{14}H_3N_2O_8) \cdot 4H_2O$ indicating the atom labeling. H atoms are omitted.

Fig. 1 shows a perspective drawing of the structure and indicates the atom-numbering scheme. In an earlier structural study of the PQQ disodium salt, PQQ formed dianions (PQQ^{2-}) (Ishida *et al.*, 1989). The charge on PQQ probably depends on the synthetic conditions. Since the present PQQ crystals were prepared under neutral conditions, the PQQ molecule formed trianions (PQQ^{3-}) with three carboxylate anion substituents. A comparison of the structures of PQQ^{3-} and PQQ^{2-} reveals very similar bond lengths and angles. The carboxylate C(2)—C(14) bond length of 1.498 (8) Å is also significantly shorter than the other two carboxylate C—C bond lengths [1.531 (8) and 1.552 (8) Å for C(7)—C(17) and C(9)—C(20), respectively]. This fact presumably results from the energetic stabilization of the carboxylate group through a resonance effect with the respective aromatic rings (Ishida *et al.*, 1989). Differences between PQQ^{2-} and PQQ^{3-} are manifest in the C(4)=O(4) bond lengths [1.204 (5) and 1.205 (6) Å in PQQ^{2-} (Ishida *et al.*, 1989) and 1.234 (8) Å in PQQ^{3-}]. The longer bond length in

PQQ³⁻ is due to the coordination of the O(4) atom to K(2) and K(3) ions.

The crystal structure analysis of the PQQ disodium salt demonstrated that the PQQ molecule possesses two metal-binding sites: a primary site in the pocket formed by the O(5), N(6) and O(19) atoms and a secondary site in the pocket formed by the O(4) and O(5) atoms (Ishida *et al.*, 1989). In the PQQ tripotassium salt, the two potassium cations [K(1) and K(2)] are also located at these sites. The other potassium cation [K(3)] is located at a novel site and is five-coordinate, with two PQQ³⁻ ions binding to it as monodentate ligands through O(4) and O(15), respectively. It seems that the location of the K(3) ion is favorable in view of the crystal packing.

The most interesting features in the crystal structure are the extensive network of hydrogen-bonding interactions and the stacking of the PQQ³⁻ ions. The stacking PQQ³⁻ ions form columns aligned along the crystallographic *b* axis. The closest interatomic distance between the PQQ³⁻ ions is 3.416 (8) Å {C(11)⋯C(13')} [symmetry code: (v) $-x, 1-y, 1-z$]. The hydrogen-bonding network and the stacking of the PQQ³⁻ ions play an essential role in stabilizing this crystal structure.

Experimental

Crystal data

K₃(C₁₄H₃N₂O₈)·4H₂O

M_r = 516.54

Monoclinic

*P*2₁/*n*

a = 17.060 (2) Å

b = 7.088 (2) Å

c = 15.332 (2) Å

β = 91.76 (1)°

V = 1853.1 (7) Å³

Z = 4

D_x = 1.852 Mg m⁻³

D_m = 1.858 Mg m⁻³

Cu *K*α radiation

λ = 1.54178 Å

Cell parameters from 20 reflections

θ = 18–20°

μ = 7.33 mm⁻¹

T = 298 K

Needles

0.42 × 0.12 × 0.10 mm

Dark red

Crystal source: crystallization from H₂O/CH₃CN

Data collection

Rigaku AFC-5R diffractometer

ω -2 θ scans

Absorption correction: empirical

T_{min} = 0.7502, *T_{max}* = 0.9940

3364 measured reflections

2957 independent reflections

2127 observed reflections

[*I* > 3 σ (*I*)]

R_{int} = 0.037

θ_{\max} = 62.5°

h = -20 → 20

k = 0 → 8

l = 0 → 18

3 standard reflections monitored every 100 reflections

intensity variation: 4%

Refinement

Refinement on *F*

R = 0.056

$w = 1/(\sigma^2 F + 0.0648|F| + 0.0002|F|^2)$

wR = 0.060

S = 1.23

2127 reflections

325 parameters

All H-atom parameters refined

(Δ/σ)_{max} = 0.33

$\Delta\rho_{\max}$ = 0.88 e Å⁻³

$\Delta\rho_{\min}$ = -0.80 e Å⁻³

Atomic scattering factors

from *International Tables for X-ray Crystallography* (1974, Vol. IV)

