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Di- μ -cyanato-bis[(cyanato- κN)(tetramethylethylenediamine- $\kappa^2 N$,N')copper(II)] and catena-poly[[μ_3 -cyanato- $\kappa^3 O$:N:N-bis[(cyanato- κN)(1,3-diaminopropane- $\kappa^2 N$,N')copper(II)]]- μ_3 -cyanato- $\kappa^3 N$:N:O]

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The two title Cu^{II} complexes, $[Cu_2(NCO)_4(tmeda)_2]$ (tmeda is tetramethylethylenediamine, C₆H₁₆N₂), (I), and [Cu(NCO)₂- $(pn)]_n$ (pn is 1,3-diaminopropane, $C_3H_{10}N_2$), (II), have been synthesized and their crystal structures determined. In (I), which lies about an inversion centre, each Cu centre possesses a distorted tetragonal-pyramidal geometry with four basal N atoms from two cyanate anions [Cu-N = 1.945(2)] and 1.948 (3) Å] and one tmeda molecule [Cu-N = 2.053 (2)] and 2.071 (2) Å], and one axial O atom [Cu-O = 2.737 (3) Å]from another cyanate anion. The two neighbouring Cu atoms in (I) are joined by a pair of cyanates in an end-to-end fashion, forming a dimer. In (II), each Cu centre adopts a distorted square-bipyramidal geometry, with four equatorial N atoms from two cyanates [Cu-N = 1.988 (2) and 2.007 (3) Å] and a pn ligand [Cu-N = 1.996 (3) and 2.011 (3) Å], and one apical N atom [Cu-N = 2.437 (3) Å] and an apical O atom [Cu-O =2.900 (3) Å] from two cyanates. In contrast with (I), the two neighbouring Cu atoms in (II) are bridged by two cyanates in an end-on fashion, to form a centrosymmetric dimeric unit. These units are further crosslinked, forming a two-dimensional network structure, via weak interactions between the bridging cyanate O atom and a neighbouring Cu atom, plus interactions of the amine H atoms with the cyanate O atoms and the terminal cyanate N atom.

Comment

The synthesis of cyanate complexes continues to be the subject of much interest, and intensive investigations have taken place as a result of their diverse structures and potential applications in magnetic materials (Diaz *et al.*, 2001; Grove *et al.*, 2001; Hernández *et al.*, 2001). It is noteworthy that, in most dinuclear Cu-cyanate complexes for which structural and magnetic data are available, the two Cu atoms are linked together by different bridging ligands (Boillot et al., 1985; Mallah et al., 1986; Kahn et al., 1989), while known homocyanate-bridged Cu₂ complexes with aliphatic amine coligands, such as substituted ethylenediamine (Mauro et al., 1990; Zukerman-Schpector et al., 1991), substituted diethylenetriamine (Escuer et al., 1999) and azacryptate (Harding et al., 1996), are scarce. To date, no two-dimensional structures linked by homocyanate-bridged Cu₂ moieties with aliphatic amine co-ligands have been reported. To gain a deeper insight into the structures and properties of homocyanate-bridged Cu₂ complexes, we report herein the syntheses and crystal structures of two new Cu-cyanate complexes with tetramethylethylenediamine (tmeda) and 1,3-diaminopropane (pn) as co-ligands, namely [Cu(tmeda)(N₂C₂O₂)]₂, (I), and [Cu₂- $(pn)_2(N_2C_2O_2)_2]_n$, (II).



Complex (I) (Fig. 1) is a centrosymmetric dinuclear structure with two end-to-end cyanate bridges, in which each Cu^{II} ion is pentacoordinated to three cyanate anions and one tmeda ligand to give a distorted tetragonal–pyramidal geometry. For atom Cu1, the basal plane is formed by one terminal cyanate atom (N1), one bridged cyanate atom (N2) and two atoms (N3 and N4) of the tmeda ligand, the apical site being occupied by one atom (O2ⁱ) of another bridged cyanate [symmetry code: (i) 2 - x, 2 - y, 2 - z]. In each dimer, the Cu···Cu distance is 5.4330 (12) Å [the closest Cu···Cu separation is 4.9814 (14) Å].

