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Key indicators

Single-crystal X-ray study
T = 292 K
Mean $\sigma(\text{C}-\text{C}) = 0.011 \text{ \AA}$
R factor = 0.073
wR factor = 0.189
Data-to-parameter ratio = 16.3

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

A three-dimensional polymeric mixed-valence copper complex: poly[tri- μ -azido- μ -di-2-pyridyl-diazene-dicopper(I,II)]

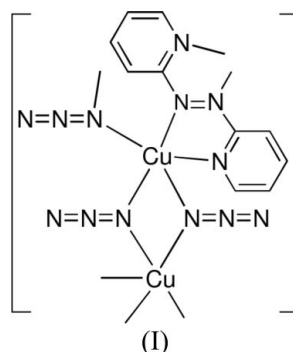
The title complex, $[\text{Cu}_2(\text{N}_3)_3(\text{C}_{10}\text{H}_8\text{N}_4)]_n$, shows a three-dimensional polymeric structure. Each Cu atom is surrounded by five N atoms from three bridging azide ligands and one di-2-pyridyldiazene ligand [$\text{Cu}-\text{N} = 2.002(5), 2.317(6), 2.020(4), 1.995(5)$ and $1.985(3) \text{ \AA}$], forming a CuN_5 trigonal bipyramid. One $\mu_{1,1}$ -azide bridges two Cu atoms and the other two $\mu_{1,1}$ -azides bridge pairs of Cu atoms, forming a $-\text{Cu}-\text{N}-\text{Cu}-\text{N}_2-$ chain. The di-2-pyridyldiazene bridges two Cu atoms of different $-\text{Cu}-\text{N}-\text{Cu}-\text{N}_2-$ chains, forming a three-dimensional polymeric structure. On the basis of charge equilibrium, the presence of one Cu^{II} and one Cu^{I} ion are deduced and the two ions are disordered because of the occurrence of only one unique Cu atom in the structure.

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Comment

μ -Azidocopper complexes have received intense attention due to their structural and magnetic diversity. The azide anion can bridge two copper ions either in an 'end-on' ($\mu_{1,1}-\text{N}_3$) fashion or an 'end-to-end' ($\mu_{1,3}-\text{N}_3$) fashion (Comarmond *et al.*, 1982; Sikorav *et al.*, 1984; Agnus *et al.*, 1979; Kahn *et al.*, 1983). Sometimes both bridging fashions are present in one complex (Felthouse & Hendrickson, 1978; Banci *et al.*, 1984). We report here the preparation and crystal structure of a three-dimensional azido mixed-valence copper complex, (I).



As shown in Fig. 1, each Cu atom in (I) is surrounded by five N atoms of three azide bridging ligands and one di-2-pyridyldiazene ligand, forming a CuN_5 trigonal bipyramid. A linear $\mu_{1,1}$ -azide ligand bridges two Cu atoms, and two linear $\mu_{1,1}$ -azide anions bridge two Cu atoms, forming a $-\text{Cu}-\text{N}-\text{Cu}-\text{N}_2-$ chain; di-2-pyridyldiazene bridges two Cu atoms of different $-\text{Cu}-\text{N}-\text{Cu}-\text{N}_2-$ chains, forming a three-dimensional structure. Charge equilibrium calculation suggests that there present one Cu^{II} and one Cu^{I} ion, which are disordered because of the existence of only one unique Cu atom in the structure. Therefore, the bond distances and the bond angles around the Cu^{I} ion should be the average values. One azide group (N6/N7/N8) lies along a twofold rotation axis.

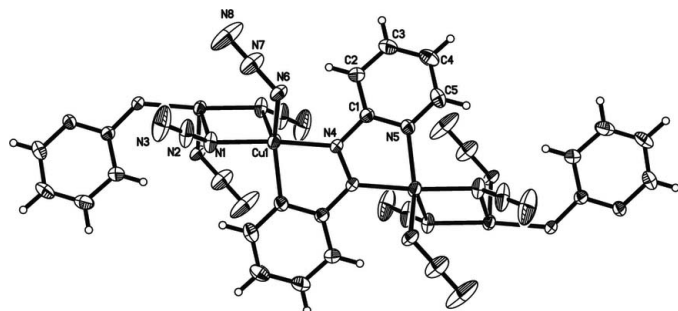


Figure 1
A view of the coordination environment in the title compound, shown with 30% probability displacement ellipsoids.

