

Catharine Esterhuysen,^{a*}
Martin W. Bredenkamp,^a
Leon van Kralingen,^a Hanelie J.
Adendorff,^b Edmund P. Jacobs^a
and Pieter Swart^c

^aDepartment of Chemistry and Polymer Science, University of Stellenbosch, Private Bag X1, Matieland 7602, South Africa, ^bCentre for Teaching and Learning, University of Stellenbosch, Private Bag X1, Matieland 7602, South Africa, and ^cDepartment of Biochemistry, University of Stellenbosch, Private Bag X1, Matieland 7602, South Africa

Correspondence e-mail: ce@sun.ac.za

Key indicators

Single-crystal X-ray study
T = 173 K
Mean $\sigma(\text{C}-\text{C}) = 0.005 \text{ \AA}$
R factor = 0.033
wR factor = 0.072
Data-to-parameter ratio = 12.4

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

Aqua[3,6-bis(methoxycarbonylmethyl)-3,6-diazaoctanedioato]copper(II) dihydrate

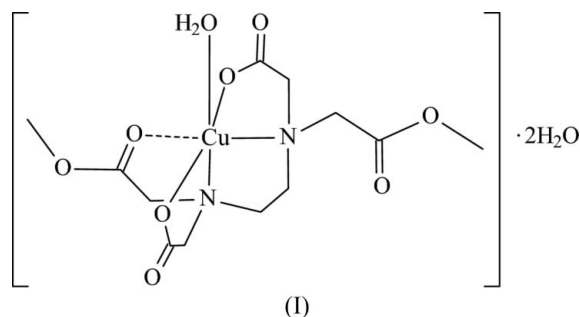
The title complex, $[\text{Cu}(\text{C}_{12}\text{H}_{18}\text{N}_2\text{O}_8)(\text{H}_2\text{O})] \cdot 2\text{H}_2\text{O}$, crystallizes with the tetradentate ethylenediaminetetraacetic acid (edta) diester derivative and one water molecule forming a pyramidal environment. The vacant site of the octahedral complex is remotely associated with an O atom of one of the carbonyl atoms of the edta derivative.

Received 13 December 2005

Accepted 11 January 2006

Comment

In our quest for alternative immobilized metal affinity chromatography (IMAC) ligands (Porath *et al.*, 1975) that may be tethered to a solid support, ethylenediaminetetraacetic acid (edta) esters have been investigated. Esterifying two of the four acid groups of the potentially hexadentate edta ligand reduces it to a tetradentate ligand, providing two open sites in the octahedral coordination for selective coordination to the target compound (usually enzymes and proteins) to be purified by IMAC, and simultaneously using the ester group to anchor the ligand to the tether. The dimethyl ester of edta has been used as a model compound in order to investigate the coordination arrangement of the ligand, since the availability of adjacent sites is advantageous in IMAC.



The dimethyl ester of edta was prepared by refluxing edta dianhydride in an excess of methanol. The resultant ligand was complexed to basic copper(II) carbonate in water at 333 K, and allowed to crystallize, yielding the title complex, (I), with two molar equivalents of water in the asymmetric unit. The tetradentate ligand has the desired configuration (Fig. 1), coordinating through the two free carboxylates and both amine groups, with the two remaining sites of the Cu coordination adjacent. A coordinated water molecule and the carbonyl O atom of one of the ester groups complete the severely Jahn–Teller distorted octahedral configuration around the Cu atom; see Table 1. The axial Cu–O distance is nearly 0.6 Å longer than the two equatorial Cu–O(carboxylate) distances, while the axial Cu–N bond is 0.25 Å longer than its equatorial counterpart. Furthermore,

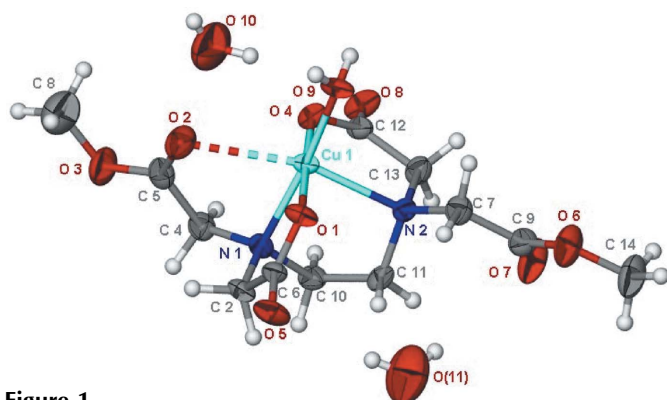


Figure 1
The asymmetric unit of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

