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Key indicators

Single-crystal X-ray study T = 299 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.040 wR factor = 0.093 Data-to-parameter ratio = 13.8

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

Trisodium citrate dihydrate

From a solution containing uranyl nitrate and sodium citrate, single crystals of trisodium citrate dihydrate, $3Na^+ \cdot C_6H_5O_7{}^{3-} \cdot 2H_2O$, were obtained. The structure consists of a complex network of citrate and sodium ions. Additionally, hydrogen bonds between the citrate ions and the water of crystallization are formed.

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Comment

Citric acid is a tricarboxylic acid, which can undergo different degrees of deprotonation and thus can form different reaction products with alkali hydroxide, depending on the molar ratio. Additionally, the amount of water of crystallization may vary depending on the stoichiometry and concentration of the different species in the reaction medium. The unit cells and space groups of a number of sodium and potassium citrates have been known for a long time (Burns & Iball, 1954). However, no structure determination of a sodium citrate dihydrate has been undertaken. Later, the structures of two other hydrates of sodium citrate were determined, *viz.* sodium dihydrogen citrate (Glusker *et al.*, 1965) and trisodium citrate 5.5-hydrate (Viossat *et al.*, 1986).



In our investigations of the uranyl/citrate system, we obtained single crystals of trisodium citrate dihydrate, (I), whose structure has been determined. It contains one citrate anion, whose geometry is unexceptional, and three Na⁺ ions in the asymmetric unit (Fig. 1). The Na⁺ ions are coordinated by both citrate ions and water molecules, yielding a coordination number of 6, with rather irregular coordination polyhedra (Fig. 1). Each citrate ion coordinates to nine Na⁺ ions as a bridging ligand, yielding a complex three-dimensional network. Additionally, the water ligands form hydrogen bonds to some of the carboxyl O atoms (Fig. 2). The coordination of the citrate ion is shown in Fig. 1, while Fig. 3 shows the unit-cell contents.

Experimental

To a solution of uranyl nitrate hexahydrate and trisodium citrate in a molar ratio of 1:3, sodium hydroxide was added until a pale-yellow product started to precipitate. From this solution, colourless crystals of the title compound were obtained upon standing for one month.

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Figure 1

The citrate ion and the three Na⁺ ions with their complete coordination environments in the structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level. [Symmetry codes: (i) x, $1 - y, z - \frac{1}{2}$; (ii) 1 - x, 1 - y, -z; (iii) $\frac{1}{2} - x, \frac{1}{2} - y, -z$; (iv) $1 - x, y, \frac{1}{2} - z$; (vi) 1 - x, -y, -z.]



Figure 2

The citrate ion and the surrounding $Na^{\scriptscriptstyle +}$ ions and water molecules. Hydrogen bonds are shown as dashed lines.

Crystal data

3Na ⁺ ·C ₆ H ₅ O ₇ ³⁻ ·2H ₂ O $M_r = 294.10$ Monoclinic, C2/c a = 15.7044 (4) Å b = 12.5010 (4) Å c = 11.2837 (4) Å $\beta = 103.5841$ (13)° V = 2153.26 (12) Å ³	$D_x = 1.814 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 3843 reflections $\theta = 4.5-27.5^{\circ}$ $\mu = 0.27 \text{ mm}^{-1}$ T = 299 K Cuboid colourless
Z = 8	$0.20 \times 0.15 \times 0.10 \text{ mm}$
Data collection	
Bruker–Nonius KappaCCD diffractometer	1643 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.055$ $\theta_{\text{int}} = 27.5^{\circ}$
Absorption correction: none	$h = -17 \rightarrow 20$
8459 measured reflections	$k = -14 \rightarrow 16$
2451 independent reflections	$l = -14 \rightarrow 12$





The unit cell contents, viewed along b. H atoms have been omitted.

Refinement

 $\begin{array}{ll} \mbox{Refinement on } F^2 & w = 1/[\sigma^2(F_o^2) + (0.0385P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.040 & w + 0.6276P] \\ wR(F^2) = 0.093 & where $P = (F_o^2 + 2F_c^2)/3$ \\ S = 1.02 & (\Delta/\sigma)_{max} < 0.001 \\ 2451 \ reflections & \Delta\rho_{max} = 0.43 \ e^{\begin{array}{c} A^{-3} \\ max = 0.43 \ e^{\begin{array}{c} A^{-3} \\ max = -0.32 \ e^{\b$

Table 1

Selected geometric parameters (Å).

Na1-O8 ⁱ	2.3404 (18)	Na2-O6	2.3819 (15)
Na1-O1	2.3442 (16)	Na2-O8	2.450 (2)
Na1-O6 ⁱⁱ	2.3621 (16)	Na2-O1	2.4588 (16)
Na1-O9 ⁱⁱⁱ	2.3819 (17)	Na2-Na3	3.3241 (13)
Na1-O5	2.4627 (17)	Na2-Na2 ^{iv}	3.5644 (15)
Na1-O5 ⁱⁱ	2.5492 (18)	Na2-Na3 ⁱⁱⁱ	4.2172 (12)
Na1-C6 ⁱⁱ	2.775 (2)	Na3–O4 ^{iv}	2.3346 (17)
Na1-Na2 ⁱ	3.3240 (12)	Na3-O7	2.3947 (15)
Na1–Na3 ⁱⁱⁱ	3.4123 (11)	Na3-O1 ⁱⁱⁱ	2.3999 (15)
Na1–Na1 ⁱⁱ	3.4358 (16)	Na3-O9	2.4013 (19)
Na2-O2 ⁱⁱⁱ	2.3163 (16)	Na3-O2 ⁱⁱⁱ	2.6184 (17)
Na2-O6 ^{iv}	2.3342 (16)	Na3-O3 ^v	2.781 (2)

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

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

$D-\mathrm{H}\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$07-H7\cdots 04$	0.82 (2)	1.85 (2)	2.6072 (19)	153 (2)
$08-H8A\cdots 05^{vi}$	0.803 (15)	2.030 (16)	2.820 (2)	168 (3)
$08-H8B\cdots 03^{vi}$	0.840 (15)	1.920 (16)	2.755 (2)	173 (2)
$09-H9A\cdots 03^{v}$	0.808 (15)	1.974 (17)	2.706 (2)	150 (2)
$09-H9B\cdots 02^{vii}$	0.810 (16)	1.953 (16)	2.762 (2)	176 (3)

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

H atoms attached to carbon were positioned geometrically (C–H = 0.97 Å) and refined in the riding-model approximation, with $U_{\rm iso}$ equal to $1.2U_{\rm eq}$ of the carrier atom. The geometry of the water

molecules were restrained to a reasonable chemical model $[O-H = 0.82 (1) \text{ Å} \text{ and } H \cdots H = 1.37 (1) \text{ Å}]$. The H-atom coordinates were then allowed to refine, with U_{iso} equal to $1.2U_{eq}$ of the O atom.

Data collection: *KappaCCD Software* (Nonius, 1997); cell refinement: *HKL SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* (Otwinowski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SIR*97 (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *DIAMOND* (Brandenburg, 2001) and *ORTEP*-3 (Farrugia, 1997); software used to prepare material for publication: *maXus* (Mackay *et al.*, 1997).

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