Table 1. Fractional atomic coordinates and equivalent isotropic thermal parameters (Å²)

$$B_{\text{eq}} = \frac{4}{3} \sum_i \sum_j \beta_{ij} a_i \cdot a_j$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>B_{eq}</i>
K(1)	-0.26077 (7)	0.1295 (2)	0.27651 (8)	2.36
K(2)	-0.08005 (7)	0.3300 (3)	0.10593 (8)	2.75
K(3)	-0.14302 (7)	0.3644 (3)	0.84090 (8)	2.83
O(4)	0.0340 (3)	0.4260 (7)	0.2566 (3)	3.41
O(5)	-0.1207 (3)	0.3875 (7)	0.2823 (3)	2.90
O(15)	0.2987 (3)	0.2586 (7)	0.4944 (3)	3.21
O(16)	0.2408 (3)	0.1914 (6)	0.6194 (3)	2.36
O(18)	-0.3297 (2)	0.2069 (7)	0.5826 (3)	2.53
O(19)	-0.3417 (2)	0.2014 (6)	0.4377 (3)	2.51
O(21)	0.0268 (3)	0.0846 (8)	0.6673 (3)	3.77
O(22)	-0.0886 (3)	0.0998 (7)	0.7261 (3)	3.07
N(1)	0.0940 (3)	0.2340 (7)	0.5360 (3)	1.98
N(6)	-0.1822 (3)	0.2693 (7)	0.4301 (3)	1.91
C(2)	0.1632 (3)	0.2718 (8)	0.4949 (4)	1.86
C(3)	0.1447 (3)	0.3298 (9)	0.4117 (4)	2.28
C(4)	0.0135 (3)	0.3713 (9)	0.3289 (4)	2.21
C(5)	-0.0754 (3)	0.3499 (9)	0.3419 (4)	2.11
C(7)	-0.2134 (3)	0.2193 (8)	0.5042 (4)	1.94
C(8)	-0.1671 (3)	0.1723 (8)	0.5771 (4)	2.01
C(9)	-0.0856 (3)	0.1798 (8)	0.5765 (4)	1.83
C(10)	0.0317 (3)	0.2656 (8)	0.4801 (4)	1.96
C(11)	0.0620 (3)	0.3279 (8)	0.4024 (4)	1.76
C(12)	-0.1041 (3)	0.2835 (8)	0.4270 (4)	1.90
C(13)	-0.0511 (3)	0.2402 (8)	0.4998 (4)	1.71
C(14)	0.2408 (3)	0.2384 (9)	0.5408 (4)	1.98
C(17)	-0.3028 (3)	0.2094 (8)	0.5083 (4)	1.60
C(20)	-0.0442 (3)	0.1191 (8)	0.6636 (4)	1.78
O(W1)	-0.2705 (3)	0.4165 (8)	0.7291 (3)	4.48
O(W2)	0.0408 (3)	0.0637 (7)	0.1189 (3)	3.07
O(W3)	-0.3640 (3)	0.4671 (7)	0.3025 (3)	3.33
O(W4)	-0.0056 (3)	0.2926 (7)	-0.0550 (3)	3.04

Table 2. Selected geometric parameters (Å)

K(1)—O(5)	3.008 (5)	O(16)—C(14)	1.250 (7)
K(1)—O(19)	2.913 (4)	O(18)—C(17)	1.242 (7)
K(1)—N(6)	2.852 (5)	O(19)—C(17)	1.253 (7)
K(1)—O(W3)	3.005 (5)	O(21)—C(20)	1.236 (8)
K(1)—O(5 ⁱⁱ)	2.781 (5)	O(22)—C(20)	1.248 (7)
K(1)—O(16 ⁱⁱⁱ)	2.724 (4)	N(1)—C(2)	1.383 (7)
K(1)—O(16 ⁱⁱⁱ)	2.794 (4)	N(1)—C(10)	1.362 (7)
K(1)—O(W3 ³)	2.736 (5)	N(6)—C(7)	1.317 (7)
K(2)—O(4)	3.052 (5)	N(6)—C(12)	1.339 (7)
K(2)—O(5)	2.842 (5)	C(2)—C(3)	1.368 (8)
K(2)—O(W2)	2.797 (5)	C(2)—C(14)	1.498 (8)
K(2)—O(W4)	2.821 (5)	C(3)—C(11)	1.414 (8)
K(2)—O(15 ⁱⁱ)	2.716 (5)	C(4)—C(5)	1.544 (9)
K(2)—O(16 ⁱⁱⁱ)	3.073 (4)	C(4)—C(11)	1.411 (8)
K(2)—O(W3 ³)	3.095 (5)	C(5)—C(12)	1.484 (8)
K(2)—O(W4 ^{iv})	3.158 (5)	C(7)—C(8)	1.389 (8)
K(3)—O(22)	2.752 (5)	C(7)—C(17)	1.531 (8)
K(3)—O(W1)	2.752 (5)	C(8)—C(9)	1.391 (8)
K(3)—O(4 ^v)	2.841 (5)	C(9)—C(13)	1.399 (7)
K(3)—O(15 ^{vi})	2.726 (5)	C(9)—C(20)	1.552 (8)
K(3)—O(W4 ^{vii})	2.841 (5)	C(10)—C(11)	1.385 (8)
O(4)—C(4)	1.234 (8)	C(10)—C(13)	1.465 (8)
O(5)—C(5)	1.209 (8)	C(12)—C(13)	1.448 (8)
O(15)—C(14)	1.244 (7)		