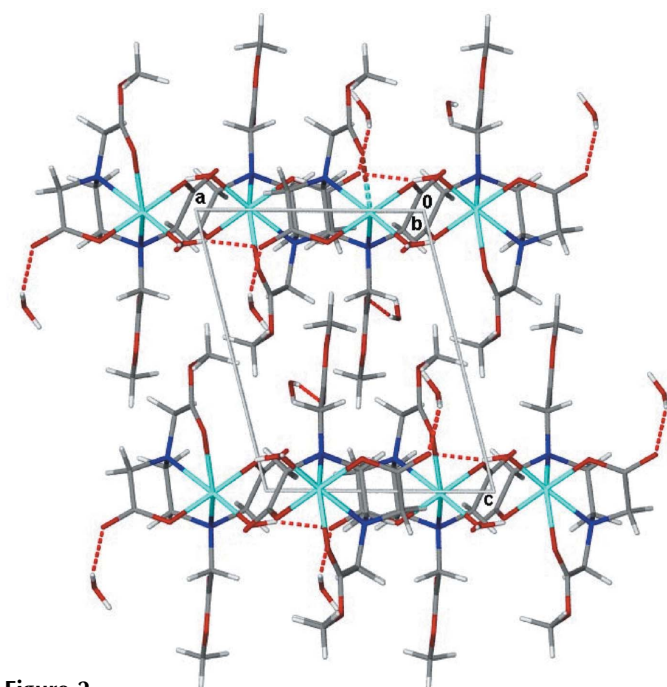


Figure 2
The packing of (I), showing the hydrogen-bonding (dashed lines) pattern of layers parallel to the *ab* plane.

the two axial ligands are bent away from the water ligand, such that the O2—Cu1—N2 angle is 154.03 (9)°.

The two five-membered N—O(carboxylate) chelate rings that coordinate through equatorial O atoms are both essentially planar, the largest deviations from planarity being 0.097 (2) Å for C13 and 0.098 (2) Å for C2. The five-membered N—O(ester carbonyl) chelate ring coordinating through the axial O is, on the other hand, twisted, with the largest deviation from planarity exhibited by N1 [0.305 (2) Å]. A further indication of the twisting can be seen in the Cu1—N1—C4—C5 torsion angle of −39.8 (3)°.

Each H atom of the coordinated water molecule participates in hydrogen bonding (see Table 2) with one of the coordinated carboxylate groups of two neighbouring complexes. This leads to the formation of layers of the complexes parallel to the *ab* plane (see Fig. 2). The terminal

ester groups of the edta derivative and the free water molecules lie between these layers, with the two water molecules hydrogen bonded to the carbonyl O atoms of a bound carboxylate and a free ester group, respectively.

Experimental

Preparation of dimethyl *N,N'*-bis(carboxymethyl)-3,6-diazaoctanedioate (DMDDO): ethylenediaminetetraacetic dianhydride (5.00 g, 20 mmol) was mixed with methanol (100 ml) and refluxed for 6 h. The solution was filtered while hot and the methanol removed under reduced pressure. The residue was dissolved in refluxing ethyl acetate (50 ml) and allowed to crystallize below 273 K. The crystals were filtered off and washed with diethyl ether and dried under high vacuum, yielding dimethyl *N,N'*-bis(carboxymethyl)-3,6-diazaoctanedioate (yield 6.27 g, 98%). ν_{\max} (cm^{−1}, KBr, Perkin–Elmer 1720 F T–IR spectrophotometer): 3474 (*w*), 3024 (*m*), 2996 (*w*), 2967 (*m*), 1750 (*vs*, ester carbonyl), 1632 (*s*, carboxylate carbonyl), 1471 (*sh*), 1444 (*s*), 1402 (*s*), 1241 (*vs*), 1216 (*s*) and 1087 (*m*); ¹H NMR (CDCl₃, Varian VXR 300, 300 MHz): δ 2.73 (*s*, 4H, −NCH₂CH₂N−), 3.43 (*s*, 4H, −CH₂-acid), 3.55 (*s*, 4H, −CH₂-ester) and 3.60 (*s*, 6H, −CH₃); ¹³C NMR (CDCl₃, Varian VXR 300, 75 MHz): δ 50.89 (*q*, −CH₃) 51.2 (*t*, −NCH₂CH₂N−), 54.1 (*t*, −CH₂-ester), 54.4 (*t*, −CH₂-acid), 171.2 (*s*, −COO-ester) and 172.2 (*s*, −COOH); *elz* (EI): 320 (100%, *M*⁺).

