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## Key indicators

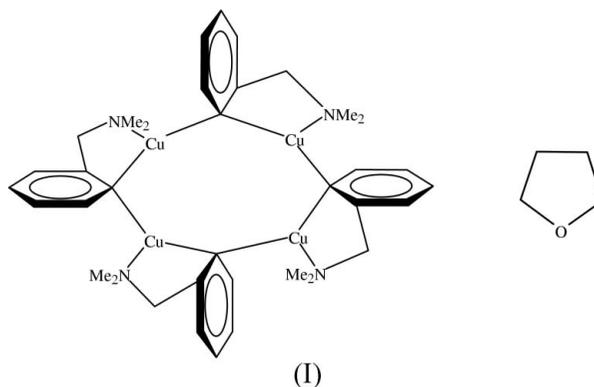
Single-crystal X-ray study  
 $T = 100\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$   
Disorder in solvent or counterion  
 $R$  factor = 0.019  
 $wR$  factor = 0.045  
Data-to-parameter ratio = 18.2For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.A second polymorph of *cyclo*-tetrakis[ $\mu_2$ -[(dimethyl-  
amino)methyl]phenyl- $\kappa^3\text{C}^2:\text{C}^2,\text{N}$ ]copper tetrahydro-  
furan solvateThe structure of the title compound,  $[\text{Cu}_4(\text{C}_9\text{H}_{12}\text{N})_4]\cdot\text{C}_4\text{H}_8\text{O}$ , has previously been reported in the monoclinic space group  $C2/c$ . We report here the structure of a second polymorph which crystallizes in the orthorhombic space group  $P2_12_12_1$ . The copper complex is located at a general position but exhibits local  $S_4$  symmetry.

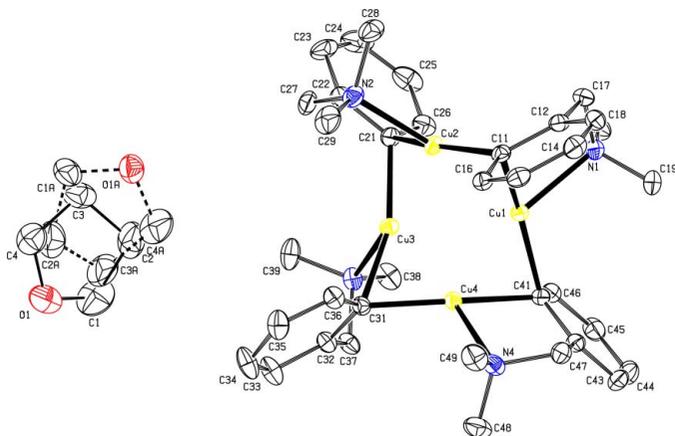
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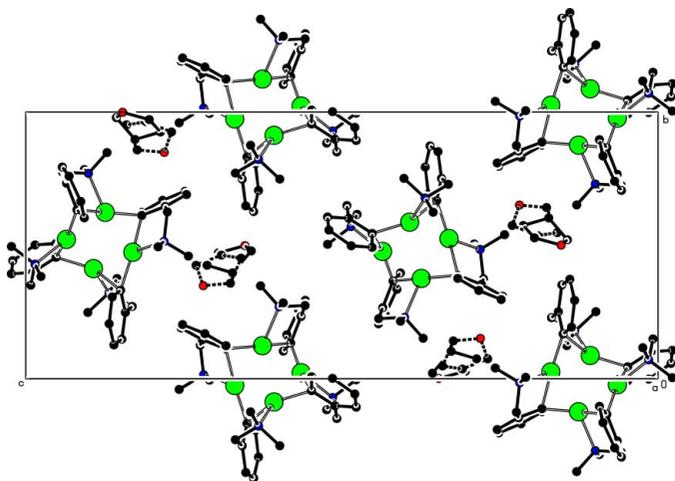
Online 31 May 2005

## Comment

The structure of the title compound, (I), has previously been reported crystallizing in the monoclinic space group  $C2/c$ , (II) (Janssen *et al.*, 1996). The asymmetric unit of (II) contains one half of the copper complex, the other half being identical by a twofold rotation, and one half-weight tetrahydrofuran (THF) molecule. The THF was found to be disordered about the twofold rotation axis. The solvent-free metal complex has also been reported, in space group  $P2_1/c$ , (III) [Spek & van Koten, 2004; Cambridge Structural Database (Version 5.26) and Conquest (Version 1.7) (Allen, 2002) refcode IQOTOX].The title compound, (I), is a second polymorph of the  $C2/c$  structure (Fig. 1) and crystallizes in the orthorhombic space group  $P2_12_12_1$ . The asymmetric unit consists of one complete copper complex and a single disordered THF molecule. The overall crystal structure is quite different from that of the previous polymorph, (II). In (II), the complex and the THF lie on a twofold rotation axis, with the THF molecules lying in the cavities of adjacent copper complexes (Fig. 2). In (I), however, whilst the complexes now line up roughly parallel to the  $a$  axis, the THF no longer sits between pairs of complexes and is instead displaced to the side (Fig. 3).The structure of the copper complex is essentially the same as previously reported, with the Cu atoms forming a 'butterfly' arrangement. The Cu—C and Cu—N bond lengths are comparable with those in (II), as are the Cu—C—Cu bond angles, which are once again very acute, at between  $70.33(8)$  and  $72.14(9)^\circ$ . The Cu...Cu distances in (II) are very short, at


**Figure 1**

A view of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms have been omitted for clarity. Both disorder components are shown.


