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Key indicators

Single-crystal X-ray study T = 297 KMean $\sigma(\text{C}-\text{C}) = 0.010 \text{ Å}$ R factor = 0.044 wR factor = 0.114 Data-to-parameter ratio = 14.0

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

Bis(μ -4-nitropyridine *N*-oxide- $\kappa^2 O$:*O*)-bis[dibromocopper(II)]

The title compound, $[Cu_2Br_4(C_5H_4N_2O_3)_2]$, is a centrosymmetric dimer that contains a central planar Cu_2O_2 rhombus with the O atom of the N-oxide group bridging two Cu atoms $[Cu-O-Cu = 109.08 (19)^{\circ}$ and $O-Cu-O = 70.92 (19)^{\circ}]$. The two-dimensional sheet structure is built up *via* semicoordinate intermolecular $Cu \cdots Br$ bonds [3.226 (1) and 3.810 (1) Å] and weak hydrogen bonding $[C \cdots Br = 3.733 (7), 3.781 (7) \text{ and } 3.830 (7) \text{ Å}]$.

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Comment

The molecule of the title compound, (I), lies on a center of inversion at (-x, 1 - y, 1 - z). As shown in Fig. 1, the central core consists of a Cu₂O₂ rhombus with the O atoms of the *N*-oxide ligand bridging the two Cu atoms. The Cu–O–Cu angles are 109.08 (19)° and O–Cu–O angles are 70.92 (19)°. The CuBr₂ planes are both inclined at 10.8 (1)° to the central Cu₂O₂ planar core. The perpendicular *N*-oxide ligands are also tilted out of the plane of the central core by 71.6 (2)°. The terminal nitro groups are almost coplanar with the pyridine rings [8.5 (1.5)°], indicating extended conjugation of the aromatic system.



The extended two-dimensional system is more complicated. There are two semi-coordinate Cu···Br bonds that complete the Cu coordination sphere, *viz*. Cu1-Br1ⁱⁱ [3.226 (1) Å; symmetry code: (ii) $\frac{1}{2} - x$, $y - \frac{1}{2}$, z] and Cu1-Br1ⁱⁱⁱ



The molecular structure of (I), showing 30% probability displacement ellipsoids. The suffix A indicates the symmetry code (-x, 1 - y, 1 - z).

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Figure 2

The complete coordination sphere around the Cu centers for the central molecule only. Dashed lines indicate semi-coordinate bonding. H atoms have been omitted for clarity.



Figure 3

Hydrogen bonding (dashed lines) within the extended sheet.

[3.810 (1) Å; symmetry code: (iii) -x, 2 - y, 1 - z; see Table 1 for the complete geometrical parameters], resulting in a very distorted octahedron which is shown in Fig. 2. This semicoordinate Cu···Br bonding links the Cu₂O₂Br₄ units into a sheet structure parallel to the (001) plane in the unit cell. The inorganic core is sheathed by the 4-nitropyridine groups, which also hold the sheet together *via* non-classical bifurcated hydrogen bonding (see Table 2) between C6 and Br1ⁱ and Br2ⁱ [symmetry code: (i) $x - \frac{1}{2}, \frac{3}{2} - y, 1 - z$] and a weak interaction between C2 and Br1ⁱⁱ. This synthon is shown in Fig. 3. The terminal nitro groups on adjacent sheets interdigitate and form a herring-bone pattern (Fig. 4).

There are relatively few *N*-oxide dihalo copper(II) structures in the Cambridge Structural Database (Version 5.26; Allen, 2002). Excluding solvent-coordinated or internally coordinated copper systems there are only nine reports: CSD refcode CUCPYO (Schäfer *et al.*, 1965), CUCPYO11 (Sager *et al.*, 1967), DURYIY (Nepveu *et al.*, 1986), HULZOD (Ivashevskaja *et al.*, 2002), PHPYOC, PYOCDB (Watson & Johnson, 1971), QQQBWD, QQQBWG, QQQBWJ (Kidd *et*



Figure 4 Packing of (I) viewed down the *a* axis.

al., 1967). Only DURYIY and PYOCDB are bromo species, the remainder are all chloro-substituted. The title compound, (I), has many similarities to the above compounds.

Experimental

A 1:1 solution of 4-nitropyridine *N*-oxide and $CuBr_2$ in ethanol was left to slowly evaporate over 7 d. Just before complete dryness, the product formed as red plates in *ca* 60% yield.

