Guan, J., Shen, Q., Jin, S. & Lin, Y. (1994). Acta Cryst. C50, 891-893.

- Herrmann, W. A., Anwander, R., Kleine, M., Ofele, K., Riede, J. & Scherer, W. (1992). Chem. Ber. 125, 2391–2397.
- Kilimann, U., Schäfer, M., Herbst-Irmer, R. & Edelmann, F. T. (1994). J. Organomet. Chem. 469, C10–C14.
- Sheldrick, G. M. (1990a). Acta Cryst. A46, 467-473.
- Sheldrick, G. M. (1990b). SHELXTL/PC Users Manual. Siemens
- Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.Sheldrick, G. M. (1993). SHELXL93. Program for the Refinement of Crystal Structures. University of Göttingen, Germany.
- Shen, Q., Cheng, Y. & Lin, Y. (1991). J. Organomet. Chem. 419, 293–298.
- Siemens (1994). XSCANS. X-ray Single Crystal Analysis System. Version 2.1. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.
- Song, S., Shen, Q. & Jin, S. (1992). Polyhedron, 11, 2863-2865.
- Walker, N. & Stuart, D. (1983). Acta Cryst. A39, 158-166.

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Diaquabis(2-quinolinecarboxylato-*N*,*O*)manganese(II)

NOBUO OKABE AND MAKIKO KOIZUMI

Faculty of Pharmaceutical Sciences, Kinki University, Kowakae 3-4-1, Higashiosaka, Osaka 577, Japan

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Abstract

The title compound, $[Mn(C_{10}H_6NO_2)_2(H_2O)_2]$, contains a six-coordinate Mn atom at a center of symmetry. The Mn atom displays distorted octahedral coordination. 2-Quinolinecarboxylate anions lie in a *trans* position with respect to one another, forming the equatorial plane, and the water ligands occupy the axial positions. In the equatorial plane, the Mn–ligand distances and the bite angle of the ligand are Mn–O 2.131 (2), Mn–N 2.315 (2) Å and O–Mn–N 74.85 (7)°. In the axial direction, Mn–OH₂ 2.185 (2) and O–Mn–N 91.10 (8)°. Molecules are linked by a three-dimensional O–H···O hydrogen-bonding network involving water molecules and carboxylate groups.

Comment

2-Quinolinecarboxylic acid (quinaldinic acid), (I), is one of the final products of tryptophan metabolism and is derived from kynurenic acid (Martin, Mayes & Rodwell, 1983). It inhibits dopamine β -hydroxylase (EC 1.14.17.1), which is a multi-subunit enzyme containing two Cu atoms per subunit (Townes, Titone & Rosenberg, 1990).

2-Quinolinecarboxylic acid is also known to be a potent chelator of copper and other transition metal ions (Martell & Smith, 1974). Accordingly, we have carried out the structure analysis of the chelate compounds of 2-quinolinecarboxylic acid and biologically important metal ions in order to clarify the binding scheme of this metabolite to metal ions. In the present study, the crystal structure of the Mn^{II} complex of 2-quinolinecarboxylic acid, (II), has been determined.



The structure of the precursor of 2-quinolinecarboxylic acid, kynurenic acid, has been reported (Okabe, Miura & Shimosaki, 1996), and recently the structure of (II) has been reported as the water and ethanol solvate (Haendler, 1996). The molecular structure of (II) with the atom-labeling scheme is shown in Fig. 1. The Mn^{II} atom is coordinated octahedrally by two trans quinoline N atoms [Mn1-N1 2.315 (2) Å], by two trans-O atoms of the carboxylate groups in position 2 of (I) [Mn1-O2 2.131 (2) Å] and by two trans-O atoms of the water molecules [Mn1-O3 2.185 (2) Å]. These Mn^{II}-ligand distances are similar to those values in the water/ethanol solvate [Mn-N 2.324(3), Mn—O 2.125 (2) and Mn—OH₂ 2.209 (3) Å; Haendler, 1996]. The carboxyl group is present in an ionized form and is almost parallel to the quinoline ring plane $[O2-C1-C2-N1 - 15.4(3)^{\circ}]$. These conformational features are also observed in the solvated complex (Haendler, 1996). The molecular packing of the crystal



Fig. 1. ORTEPII (Johnson, 1976) drawing of the title compound with the atom-numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 50% probability level.