In (II), atom Cu1 is bonded to one terminal cyanate atom (N3), one bridged cyanate atom (N4ⁱⁱ) and two atoms (N1 and N2) of one pn ligand in the basal plane, along with one bridged cyanate N atom (N4) and one bridged cyanate O atom (O2ⁱⁱⁱ) in the apical sites, to form a distorted tetragonal–bipyramidal geometry [symmetry codes: (ii) 2 - x, 2 - y, -z; (iii) x, y - 1, z]. Two Cu atoms are linked by two cyanates in an end-on





A view of the molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms have been omitted for clarity [symmetry code: (i) 2 - x, 2 - y, 2 - z].

bonding mode to define a dimer (Fig. 2). These dimeric units are linked together to form a one-dimensional chain (Fig. 3) *via* weak interactions [Cu···O = 2.900 (3) Å]. These chains are then crosslinked *via* hydrogen bonds (Table 3) involving the terminal cyanate N atom [N···N = 3.188 (4) Å], the bridging cyanate O atom [N···O = 3.034 (4) Å] and the terminal cyanate O atoms [N···O = 3.120 (4) and 3.162 (4) Å], thus forming a two-dimensional network (Fig. 4). To our knowledge, such two-dimensional networks are rare in Cu–cyanate complexes. In (II), the cyanates exhibit two types of coordination mode, one of which is end-on within the dimer, while the second cyanate is end-to-end between dimers. The Cu···Cu distance within the dimers in (II) is 3.2137 (9) Å and that between neighbouring units is 5.375 (2) Å.

Interestingly, by replacing tmeda with pn as the co-ligand, the title Cu complexes change from a simple dimer to a twodimensional structure, and the bonding model of the bridging cyanate also changes. This may be attributed to the steric difference between the two aliphatic amine ligands. As tmeda



Figure 2

A view of the Cu^{II} coordination environment of (II), showing the atomlabelling scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms have been omitted for clarity [symmetry codes: (ii) 2 - x, 2 - y, -z; (iii) x, y - 1, z; (iv) 1 - x, 2 - y, -z].



Figure 3

The one-dimensional chain structure of (II) linked *via* the weak interactions, with a view of the unit cell.

is more sterically crowded than pn, in (I), the bridging cyanates adopt an end-to-end bonding fashion to reduce the steric repulsion between two tmeda ligands. In contrast with (I), the metal ion should be sterically unsaturated if (II) has a similar structure to (I). As a result, the smaller pn ligand allows the bonding model of the bridged cyanate to change from end-to-end to end-on and forces the Cu atom to coordinate further to the neighbouring O atom, thus forming a one-dimensional infinite chain with a higher coordination number.

In (I), the Cu–N(tmeda) bond lengths [2.053 (2)-2.071 (2) Å] are longer than the Cu–N(cyanate) distances [1.945 (2)-1.948 (3) Å; Table 1]. These data are similar to the corresponding distances in $\{[CuL(NCO)_n](ClO_4)\}_n$ (*L* is *N*,*N*,*N''*,*N''*-pentamethyl-3-azapentane-1,5-diamine; Vicente *et al.*, 1994). In contrast with (I), the Cu–N bond lengths in the basal plane of (II) [1.988 (2)–2.011 (3) Å; Table 2] show no distinct differences and are comparable with the corre-



Figure 4

The two-dimensional network of (II) crosslinked via hydrogen-bond interactions, viewed along the c axis.

distances in $[Cu_2(N_2C_2O_2)(medien)_2](ClO_4)_2$ sponding (medien is 4-methyldiethylenetriamine; Escuer et al., 1999). The apical Cu-N(cyanate) [2.437 (3) Å] or Cu-O(cyanate) distances [2.900 (3) Å] in (II) are longer than the Cu–N distances in the basal plane. A similar situation is also seen in (I). The N-Cu-N angles (two neighbouring N atoms) are in the range $85.35(8)-94.16(12)^{\circ}$ for (I) and 84.56(11)- $97.19 (11)^{\circ}$ for (II). Both ranges deviate from the ideal value of 90° , but the N(basal)-Cu-O(apical) angles in (I) $[87.20 (8)-98.81 (10)^{\circ}]$ deviate less than those in (II) $[74.01 (10)-100.84 (9)^{\circ}]$, indicating that the geometry of (II) is more distorted than that of (I).

In (I) and (II), the geometric parameters for the tmeda [1.480 (4)-1.505 (4) Å and $107.9 (2)-111.6 (2)^{\circ}]$ and pn $[1.460 (5)-1.466 (4) \text{ Å} and 113.9 (3)-119.2 (3)^{\circ}]$ ligands are normal for aliphatic amine-containing complexes (Kovbasyuk et al., 1997; Zukerman-Schpector et al., 1991). The cyanates are almost linear, with N-C and C-O bond lengths in the ranges 1.146(3)-1.169(4) and 1.188(3)-1.211(3) Å, respectively. The N-C-O angles $[177.8 (3)-178.8 (3)^{\circ}]$ are similar to those in other cyanate complexes (Rojo et al., 1989; Real et al., 1993; Otieno et al., 1993).

