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Key indicators

Single-crystal X-ray study

 $T = 295$ KMean $\sigma(C-C) = 0.004$ Å R factor = 0.034 wR factor = 0.080

Data-to-parameter ratio = 14.2

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.A Cu–Mn complex of a Schiff base containing alanylglycine: hexaaquamanganese(II) bis{[*N*-(2-oxidobenzylidene)alanylglycinato]cuprate(II)} dodecahydrate

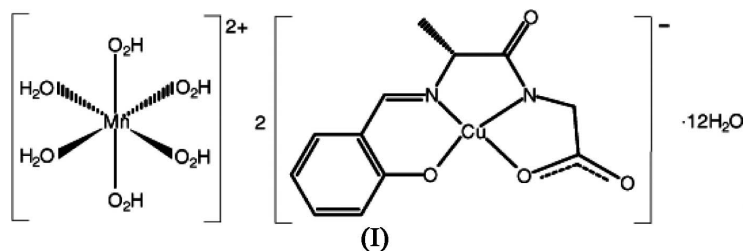
In the title compound, $[Mn(H_2O)_6][Cu(C_{12}H_{11}N_2O_4)]_2 \cdot 12H_2O$, the Cu atom has a square-planar coordination with two N and two O atoms of the tetradentate Schiff base ligand. The centrosymmetric cation $[Mn(H_2O)_6]^{2+}$ and anion $[CuL]^-$ (L is a Schiff base derived from alanylglycine and salicylaldehyde) occupy a stacking structure within well separated columns along the c axis.

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Comment

The title compound, (I), is one of the copper(II) Schiff base complexes derived from alanylglycine and salicylaldehyde. The study of copper(II) Schiff base complexes formed between pyridoxal or analogs and amino acids or small peptides (Koh *et al.*, 1996) is a subject of considerable interest owing to their structural and magnetic properties, and potential antimicrobial, anti-inflammatory and antipyretic activities, together with a superoxide dismutase-like activity (Garcia-Raso *et al.*, 1999).



The complex crystallizes in the monoclinic space group $C2/c$. The asymmetric unit consists of one $[CuL]^-$ anion (L is a Schiff base derived from alanylglycine and salicylaldehyde), one half-cation (Mn1, O1W, O2W, O3W), five uncoordinated water molecules in general positions and two uncoordinated water molecules (O4W and O9W) in special positions. $[CuL]^-$ has an approximately square-planar structure. The deprotonated Schiff base ligand is a triple negatively charged quadridentate ONNO chelant, coordinating to the Cu^{II} ion *via* one phenol O atom (O1), one deprotonated amide N atom (N2), one imine N atom (N1) and one carboxylate O atom (O3). The O1–Cu1–N2 angle of $179.15(8)^\circ$ is nearly linear. The benzene ring (C1–C6) and the O1/C1/C6/C7/N1/Cu1 ring are almost coplanar, with a small dihedral angle of $1.13(10)^\circ$. The Mn^{II} ion lies on an inversion center and exhibits essentially octahedral coordination by six aqua ligands.

The anions and cations form well separated columns along the c axis in the stacking structure of (I). Intermolecular hydrogen bonds play an important role in the stabilization of the whole structure (Table 2).