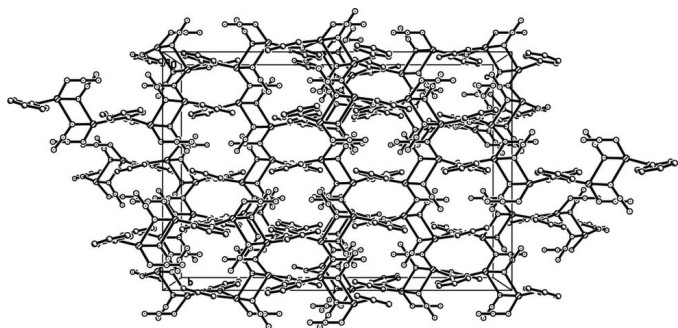


Figure 2
The crystal structure of the title compound, viewed along the *a* axis. H atoms have been omitted.

The Cu–N distances found in (I) are in accord with the Cu–N distances found in other copper complexes containing an azide bridging ligand (You, 2005; Ghoshal *et al.*, 2004). The coordinating azide anions are nearly linear and show bent coordination modes with the metal atoms (Table 1). The N–N bond of di-2-pyridyldiazene is longer than the N–N bonds of the azide ligand. The Cu···Cu distances, which are bridged by one azide bridging ligand, two azide bridging ligands and di-2-pyridyldiazene, are 3.492 (4), 3.342 (3) and 4.678 (4) Å, respectively. The dihedral angle between the two pyridine ring planes of di-2-pyridyldiazene is 29.8 (1)°.

Experimental

Crystals of (I) were prepared by the multilayer diffusion method. In a typical preparation, a solution of NaN₃ (19.5 mg, 0.3 mmol) in water (10 ml) was placed in a tube to which was added a buffer layer of ethanol/water (20 ml, 1:1 *v/v*). Finally, a solution of CuNO₃ (25.1 mg, 0.2 mmol) and di-2-pyridyldiazene (18.4 mg, 0.1 mmol) in ethanol (10 ml) was added to form a third layer. Black block-shaped crystals were formed on diffusion of the reactants over a period of two weeks. Analysis found: C 27.21, H 1.80, N 41.26%; calculated for C₁₀H₈Cu₂N₁₃: C 27.46, H 1.84, N 41.63%.

Crystal data

[Cu₂(N₃)₃(C₁₀H₈N₄)]
M_r = 437.37
 Orthorhombic, *Fddd*
a = 13.520 (3) Å
b = 18.127 (4) Å
c = 26.501 (5) Å
V = 6495 (2) Å³

Z = 16
D_x = 1.789 Mg m⁻³
 Mo *K*α radiation
 μ = 2.64 mm⁻¹
T = 292 (3) K
 Block, black
 0.12 × 0.08 × 0.06 mm

Data collection

Rigaku R-Axis RAPID IP
 diffractometer
 ω scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
*T*_{min} = 0.774, *T*_{max} = 0.852

18086 measured reflections
 1876 independent reflections
 1808 reflections with *I* > 2σ(*I*)
*R*_{int} = 0.077
 θ _{max} = 27.5°

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.074
wR (*F*²) = 0.189
S = 1.24
 1876 reflections
 115 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0802P)^2 + 91.4266P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 (Δ/σ)_{max} = 0.003
 $\Delta\rho$ _{max} = 1.35 e Å⁻³
 $\Delta\rho$ _{min} = -0.45 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Cu1–N1	2.001 (5)	Cu1–N6	1.986 (3)
Cu1–N1 ⁱ	2.318 (6)	N1–N2	1.213 (9)
Cu1–N4 ⁱⁱ	2.020 (4)	N4–N4 ⁱⁱ	1.319 (8)
Cu1–N5	1.995 (5)	N6–N7	1.210 (10)
N1–Cu1–N4 ⁱⁱ	167.3 (2)	N4 ⁱⁱ –Cu1–N1 ⁱ	93.60 (19)
N1–Cu1–N1 ⁱ	78.8 (2)	N5–Cu1–N1	95.5 (2)
N2–N1–Cu1	123.7 (5)	N5–Cu1–N1 ⁱ	113.5 (2)
N2–N1–Cu1 ⁱ	116.9 (5)	N7–N6–Cu1 ⁱⁱⁱ	118.44 (16)
N3–N2–N1	177.4 (10)	N8–N7–N6	180

Symmetry codes: (i) $-x + 1, -y + 1, -z + 1$; (ii) $-x + \frac{5}{4}, y, -z + \frac{5}{4}$; (iii) $-x + \frac{5}{4}, -y + \frac{5}{4}, z$.

All H atoms were placed in idealized positions and constrained to ride on their parent C atoms, with C–H = 0.93 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C). The highest residual density peak is 0.96 Å from the Cu1 atom.

Data collection: *CrystalStructure* (Rigaku/MS, 2004); cell refinement: *CrystalStructure*; data reduction: *CrystalStructure*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 1998); software used to prepare material for publication: *SHELXTL*.

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