Experimental

To a solution of copper acetate (0.60 mmol, 119.79 mg) in ethanol (4 ml) was added tmeda (0.60 mmol, 69.73 mg) in ethanol (4 ml). After stirring for 5 min, an aqueous solution of sodium cyanate (2 ml; 0.60 mmol, 39.01 mg) was added and the mixture stirred for a further 5 min. The solution was filtered and the filtrate left to evaporate slowly in air. After one week, blue block-shaped crystals of (I) were isolated in 67% yield. Analysis calculated for C₁₆H₃₂Cu₂N₈O₄: C 36.43, H 6.11, N 21.24%; found: C 36.02, H 5.82, N 21.38%. Crystals of (II) were prepared in a similar manner, using the pn ligand as a starting material, resulting in an 11% yield. Analysis calculated for C₁₀H₂₀Cu₂N₈O₄: C 27.09, H 4.55, N 25.27%; found: C 26.32, H 4.61, N 24.73%.

> $D_x = 1.496 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 2262

reflections $\theta = 2.4 - 27.0^{\circ}$

 $\mu = 1.85~\mathrm{mm}^{-1}$

T = 293 (2) KBlock, blue

 $0.50 \times 0.45 \times 0.45 \text{ mm}$

Compound (I)

Crystal data

$[Cu_2(NCO)_4(C_6H_{16}N_2)_2]$
$M_r = 527.58$
Triclinic, P1
a = 7.656 (2) Å
b = 8.613(2) Å
c = 9.671 (2) Å
$\alpha = 78.893 \ (3)^{\circ}$
$\beta = 71.433 \ (3)^{\circ}$
$\gamma = 77.984 \ (3)^{\circ}$
$V = 585.8 (2) \text{ Å}^3$
Z = 1

Data collection

Bruker SMART APEX CCD area-	1883 reflections with $I > 2\sigma(I)$
detector diffractometer	$R_{\rm int} = 0.016$
ω scans	$\theta_{\rm max} = 25.0^{\circ}$
Absorption correction: multi-scan	$h = -9 \rightarrow 9$
(SADABS; Sheldrick, 1996)	$k = -10 \rightarrow 9$
$T_{\min} = 0.406, T_{\max} = 0.434$	$l = -11 \rightarrow 9$
2462 measured reflections	
2033 independent reflections	

Refinement

F ²	(1) 0.001
Refinement on F ²	$(\Delta/\sigma)_{\rm max} < 0.001$
$R[F^2 > 2\sigma(F^2)] = 0.030$	$\Delta \rho_{\rm max} = 0.53 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.082$	$\Delta \rho_{\rm min} = -0.37 \text{ e } \text{\AA}^{-3}$
S = 1.06	Extinction correction: SHELXL97
2033 reflections	(Sheldrick, 1997)
141 parameters	Extinction coefficient: 0.066 (5)
H-atom parameters constrained	
$w = 1/[\sigma^2(F_o^2) + (0.0519P)^2]$	
+ 0.0912P]	
where $P = (F_o^2 + 2F_c^2)/3$	

Table 1

Selected geometric parameters (Å, °) for (I).

Cu1-N2	1.945 (2)	Cu1-N4	2.071 (2)
Cu1-N1	1.948 (3)	Cu1-O2 ⁱ	2.737 (3)
Cu1-N3	2.053 (2)	Cu1-Cu1 ⁱ	5.4330 (12)
N2-Cu1-N1	94.16 (12)	N3-Cu1-N4	85.35 (8)
N2-Cu1-N3	174.55 (10)	N2-Cu1-O2 ⁱ	89.48 (9)
N1-Cu1-N3	90.61 (10)	N1-Cu1-O2 ⁱ	98.81 (10)
N2-Cu1-N4	90.68 (10)	N3-Cu1-O2 ⁱ	87.20 (8)
N1-Cu1-N4	164.90 (11)	N4-Cu1-O2 ⁱ	95.52 (8)

Symmetry code: (i) 2 - x, 2 - y, 2 - z.