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

PQQ was purchased from Ube Industries (Tokyo, Japan). PQQ was dissolved in KOH solution and the pH of the solution was then adjusted to 7.0 by the addition of HCl solution. The addition of CH₃CN to the solution yielded the crystals in a few weeks. The density was measured by flotation in CCl₄-CH₂BrCH₂CH₂Br. A crystal was sealed in a glass capillary for the collection of the diffraction data. The unit-cell parameters indicated a monoclinic crystal system and systematic absences indicated the space group. Lattice constants were obtained by least-squares refinement. Intensity data were collected with an ω -scan width of $(1.50 + 0.15 \tan \theta)^\circ$ and a scan speed of $4.0^\circ \text{ min}^{-1}$. Data were corrected for Lorentz and polarization effects and for absorption.

The structure was solved by direct methods using *MULTAN78* (Main *et al.*, 1978) and refined by block-diagonal least squares using the program *HBL5-V* (Ashida, 1979). Anisotropic thermal parameters were refined for non-H atoms. The positions of the H atoms of PQQ were calculated on the basis of their molecular geometry. Difference Fourier syntheses calculated at intermediate stages of refinement showed maxima consistent with the expected positions of H atoms of water. H atoms were included with isotropic temperature factors in subsequent refinements. All computations were carried out on an ACOS 2020 computer at the Computation Center, Osaka University, and on an ACOS S-850 computer at the Crystallographic Research Center, Institute for Protein Research, Osaka University.

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Lists of structure factors, anisotropic thermal parameters, H-atom coordinates and complete geometry have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 71370 (14 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: OH1023]

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Dimeric Copper(II) 2-Methyl-2-phenylpropanoate Adducts with Water or 2,6-Lutidine and a Monomeric Copper(II) Triphenylacetate Adduct with 2,6-Lutidine

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Abstract

The structures of diaquatetrakis(μ -2-methyl-2-phenylpropanoato-*O*:*O'*)dicopper(II), [Cu(C₁₀H₁₁O₂)₂(H₂O)]₂ (I), bis(2,6-dimethylpyridine)tetrakis(μ -2-methyl-2-phenylpropanoato-*O*:*O'*)dicopper(II) benzene solvate, [Cu(C₁₀H₁₁O₂)₂(C₇H₆N)]₂·C₆H₆ (II) and *trans*-bis(2,6-dimethylpyridine)bis(triphenylacetato-*O*:*O'*)copper(II), [Cu(C₂₀H₁₅O₂)₂(C₇H₉N)₂] (III), have been determined by single-crystal X-ray diffraction. (I) and (II) are dinuclear Cu^{II} complexes having cage structures and the coordination geometries around the Cu atoms are square pyramidal in (I) and distorted trigonal bipyramidal in (II). The Cu...Cu distances are 2.589 (1) and 2.907 (1) Å for (I) and (II), respectively. The coordination geometry in the monomeric copper complex (III) is square planar. The $-2J$ values of (I) and (II) are 348 and 310 cm⁻¹, respectively.

Comment

The antiferromagnetic interaction in dimeric copper(II) carboxylates decreases as the coordination geometry around the Cu atoms is distorted from square pyramidal (SP) toward trigonal bipyramidal (TBP) (Uekusa *et al.*, 1992). The distorted TBP