Acta Crystallographica Section C ISSN 0108-2701 © 1997 structure is shown in Fig. 2. The crystal structure is stabilized by stacking interaction between quinoline rings and a three-dimensional hydrogen-bonding network between water molecules and carboxylate groups of the neighboring complex: O3-H31...O1ⁱ 2.753 (3) Å and O3—H32···O1ⁱⁱ 2.845 (3) Å [symmetry codes: (i) x, y, -1 + z; (ii) x, $\frac{1}{2} - y$, $-\frac{1}{2} + z$].



Fig. 2. Packing diagram of the title compound as viewed down the caxis.

Experimental

The light-yellow plate crystal was obtained by the slow evaporation of an aqueous solution of a mixture of 2-quinolinecarboxylic acid and [Mn(CH₃CO₂)₂].4H₂O in the molar ratio of 4:1 at room temperature.

Crystal data

$[Mn(C_{10}H_6NO_2)_2(H_2O)_2]$	Mo $K\alpha$ radiation
$M_r = 435.30$	$\lambda = 0.7107 \text{ Å}$
Monoclinic	Cell parameters from 24
$P2_{1}/c$	reflections
a = 10.028 (6) Å	$\theta = 18.25 - 23.40^{\circ}$
b = 13.118(7) Å	$\mu = 0.729 \text{ mm}^{-1}$
c = 7.321(5) Å	T = 296 K
$\beta = 107.33 (4)^{\circ}$	Plate
V = 919.3 (9) Å ³	$0.40 \times 0.40 \times 0.15$ mm
Z = 2	Light yellow
$D_x = 1.572 \text{ Mg m}^{-3}$	
D_m not measured	
Data collection	
Rigaku AFC-6S diffractom-	1666 reflections with
eter	$I > 3\sigma(I)$
ω –2 θ scans	$R_{\rm int} = 0.019$

intensity decay: none

eter	$I > 3\sigma(I)$
ω –2 θ scans	$R_{\rm int} = 0.019$
Absorption correction:	$\theta_{\rm max} = 27.5^{\circ}$
ψ scans (North, Phillips	$h = 0 \rightarrow 12$
& Mathews, 1968)	$k = 0 \rightarrow 16$
$T_{\rm min} = 0.582, T_{\rm max} = 0.896$	$l = -9 \rightarrow 9$
2333 measured reflections	4 standard reflections
2215 independent reflections	every 150 reflections

Refinement

Refinement on F	$(\Lambda/\sigma) = 0.002$
P = 0.025	$\Delta_{10} = 0.46 \text{ s}^{-3}$
K = 0.055	$\Delta \rho_{\rm max} = 0.40 \ {\rm e} \ {\rm A}$
wR = 0.043	$\Delta \rho_{\rm min} = -0.56 \ {\rm e} \ {\rm A}^{-3}$
S = 1.670	Extinction correction: none
1666 reflections	Scattering factors from Inter-
165 parameters	national Tables for X-ray
H atoms refined isotropically	Crystallography (Vol. IV)
$w = 4F_o^2/\sigma^2(F_o^2)$	

Table	1.	Fraci	tional	atomic	<i>CO0</i>	rdinates	and	equival	ent
		isotro	pic di	splacem	ent p	paramete	ers (Å	²)	

$B_{\text{eq}} = (8\pi^2/3)\sum_i \sum_j U^{ij} a_i^* a_i^* \mathbf{a}_i . \mathbf{a}_j.$

	x	у	z	Ben
Mnl	1/2	0	1/2	2.31 (2)
01	0.5718(2)	0.1537(1)	1.0170(2)	3.06 (6)
02	0.5952 (2)	0.0525(1)	0.7845(2)	3.14 (6)
03	0.5607 (2)	0.1411 (2)	0.3876 (3)	4.2 (1)
N1	0.3241 (2)	0.0838(1)	0.5851 (2)	2.11 (6)
C1	0.5241 (2)	0.1045 (2)	0.8652 (3)	2.32 (8)
C2	0.3680(2)	0.1120(2)	0.7674(3)	2.26 (8)
C3	0.2794 (3)	0.1503 (2)	0.8672 (3)	2.85 (9)
C4	0.1403 (3)	0.1597 (2)	0.7737(3)	2.88 (9)
C5	-0.0529(3)	0.1461 (2)	0.4710(4)	3.1 (1)
C6	-0.0959(3)	0.1244 (2)	0.2809 (4)	3.4(1)
C7	-0.0002(3)	0.0890(2)	0.1905 (3)	3.2 (1)
C8	0.1371 (3)	0.0736(2)	0.2892 (3)	2.70 (8)
C9	0.1849(2)	0.0967 (2)	0.4867 (3)	2.24 (7)
C10	0.0893 (2)	0.1341 (2)	0.5788(3)	2.54 (8)