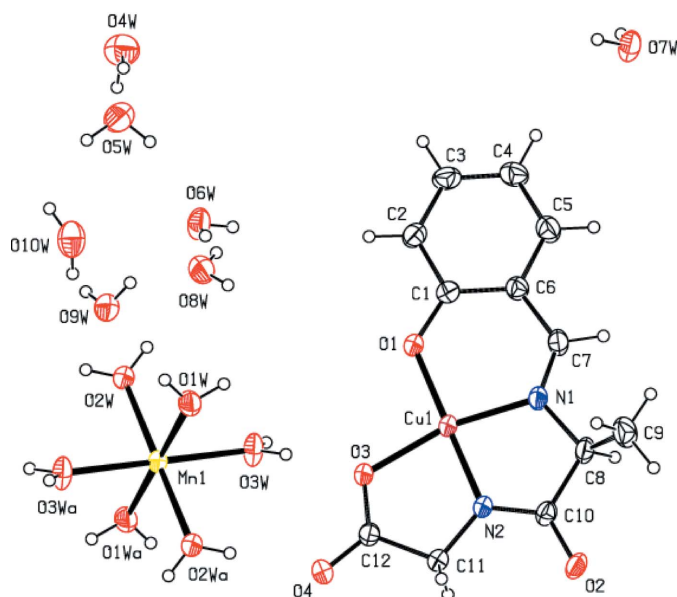


Figure 1
View of the structure of (I), showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 30% probability level.

Experimental

The title compound, (I), was prepared from the copper(II) Schiff base complex [CuL] with manganese perchlorate, as described previously (Liu *et al.*, 2004). Suitable crystals of (I) were obtained by evaporation of an aqueous solution. Analysis calculated for $C_{24}H_{58}Cu_2MnN_4O_{26}$: C 28.80, H 5.84, N 5.60%; found: C 28.74, H 5.90, N 5.54%.

Crystal data

[Mn(H₂O)₆][Cu(C₁₂H₁₁N₂O₄)₂·12H₂O]
M_r = 1000.78
 Monoclinic, *C*2/*c*
a = 28.479 (6) Å
b = 11.781 (2) Å
c = 14.483 (3) Å
 β = 117.280 (10)°
V = 4318.8 (15) Å³
Z = 4
D_x = 1.539 Mg m⁻³
 Mo Kα radiation
 μ = 1.35 mm⁻¹
T = 295 (2) K
 Block, red-violet
 0.50 × 0.35 × 0.20 mm

Data collection

Bruker SMART APEX CCD diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 2000)
T_{min} = 0.551, *T_{max}* = 0.774
 12220 measured reflections
 4719 independent reflections
 3482 reflections with *I* > 2σ(*I*)
R_{int} = 0.051
θ_{max} = 27.1°

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.034
wR(*F*²) = 0.080
S = 0.93
 4719 reflections
 333 parameters
 H atoms treated by a mixture of independent and constrained refinement
w = 1/[σ²(*F_o*²) + (0.0374*P*)²]
 where *P* = (*F_o*² + 2*F_c*²)/3
 (Δ/σ)_{max} < 0.001
 Δρ_{max} = 0.39 e Å⁻³
 Δρ_{min} = -0.34 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Cu1—O1	1.8677 (16)	O2—C10	1.258 (3)
Cu1—O3	1.9869 (18)	O3—C12	1.275 (3)
Cu1—N1	1.925 (2)	O4—C12	1.235 (3)
Cu1—N2	1.8698 (19)	N1—C8	1.478 (3)
Mn1—O3W	2.141 (3)	N1—C7	1.267 (3)
Mn1—O1W	2.190 (2)	N2—C11	1.434 (3)
Mn1—O2W	2.1721 (18)	N2—C10	1.304 (4)
O1—C1	1.321 (3)		
O1—Cu1—O3	97.25 (7)	Cu1—N1—C7	124.95 (18)
O1—Cu1—N1	95.87 (8)	Cu1—N1—C8	114.08 (15)
O1—Cu1—N2	179.15 (8)	C7—N1—C8	120.8 (2)
O3—Cu1—N1	166.79 (7)	C10—N2—C11	123.33 (19)
O3—Cu1—N2	83.03 (8)	Cu1—N2—C10	119.20 (15)
N1—Cu1—N2	83.83 (9)	Cu1—N2—C11	117.18 (17)
O2W ⁱ —Mn1—O3W	90.33 (8)	O1—C1—C2	118.6 (2)
O3W—Mn1—O3W ⁱ	180	O1—C1—C6	124.4 (2)
O2W—Mn1—O3W	89.67 (8)	N1—C7—C6	125.4 (3)
O1W ⁱ —Mn1—O2W	85.14 (7)	N1—C8—C9	112.4 (2)
O1W—Mn1—O2W	94.86 (7)	N1—C8—C10	108.0 (2)
O1W—Mn1—O3W	88.91 (9)	O2—C10—N2	125.7 (2)
O1W—Mn1—O1W ⁱ	180	O2—C10—C8	119.7 (2)
O2W—Mn1—O3W ⁱ	90.33 (8)	N2—C10—C8	114.6 (2)
O1W ⁱ —Mn1—O3W ⁱ	91.09 (9)	N2—C11—C12	107.98 (19)
O2W—Mn1—O2W ⁱ	180	O3—C12—O4	124.0 (2)
Cu1—O1—C1	125.71 (17)	O3—C12—C11	117.4 (2)
Cu1—O3—C12	114.25 (17)	O4—C12—C11	118.7 (2)

