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#### Key indicators

Single-crystal X-ray study T = 273 KMean  $\sigma(C-C) = 0.006 \text{ Å}$ Disorder in main residue R factor = 0.049 wR factor = 0.140 Data-to-parameter ratio = 16.0

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

# Tetrakis( $\mu$ -4-tolylsulfanylacetato- $\kappa^2 O:O'$ )bis[(N,N'-dimethylformamide- $\kappa O$ )copper(II)]

Molecules of the title dinuclear complex,  $[Cu_2(C_9H_9O_2S)_4-$ (C<sub>3</sub>H<sub>7</sub>NO)<sub>2</sub>], lie on centers of inversion located midway between the Cu atoms. Four carboxylate anions function as bridges to the two Cu atoms and each Cu atom is coordinated by an N,N'-dimethylformamide molecule.

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### Comment

Interest in the structural characterization of tetracarboxylatebridged dimeric copper(II) complexes has been increased since the crystal structure of copper acetate monohydrate was reported by van Niekerk & Schoening (1953). In general, a large number of copper(II) carboxylates can be formulated as  $[Cu_2(RCO_2)_4L_2]$ , where L denotes an axial oxygen- or nitrogen-donor ligand (Tong et al., 2002; Morgan et al., 2001; Tian et al., 2002; Calderazzo et al., 2003; Ponikiewski & Rothenberger, 2005). Among these dinuclear copper tetracarboxylate adducts are several having a mercapto entity in the acetate residue, such as, for example, the adducts phenylthioacetate pyridine (O'Reilly et al., 1984), isopropylthioacetate quinoline (Ouchi et al., 1983), 2,4-dichloro-5-methylphenylthioacetate acetone (Kennard et al., 1985), 2-(2-chlorophenylthio)-2methylpropionate water (Chan, Mak, Yip, Kennard et al., 1987), 2,4-dichloro-5-methylphenylthioacetate acetone (Smith et al., 1985) and 2-methyl-2-(phenylthio)propionate ethanol (Chan, Mak, Yip, Smith et al., 1987). In order to explore



The structure of (I). Displacement ellipsoids are drawn at the 30% probability level. Unlabeled atoms and atoms labeled with the suffix A are generated by the symmetry operation (1 - x, 1 - y, -z). Both the major and the minor components of the disordered 4-tolyl group are shown.

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further the coordination behavior of the dinuclear copper(II) complexes with tetracarboxylate which have a mercapto entity, we synthesized a new dinuclear copper(II) complex,  $[Cu_2(4-MPTA)_4(DMF)_2]$  (where 4-MPTA is the 4-methylphenylthioacetate anion and DMF is N,N'-dimethylformamide), (I).



Molecules of (I) lie on centers of inversion. As illustrated in Fig. 1, the two Cu<sup>II</sup> atoms are bridged by four carboxylate groups of four 4-MPTA anions forming a dinuclear structure, the Cu···Cu distance being 2.6409 (15) Å. Each Cu atom is also coordinated by an N,N'-dimethylformamide molecule. Each Cu<sup>II</sup> atom displays an approximate square-pyramidal geometry with four carboxylate O atoms from four 4-MPTA anions in the basal plane [Cu-O<sub>carboxyl</sub> = 1.961 (2)–1.974 (2) Å], and the O atom of the N,N'-dimethylformamide molecule in the apical position [Cu-O = 2.149 (2) Å]. The C-O bond distances of the carboxyl groups are nearly identical (Table 1), suggesting delocalization of electrons in these groups. No significant hydrogen-bonding interaction is observed in the crystal structure.

# **Experimental**

Blue vitriol (0.25 g, 1 mmol) and 4-methylphenylthioacetic acid (0.36 g, 2 mmol) were dissolved separately in a water–DMF (1:1) solution, and the two solutions were mixed slowly with stirring at room temperature. The pH was adjusted to 6 with triethylamine. Green crystals were obtained from the filtrate at room temperature over a period of several days. Analysis calculated for

 $C_{42}H_{50}Cu_2N_2O_{10}S_4{:}$ C 50.54, H 5.05, N 2.81%; found: C 50.64, H 5.15, N 2.71%.