Table 2. Selected geometric parameters (Å, °)

Mn1O2	2.131 (2)	C3C4	1.364 (4)
Mn1—O3	2.185 (2)	C4C10	1.406 (3)
Mn1—N1	2.315(2)	C5—C6	1.359 (4)
01—C1	1.250(3)	C5-C10	1.418 (4)
02C1	1.254 (3)	C6—C7	1.397 (4)
N1-C2	1.327 (3)	C7—C8	1.366 (3)
N1	1.378 (3)	C8C9	1.414 (3)
C1C2	1.518(3)	C9-C10	1.414 (3)
C2C3	1.401 (3)		
O2-Mn1-O3	90.01 (8)	C2—C3—C4	118.9 (2)
O2-Mn1-O3 ⁱ	89.99 (8)	C3-C4-C10	119.8 (2)
O2-Mn1-N1	74.85 (8)	C6-C5-C10	120.4 (2)
O3—Mn1—N1	91.10 (8)	C5C6C7	120.1 (2)
O2—Mn1—N1 ⁱ	105.15 (8)	C6—C7—C8	121.7 (2)
Mn1-02-C1	119.0(1)	C7C8C9	119.4 (2)
C2-N1-C9	118.2 (2)	N1-C9-C8	119.4 (2)
01C1O2	125.3 (2)	N1-C9-C10	121.2 (2)
01—C1—C2	117.0(2)	C8-C9-C10	119.4 (2)
02C1C2	117.7 (2)	C4-C10-C5	122.7 (2)
N1-C2-C1	116.2 (2)	C4C10C9	118.3 (2)
N1-C2-C3	123.6(2)	C5-C10-C9	119.0 (2)
C1C2C3	120.1 (2)		

Symmetry code: (i) 1 - x, -y, 1 - z.

All H atoms were located from difference Fourier maps and included in the refinement calculations isotropically.

Data collection: MSC/AFC Diffractometer Control Software (Molecular Structure Corporation, 1988). Cell refinement: MSC/AFC Diffractometer Control Software. Data reduction: TEXSAN (Molecular Structure Corporation, 1985). Program(s) used to solve structure: SHELXS86 (Sheldrick, 1985) and DIRDIF (Beurskens, 1984). Program(s) used to refine structure: TEXSAN. Molecular graphics: ORTEPII (Johnson, 1976).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1028). Services for accessing these data are described at the back of the journal.

References

Beurskens, P. T. (1984). DIRDIF. Direct Methods for Difference Structures – an Automatic Procedure for Phase Extension and Refinement of Difference Structure Factors. Technical Report 1984/1. Crystallography Laboratory, Toernooiveld, 6525 ED Nijmegen, The Netherlands.

Haendler, H. M. (1996). Acta Cryst. C52, 801-803.

- Johnson, C. K. (1976). *ORTEPII*. Report ORNL-5138. Oak Ridge National Laboratory, Tennessee, USA.
- Martell, A. E. & Smith, R. M. (1974). *Critical Stability Constants*, Vol. 1, pp. 78, 372; Vol. 2, p. 219. New York: Plenum Press.
- Martin, D. W., Mayes, P. A. & Rodwell, V. W. (1983). Harper's Review of Biochemistry, pp. 464-479. Tokyo: Maruzen Asia.
- Molecular Structure Corporation (1985). TEXSAN. TEXRAY Structure Analysis Package. MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.
- Molecular Structure Corporation (1988). MSC/AFC Diffractometer Control Software. MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.
- North, A. C. T., Phillips, D. C. & Mathews, F. S. (1968). Acta Cryst. A24, 351-359.
- Okabe, N., Miura, J. & Shimosaki, A. (1996). Acta Cryst. C52, 1827– 1829
- Sheldrick, G. M. (1985). SHELXS86. Crystallographic Computing 3, edited by G. M. Sheldrick, C. Krüger & R. Goddard, pp. 175–189. Oxford University Press.
- Townes, S., Titone, C. & Rosenberg, R. C. (1990). *Biochim. Biophys.* Acta, 1037, 240–247.