Preparation of (I): DMDDO (1.00 g, 3.1 mmol) was dissolved in distilled water (5 ml) at 333 K. CuCO₃·Cu(OH)₂ (238 mg, 1.6 mmol) was added with stirring until dissolved and the evolution of carbon dioxide ceased. Deep-blue crystals formed at ambient temperature. The crystals were filtered off and washed with ice-cold ethanol, yielding 1.05 g (89%) of (I). ν_{\max} (cm^{−1}, KBr): 3443 (*s, br*), 2956 (*m*), 1739 (*vs*, ester carbonyl), 1610 (*vs*, carboxylate carbonyl), 1436 (*m*), 1381–2 (*s*), 1255 (*sh*), 1221–2 (*s*), 1117–8 (*s*); *elz* (EI): 382 (*M*⁺).

Crystal data

[Cu(C₁₂H₁₈N₂O₈)(H₂O)]·2H₂O
 $M_r = 435.87$
 Monoclinic, *P*2₁
 $a = 7.5208$ (15) Å
 $b = 13.158$ (3) Å
 $c = 9.5684$ (19) Å
 $\beta = 104.31$ (3)°
 $V = 917.5$ (4) Å³
 $Z = 2$

$D_x = 1.578$ Mg m^{−3}
 Mo $K\alpha$ radiation
 Cell parameters from 1895 reflections
 $\theta = 1.0$ –27.5°
 $\mu = 1.25$ mm^{−1}
 $T = 173$ (2) K
 Prism, deep-blue
 0.37 × 0.25 × 0.12 mm

Data collection

Nonius KappaCCD diffractometer
 φ and ω scans
 Absorption correction: multi-scan
 (SCALEPACK; Otwinowski & Minor, 1997)
 $T_{\min} = 0.70$, $T_{\max} = 0.86$
 4595 measured reflections

3207 independent reflections
 2958 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$
 $\theta_{\max} = 26.0^\circ$
 $h = -9 \rightarrow 8$
 $k = -16 \rightarrow 14$
 $l = -7 \rightarrow 11$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.072$
 $S = 1.04$
 3207 reflections
 259 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0345P)^2 + 0.0434P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.40$ e Å^{−3}
 $\Delta\rho_{\min} = -0.29$ e Å^{−3}
 Absolute structure: Flack (1983),
 1318 Friedel pairs
 Flack parameter: 0.019 (14)

Table 1

Selected geometric parameters (Å, °).

Cu1—O1	1.918 (2)	Cu1—O9	1.956 (2)
Cu1—O2	2.545 (3)	Cu1—N1	2.080 (3)
Cu1—O4	1.951 (2)	Cu1—N2	2.326 (3)
O1—Cu1—O4	176.58 (11)	O9—Cu1—N1	170.86 (12)
O2—Cu1—O9	101.32 (10)	O9—Cu1—N2	102.95 (10)
O2—Cu1—N2	154.03 (9)		

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>
O10—H102...O7 ⁱ	0.81 (4)	2.23 (3)	3.024 (5)	167 (6)
O9—H92...O8 ⁱⁱ	0.844 (18)	1.851 (19)	2.695 (4)	178 (4)
O9—H91...O5 ⁱⁱ	0.827 (17)	1.88 (2)	2.638 (3)	153 (3)
O11—H112...O5	0.82 (4)	2.21 (3)	2.981 (5)	157 (7)

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

The H atoms in the complexed and uncoordinated water molecules were located in a difference Fourier map, and restrained with O—H and H...H distances of 0.82 (2) and 1.39 (2) Å, respectively, with isotropic displacement parameters $1.2U_{eq}(O)$. Other H atoms were placed in calculated positions and constrained to ride on their parent

atoms at distances of 0.99 (methylene) and 0.98 Å (methyl). Their isotropic displacement parameters were fixed at 1.2 or 1.5 (for methyl groups) times $U_{eq}(\text{parent atom})$.

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO-SMN*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *X-SEED*.

The authors thank the Water Research Commission of South Africa, the NRF and the University of Stellenbosch for financial support.

References

- Barbour, L. J. (2001). *J. Supramol. Chem.* **1**, 189–191.
 Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
 Nonius (1998). *COLLECT*. Nonius BV, Delft, The Netherlands.
 Otwinowski, Z. & Minor, W. (1997). *Methods in Enzymology*, Vol. 276, *Macromolecular Crystallography*, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.
 Porath, J., Carlsson, J., Olsson, I. & Belfrage, G. (1975). *Nature (London)*, **258**, 598–599.
 Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. Versions 97-1. University of Göttingen, Germany.