14547 measured reflections

 $R_{\rm int} = 0.043$ 

 $\theta_{\rm max} = 27.8^\circ$ 

5397 independent reflections

3383 reflections with  $I > 2\sigma(I)$ 

#### Crystal data

 $\begin{bmatrix} Cu_2(C_9H_9O_2S)_4(C_3H_7NO)_2 \end{bmatrix} & Z = 2 \\ M_r = 984.11 & D_x = 1.418 \text{ Mg m}^{-3} \\ \text{Monoclinic, } P_1/n & \text{Mo } K\alpha \text{ radiation} \\ a = 10.997 (7) \text{ Å} & \mu = 1.16 \text{ mm}^{-1} \\ b = 7.808 (5) \text{ Å} & T = 273 (2) \text{ K} \\ c = 27.066 (17) \text{ Å} & \text{Prism, green} \\ \beta = 97.49 (1)^\circ & 0.30 \times 0.20 \times 0.18 \text{ mm} \\ V = 2304 (3) \text{ Å}^3 \\ \end{bmatrix}$ 

#### Data collection

Bruker SMART CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $T_{\min} = 0.757, T_{\max} = 0.812$ 

### Refinement

 Refinement on  $F^2$  H-atom parameters constrained

  $R[F^2 > 2\sigma(F^2)] = 0.049$   $w = 1/[\sigma^2(F_o^2) + (0.0643P)^2]$ 
 $wR(F^2) = 0.140$  where  $P = (F_o^2 + 2F_c^2)/3$  

 S = 1.09  $(\Delta/\sigma)_{max} = 0.001$  

 5397 reflections
  $\Delta\rho_{max} = 0.89$  e Å<sup>-3</sup>

 338 parameters
  $\Delta\rho_{min} = -0.47$  e Å<sup>-3</sup>

# Table 1

Selected geometric parameters (Å, °).

Cu1-O1	1.961 (2)	O1-C9	1.245 (4)
Cu1-O3	1.962 (3)	O2-C9	1.253 (4)
Cu1-O4 <sup>i</sup>	1.965 (3)	O3-C18	1.263 (4)
$Cu1 - O2^{i}$	1.974 (2)	O4-C18	1.250 (4)
Cu1-O5	2.149 (2)	O5-C19	1.263 (4)
Cu1-Cu1 <sup>i</sup>	2.6409 (15)		
O1-Cu1-O3	87.68 (10)	$O4^i$ -Cu1-O2 <sup>i</sup>	89.07 (11)
$O1-Cu1-O4^{i}$	90.54 (11)	O1-Cu1-O5	98.89 (9)
$O3-Cu1-O4^{i}$	167.97 (9)	O3-Cu1-O5	99.79 (10)
$O1-Cu1-O2^{i}$	167.81 (9)	O4 <sup>i</sup> -Cu1-O5	92.23 (10)
$O3-Cu1-O2^i$	90.18 (10)	$O2^i$ -Cu1-O5	93.30 (9)

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

One of the 4-tolyl groups was found to be disordered over two orientations. The occupancy factors for the disordered atoms C10–C16/C10'-C16' refined to 0.485 (4)/0.515 (4). The C-C bond lengths involving the disordered atoms were restrained to be equal. In addition, atoms in each disordered group were restrained to be coplanar. H atoms were placed in idealized positions and allowed to ride on their parent atoms, with C-H = 0.93-0.97 Å and  $U_{\rm iso}(H) = 1.2-1.5U_{\rm eq}(C)$ . A rotating group model was used for the methyl groups. Large  $U_{\rm eq}$  values for the methyl C atoms of the DMF molecule suggest disorder, but no suitable disorder model was found.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2002); software used to prepare material for publication: *SHELXL97*.

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