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Tetrakis(μ -benzoato-O:O')bis(urea)-1 κO ,2 κO -dicopper(II)

Ivan Leban,^a Nikolaj Grgurevič,^a Joachim Sieler^b and Primož Šegedin^a

^a Faculty of Chemistry and Chemical Technology, University of Ljubljana, Aškerčeva 5, PO Box 537, SLO-1001 Ljubljana, Slovenia, and ^bInstitut für Anorganische Chemie, Universität Leipzig, Linnéstrasse 3, D-04103 Leipzig, Germany. E-mail: ivan.leban@uni-lj.si

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Abstract

The adduct of tetra- μ -benzoato-dicopper(II) with urea, $[Cu(C_7H_5O_2)_2(CH_4N_2O)]_2$, is dinuclear. Each Cu atom is displaced by 0.195 (3) Å from the plane of the four benzoato O atoms to which it is bonded. The Cu- \cdot Cu distance is 2.633 (2) Å. The average Cu-O(benzoate) distance is 1.964 (3) Å and the Cu-O(urea) bond length is 2.156 (6) Å.

Comment

Carboxylic acid complexes of copper(II) are of special chemical and biological interest and accordingly have been extensively studied. Their preparation, structural, magnetic and spectral properties have frequently been described (see Melnik, 1982; Kato & Muto, 1988; Valko, Bilton, Morris & Melnik, 1993; Sundberg, Uggla & Melnik, 1996, and references therein). Our interest in copper(II) carboxylates evolves from their biological activity (Bergant, Petrič, Pohleven, Reberšek & Šegedin, 1994; Grgurevič, Petrič, Pohleven & Šegedin, 1996). Different pharmacological effects have been established for solvated mononuclear and dinuclear complexes (Sorensen, 1983) and fungicidal and insecticidal action is to be anticipated.

Several dinuclear tetrabenzoatocopper(II) complexes of general formula $Cu_2(benz)_4.2L$ (benz = $C_6H_5COO^-$; L = axially bonded neutral ligand) have been structurally characterized. The axial ligands L were: methanol, CH₃OH (DAYJIW); dimethylsulfoxide, (CH₃)₂SO (CO-CYUO); pyridine, C_5H_5N (BZPRCV, BZPRCV01); quinoline, C₉H₇N (TBZOQC); triphenylphosphine oxide, (C₆H₅)₃PO (PILBIV); benzoic acid, C₆H₅COOH (KUB-HOE); β -picoline, C₆H₇N (KUBHUK); 4-methylquinoline, $C_{10}H_9N$ (KUBJAS); 7-methylquinoline, $C_{10}H_9N$ (KUBJIA); 4,7-dichloroquinoline, C₉H₅Cl₂N (KUB-JOG); caffeine, C₈H₁₀N₄O₂ (KUBJUM) (from the Cambridge Structural Database, October 1995 release, with refcodes in parentheses; Allen & Kennard, 1993). Additionally, the dimeric copper(II) benzoate with N,N-diethylnicotinamide has been described recently (Hökelek, Necefoğlu & Balci, 1995).

In this paper, we report the structure determination of a dinuclear copper(II) benzoate complex with two urea molecules [urea = $CO(NH_2)_2$, CH_4N_2O] acting as axial ligands (Fig. 1), $Cu_2(benz)_4.2(urea)$, (I).



The structure analysis confirmed the expected dimeric form. Four bidentate benzoate anions form bridges between Cu atoms which have a 4+1 environment. The coordination sphere around each Cu atom is made up of four O atoms of benzoate groups in the basal plane, the apical position being occupied by the urea O atom. The average Cu—O(basal) distance is 1.964 (3) Å, which is a typical value in dimeric copper carboxylates complexes. The Cu atom is shifted by 0.195 (3) Å out of the plane of the four O atoms towards the axial ligand.

As suggested by Kawata *et al.* (1992), φ_{rot} (the dihedral angle between the phenyl and carboxyl planes