Compound (II)

Crystal data

$[Cu(NCO)_2(C_3H_{10}N_2)]$	Mo $K\alpha$ radiation
$M_r = 221.72$	Cell parameters from 2190
Monoclinic, $P2_1/n$	reflections
a = 6.711 (2) Å	$\theta = 3.1 - 26.8^{\circ}$
b = 6.636 (2) Å	$\mu = 2.56 \text{ mm}^{-1}$
c = 19.230 (6) Å	T = 298 (2) K
$\beta = 100.049 \ (3)^{\circ}$	Block, blue
$V = 843.3 (4) \text{ Å}^3$	$0.20 \times 0.10 \times 0.10 \text{ mm}$
Z = 4	
$D_x = 1.746 \text{ Mg m}^{-3}$	

Data collection

Bruker SMART APEX CCD area-	1517 reflections with $I > 2\sigma(I)$
detector diffractometer	$R_{\rm int} = 0.033$
ω scans	$\theta_{\rm max} = 27.2^{\circ}$
Absorption correction: multi-scan	$h = -8 \rightarrow 8$
(SADABS; Sheldrick, 1996)	$k = -8 \rightarrow 7$
$T_{\min} = 0.629, \ T_{\max} = 0.784$	$l = -17 \rightarrow 24$
4002 measured reflections	
1859 independent reflections	

Table 2

Selected geometric parameters (Å, $^{\circ}$) for (II).

Cu1-N1	2.011 (3)	Cu1-N4	2.437 (3)
Cu1-N2	1.996 (3)	Cu1-O2 ⁱⁱⁱ	2.900 (3)
Cu1-N3	1.988 (2)	Cu1-Cu1 ⁱⁱ	3.2137 (9)
Cu1-N4 ⁱⁱ	2.007 (3)	Cu1-Cu1 ^{iv}	5.375 (2)
N3-Cu1-N2	88.59 (11)	N4 ⁱⁱ -Cu1-N4	87.89 (10)
N3-Cu1-N4 ⁱⁱ	91.74 (11)	N1-Cu1-N4	91.02 (11)
N2-Cu1-N4 ⁱⁱ	174.85 (12)	N3-Cu1-O2 ⁱⁱⁱ	93.13 (10)
N3-Cu1-N1	172.23 (12)	N2-Cu1-O2 ⁱⁱⁱ	74.01 (10)
N2-Cu1-N1	94.50 (12)	N4 ⁱⁱ -Cu1-O2 ⁱⁱⁱ	100.84 (9)
N4 ⁱⁱ -Cu1-N1	84.56 (11)	N1-Cu1-O2 ⁱⁱⁱ	80.90 (10)
N3-Cu1-N4	95.68 (11)	N4-Cu1-O2 ⁱⁱⁱ	167.40 (8)
N2-Cu1-N4	97.19 (11)		

Symmetry codes: (ii) 2 - x, 2 - y, -z; (iii) x, y - 1, z; (iv) 1 - x, 2 - y, -z.

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0497P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.035$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.090$	$(\Delta/\sigma)_{\rm max} < 0.001$
S = 1.04	$\Delta \rho_{\rm max} = 0.58 \ {\rm e} \ {\rm \AA}^{-3}$
1859 reflections	$\Delta \rho_{\rm min} = -0.38 {\rm e} {\rm \AA}^{-3}$
125 parameters	
H atoms treated by a mixture of	
independent and constrained	
refinement	

Table 3

Hydrogen-bonding and contact geometry (Å, °) for (II).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
		/->	/	
$N1 - H1A \cdots O1^{v}$	0.86 (3)	2.28 (3)	3.120 (4)	163 (3)
$N2-H2A\cdots N3^{1}$	0.70 (3)	2.55 (3)	3.188 (4)	153 (3)
$N2-H2B\cdots O1^{v_1}$	0.81 (4)	2.42 (4)	3.162 (4)	153 (3)
$N2-H2B\cdots O2^{m}$	0.81 (4)	2.56 (3)	3.034 (4)	119 (3)
$N1 - H1B \cdot \cdot \cdot N3^{ii}$	0.84 (4)	2.53 (4)	3.318 (4)	156 (4)

Symmetry codes: (ii) 2-x, 2-y, -z; (iii) x, y-1, z; (iv) 1-x, 2-y, -z; (v) 2-x, 1-y, -z; (vi) 1-x, 1-y, -z.

In (I), all the H atoms were constrained to an ideal geometry, with C3–H and C4–H distances of 0.97 Å and all other C–H distances 0.96 Å. In (II), the amine H atoms were found from a difference Fourier map and refined freely, with all other H atoms placed in geometrically idealized positions and constrained to ride on their parent atoms, with C–H distances of 0.97 Å.

For both compounds, data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2000); software used to prepare material for publication: *SHELXTL*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: GG1183). Services for accessing these data are described at the back of the journal.

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