Crystal data

$\begin{bmatrix} Cu_2Br_4(C_5H_4N_2O_3)_2 \end{bmatrix} \\ M_r = 726.92 \\ \text{Orthorhombic, } Pbca \\ a = 10.044 (2) \text{ Å} \\ b = 7.2816 (15) \text{ Å} \\ c = 25.006 (5) \text{ Å} \\ V = 1828.9 (6) \text{ Å}^3 \\ Z = 4 \end{bmatrix}$	Mo K α radiation Cell parameters from 5682 reflections $\theta = 2.6-27.0^{\circ}$ $\mu = 11.10 \text{ mm}^{-1}$ T = 297 (2) K Plate, red 0.44 × 0.15 × 0.01 mm
$D_x = 2.640 \text{ Mg m}^{-3}$	
Data collection	
Bruker–Siemens SMART APEX diffractometer ω scans Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2001) $T_{min} = 0.081, T_{max} = 0.898$ 24047 measured reflections	1654 independent reflections 1400 reflections with $I > 2\sigma(I)$ $R_{int} = 0.080$ $\theta_{max} = 25.2^{\circ}$ $h = -12 \rightarrow 12$ $k = -8 \rightarrow 8$ $l = -30 \rightarrow 30$
Refinement	
Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.044$ $wR(F^2) = 0.114$ S = 1.07 1654 reflections 118 parameters H-atom parameters constrained	$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.0511P)^2 \\ &+ 9.3212P] \\ &\text{where } P = (F_o^2 + 2F_c^2)/3 \\ &(\Delta/\sigma)_{\text{max}} < 0.001 \\ &\Delta\rho_{\text{max}} = 1.92 \text{ e } \text{\AA}^{-3} \\ &\Delta\rho_{\text{min}} = -0.62 \text{ e } \text{\AA}^{-3} \end{split}$

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Selected	geometric	parameters	(Å,	°)	

Table 1

Br1-Cu1	2.3403 (10)	Cu1-O1 ⁱ	2.034 (4)
Br2-Cu1	2.3178 (11)	Cu1-Br1 ⁱⁱ	3.2261 (11)
Cu1-O1	1.982 (4)	Cu1-Br1 ⁱⁱⁱ	3.8103 (12)
$O1-Cu1-O1^{i}$	70.92 (19)	Br2-Cu1-Br1 ⁱⁱ	110.31 (4)
O1-Cu1-Br2	162.07 (13)	Br1-Cu1-Br1 ⁱⁱ	88.09 (3)
O1 ⁱ -Cu1-Br2	93.35 (12)	O1-Cu1-Br1 ⁱⁱⁱ	84.98 (13)
O1-Cu1-Br1	97.70 (13)	O1 ⁱ -Cu1-Br1 ⁱⁱⁱ	115.93 (13)
O1 ⁱ -Cu1-Br1	166.38 (13)	Br2-Cu1-Br1 ⁱⁱⁱ	94.74 (4)
Br2-Cu1-Br1	98.93 (4)	Br1-Cu1-Br1 ⁱⁱⁱ	69.07 (3)
O1-Cu1-Br1 ⁱⁱ	76.86 (13)	Br1 ⁱⁱ -Cu1-Br1 ⁱⁱⁱ	148.65 (3)
O1 ⁱ -Cu1-Br1 ⁱⁱ	82.11 (13)	Cu1-O1-Cu1 ⁱ	109.08 (19)

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

Table 2Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$C6-H6A\cdots Br1^{iv}$	0.93	2.89	3.733 (7)	151 136
$C2-H2A\cdots Br1^{ii}$	0.93	3.05	3.830 (7)	143

Symmetry codes: (ii) $\frac{1}{2} - x$, $y - \frac{1}{2}$, z; (iv) $x - \frac{1}{2}$, $\frac{3}{2} - y$, 1 - z.

All H atoms were positioned geometrically and refined using a riding model, with $U_{\rm iso}({\rm H})$ values constrained to be $1.2U_{\rm eq}$ of the carrier atom and C-H = 0.93 Å. There are large residuals of *ca* 1.92 e Å⁻³ approximately 1.1 Å from Br2 and Br1. These are due to imperfectly corrected absorption effects and could not be eliminated.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to solve structure: *XS* in *SHELXTL* (Bruker, 2001); program(s) used to refine structure: *XL* in *SHELXTL*; molecular graphics: *XP* in *SHELXTL*; software used to prepare material for publication: *XCIF* in *SHELXTL*.

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