**Figure 2**

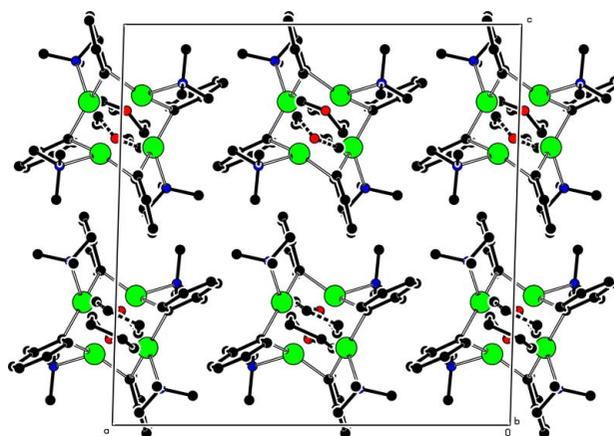
A packing diagram for the  $P2_12_12_1$  polymorph, (I), viewed along the  $a$  axis. The THF molecule is disordered over two positions, with relative occupancies of 73% and 27%; dashed lines indicate the minor disorder component.

2.3728 (3) and 2.3885 (3) Å. Polymorphs (I) and (III) also have very short Cu...Cu distances for Cu atoms that are adjacent in the metallocycle, at between 2.3773 (4) and 2.4118 (4) Å for (I), and between 2.372 (2) and 2.396 (2) Å for (III). However, in both cases, opposite Cu atoms have longer Cu...Cu distances, with one distinctly longer than the other [3.0065 (4) and 3.3810 (4) Å in (I), and 3.040 (2) and 3.380 (3) Å in (III)].

As a consequence of the different packing, the intermolecular Cu...Cu distances are different for the three crystal structures: in (I), the shortest Cu...Cu distance is 8.1780 (4) Å, in (II) it is 8.0630 (6) Å, and in (III) it is 6.508 (2) Å.

## Experimental

The title compound was obtained unintentionally during an attempt to synthesize a mixed Cu–Li complex. It was recrystallized from THF, yielding pale-yellow crystals of (I) suitable for X-ray diffraction.


**Figure 3**

A packing diagram for the  $C2/c$  polymorph, (II), viewed along the  $b$  axis. The THF molecule is disordered about the twofold axis.

### Crystal data

[Cu<sub>4</sub>(C<sub>9</sub>H<sub>12</sub>N<sub>4</sub>)<sub>4</sub>·C<sub>4</sub>H<sub>8</sub>O]  
 $M_r = 863.05$   
 Orthorhombic,  $P2_12_12_1$   
 $a = 9.4952$  (2) Å  
 $b = 13.2026$  (3) Å  
 $c = 31.5269$  (6) Å  
 $V = 3952.25$  (14) Å<sup>3</sup>  
 $Z = 4$   
 $D_x = 1.450$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation  
 Cell parameters from 53905 reflections  
 $\theta = 1.3$ – $27.5^\circ$   
 $\mu = 2.16$  mm<sup>-1</sup>  
 $T = 100$  (2) K  
 Block, pale yellow  
 $0.42 \times 0.24 \times 0.15$  mm

### Data collection

Bruker Nonius KappaCCD area-detector diffractometer  
 $\omega$  and  $\varphi$  scans  
 Absorption correction: multi-scan (MULABS; Blessing, 1995)  
 $T_{\min} = 0.59$ ,  $T_{\max} = 0.72$   
 53905 measured reflections

9022 independent reflections  
 8758 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.047$   
 $\theta_{\max} = 27.5^\circ$   
 $h = -12 \rightarrow 12$   
 $k = -17 \rightarrow 17$   
 $l = -40 \rightarrow 40$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.019$   
 $wR(F^2) = 0.045$   
 $S = 1.04$   
 9022 reflections  
 496 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0157P)^2 + 1.2769P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.32$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.26$  e Å<sup>-3</sup>  
 Absolute structure: Flack (1983), with 3956 Friedel pairs  
 Flack parameter:  $-0.013$  (6)

All H atoms were placed in geometrically idealized positions, with C–H distances in the range 0.95–0.99 Å, and constrained to ride on their parent atoms, with  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$  for methyl H atoms and  $1.2U_{\text{eq}}(\text{C})$  for all other H atoms. The THF molecule was refined using a disorder model, with relative occupancies of 73.1 (4)% and 26.9 (4)%.

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *DENZO* (Otwinowski & Minor, 1997); data reduction: *DENZO*; program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

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