Symmetry code: (i) -*x*, -*y*, -*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>
O1W—H1A...O9W	0.75 (3)	2.03 (3)	2.767 (3)	173 (3)
O1W—H1A...O9W	0.75 (3)	2.03 (3)	2.767 (3)	173 (3)
O1W—H1B...O7W ⁱ	0.83 (4)	1.91 (3)	2.724 (3)	168 (3)
O2W—H2A...O8W	0.79 (2)	1.98 (2)	2.759 (3)	172 (3)
O2W—H2B...O4 ⁱ	0.79 (2)	1.93 (2)	2.722 (3)	179 (4)
O3W—H3A...O10W ⁱⁱⁱ	0.76 (3)	1.96 (3)	2.724 (4)	179 (4)
O3W—H3B...O3	0.81 (3)	1.91 (3)	2.714 (3)	174 (3)
O4W—H4W...O5W	0.98 (4)	1.85 (4)	2.808 (4)	168 (4)
O5W—H5A...O8W ^v	0.86 (5)	1.97 (5)	2.797 (4)	164 (5)
O5W—H5B...O6W	0.80 (5)	1.95 (5)	2.718 (4)	159 (5)
O6W—H6A...O2 ^v	0.75 (4)	1.90 (4)	2.653 (3)	177 (1)
O6W—H6B...O4 ^{vi}	0.79 (3)	2.00 (3)	2.789 (3)	179 (5)
O7W—H7A...O2 ^{vii}	0.79 (3)	1.90 (3)	2.689 (3)	174 (3)
O7W—H7B...O1 ^v	0.76 (4)	2.01 (4)	2.765 (3)	174 (3)
O8W—H8A...O5W ^{viii}	0.78 (4)	1.97 (4)	2.734 (4)	167 (4)
O8W—H8B...O7W ⁱⁱ	0.76 (3)	1.99 (3)	2.749 (4)	174 (3)
O9W—H9W...O6W	0.78 (3)	2.03 (3)	2.803 (3)	174 (3)

Symmetry codes: (i) -*x*, -*y*, -*z*; (ii) -*x* + ½, *y* - ½, -*z* + ½; (iii) *x*, *y*, *z* - 1; (iv) -*x*, *y*, -*z* + ½; (v) -*x* + ½, *y* + ½, -*z* + ½; (vi) *x*, -*y*, *z* + ½; (vii) *x*, *y* + 1, *z*; (viii) *x*, -*y* + 1, *z* - ½.

The water H atoms were located in difference maps and refined freely [O—H = 0.75 (3)–0.98 (4) Å]. The other H atoms were placed in geometrically idealized positions with C—H distances in the range 0.93–0.98 Å and constrained to ride on their parent atoms with *U*_{iso}(H) = 1.2–1.5*U*_{eq}(C).

Data collection: SMART for WNT/2000 (Bruker, 2002); cell refinement: SAINT-Plus (Bruker, 2003); data reduction: SAINT-Plus; program(s) used to solve structure: SIR97 (Altomare *et al.*, 1999); program(s) used to refine structure: SHELXTL (Bruker, 2000); molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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