

Poly[[diaquabis(μ_3 -maleato- κ^4 O¹:O^{1'},O⁴:-O^{4'})dicopper(II)] trihydrate]

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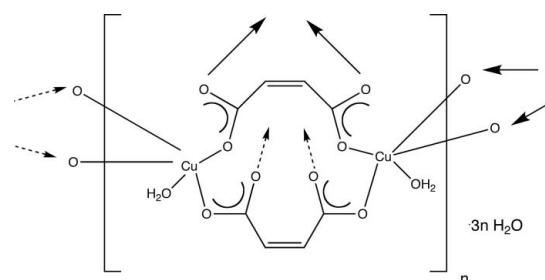
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Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(C-C) = 0.004$ Å; R factor = 0.023; wR factor = 0.055; data-to-parameter ratio = 11.1.

In the title compound, $\{[Cu_2(C_4H_2O_4)_2(H_2O)_2]\cdot 3H_2O\}_n$, Cu^{II} ions with square-pyramidal coordination are bridged by exot Tridentate maleate dianions into $[Cu_2(maleate)_2(H_2O)_2]_n$ layers coincident with the bc crystal plane. The interlamellar regions contain hydrogen-bonded cyclic water hexamers which facilitate layer stacking into a pseudo-three-dimensional crystal structure. The water hexamers themselves are formed by the operation of crystallographic inversion centers on sets of three crystallographically distinct water molecules of hydration.

Related literature

For recent dpa coordination polymers, see: Brown *et al.* (2008). For the preparation of dpa, see: Zapf *et al.* (1998). For the determination of the τ factor for five-coordinate geometries, see: Addison *et al.* (1984).



Experimental

Crystal data

$[Cu_2(C_4H_2O_4)_2(H_2O)_2]\cdot 3H_2O$
 $M_r = 445.27$
Monoclinic, $P2_1/c$
 $a = 8.8835$ (14) Å

$b = 8.7700$ (14) Å
 $c = 18.814$ (3) Å
 $\beta = 97.994$ (3) $^\circ$
 $V = 1451.5$ (4) Å³

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 3.00$ mm⁻¹

$T = 173$ (2) K
 $0.30 \times 0.28 \times 0.05$ mm

Data collection

Bruker APEXII diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{min} = 0.470$, $T_{max} = 0.860$

9585 measured reflections
2643 independent reflections
2331 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.026$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.023$
 $wR(F^2) = 0.055$
 $S = 1.03$
2643 reflections
238 parameters
15 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.31$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.31$ e Å⁻³

Table 1
Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1W-H1WA..O5	0.878 (16)	2.034 (17)	2.910 (3)	175 (3)
O1W-H1WB..O2W	0.864 (16)	2.034 (18)	2.863 (3)	161 (3)
O2W-H2WA..O7	0.861 (16)	1.967 (17)	2.827 (3)	177 (3)
O2W-H2WB..O3W	0.851 (16)	2.014 (18)	2.854 (3)	169 (3)
O3W-H3WA..O2 ⁱ	0.871 (16)	1.995 (19)	2.847 (2)	166 (3)
O3W-H3WB..O1W ⁱ	0.857 (16)	2.17 (2)	2.928 (3)	148 (2)
O9-H9A..O1W ⁱⁱ	0.853 (16)	1.987 (18)	2.831 (3)	170 (3)
O9-H9B..O10 ⁱⁱⁱ	0.851 (16)	2.023 (19)	2.855 (3)	165 (2)
O10-H10A..O2W ^{iv}	0.867 (16)	1.943 (17)	2.797 (3)	168 (3)
O10-H10B..O3W	0.846 (16)	2.059 (18)	2.879 (3)	163 (2)

Symmetry codes: (i) $-x + 1, -y + 1, -z$; (ii) $-x + 2, -y + 1, -z$; (iii) $x + 1, y + 1, z$; (iv) $-x + 1, -y, -z$.

Data collection: *COSMO* (Bruker, 2006); cell refinement: *APEX2* (Bruker, 2006); data reduction: *SAINT* (Bruker, 2006); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *CrystalMaker* (Palmer, 2007); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SJ2520).

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