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Key indicators

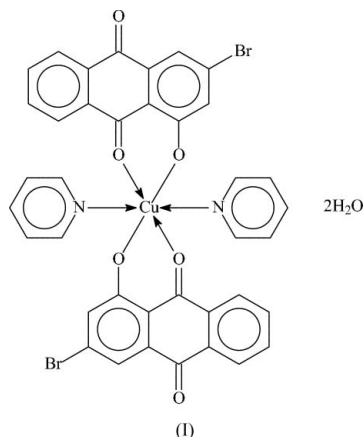
Single-crystal X-ray study
 $T = 295$ K
Mean $\sigma(\text{C}-\text{C}) = 0.007$ Å
 R factor = 0.048
 wR factor = 0.144
Data-to-parameter ratio = 15.0For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Bis(3-bromo-1-oxidoanthraquinone- $\kappa^2\text{O}^1, \text{O}^9$)-
bis(pyridine- κN)copper(II) dihydrateThe Cu atom in the title compound, $[\text{Cu}(\text{C}_{14}\text{H}_6\text{BrO}_3)_2(\text{C}_5\text{H}_5\text{N})_2] \cdot 2\text{H}_2\text{O}$, lies on a center of inversion; it is chelated by the bromo-substituted hydroxyanthraquinone molecule and is coordinated by the pyridine molecules in an all-*trans* octahedral geometry.

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Comment

Our previous study has documented the structure of the zinc derivative of a bromo-substituted 1-hydroxyanthraquinone anion (Ali *et al.*, 2005). This zinc complex crystallizes from pyridine as a bis-pyridine adduct in an all-*trans* octahedral environment. The corresponding title copper complex, (I), adopts the identical geometry, but the compound crystallizes as a dihydrate (Fig. 1). The Cu atom lies on a special position of $\bar{1}$ site symmetry; the chelating O atoms form a square, and the N atoms of the heterocycle occupy the other two octahedral sites. Hydrogen bonds (Table 2) link the molecule to the uncoordinated water molecule to give rise to a linear chain.

Experimental

3-Bromo-1-hydroxyanthraquinone (0.50 g, 1.65 mmol) and copper acetate monohydrate (0.16 g, 0.82 mmol) were heated in ethanol for several hours. The solid that was isolated upon removal of the solvent was recrystallized from pyridine to furnish brown prisms.

Crystal data

 $[\text{Cu}(\text{C}_{14}\text{H}_6\text{BrO}_3)_2(\text{C}_5\text{H}_5\text{N})_2] \cdot 2\text{H}_2\text{O}$ $M_r = 861.97$ Triclinic, $P\bar{1}$ $a = 8.8121$ (8) Å $b = 10.0253$ (9) Å $c = 10.1720$ (9) Å $\alpha = 69.474$ (1)° $\beta = 80.252$ (1)° $\gamma = 81.477$ (1)° $V = 825.6$ (1) Å³ $Z = 1$ $D_x = 1.734$ Mg m⁻³Mo $K\alpha$ radiation

Cell parameters from 2123

reflections

 $\theta = 2.4$ – 25.4 ° $\mu = 3.14$ mm⁻¹ $T = 295$ (2) K

Prism, brown

 $0.36 \times 0.18 \times 0.18$ mm

Data collection

Bruker SMART area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 1999)
 $T_{\min} = 0.347$, $T_{\max} = 0.568$
 6931 measured reflections

3565 independent reflections
 2240 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.029$
 $\theta_{\text{max}} = 27.2^\circ$
 $h = -10 \rightarrow 11$
 $k = -12 \rightarrow 12$
 $l = -12 \rightarrow 13$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.144$
 $S = 1.01$
 3565 reflections
 238 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0832P)^2 + 0.0828P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.85 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.47 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

Cu1—O1	1.932 (3)	Cu1—O2	2.229 (3)
Cu1—N1	2.077 (4)		
O1—Cu1—O1 ⁱ	180	O2—Cu1—O2 ⁱ	180
O1—Cu1—O2	86.3 (1)	O2—Cu1—N1	91.4 (1)
O1—Cu1—O2 ⁱ	93.7 (1)	O2—Cu1—N1 ⁱ	88.6 (1)
O1—Cu1—N1	90.8 (1)	N1—Cu1—N1 ⁱ	180
O1—Cu1—N1 ⁱ	89.2 (1)		

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

Table 2

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1w—H1w1 \cdots O1	0.86 (1)	2.23 (4)	2.948 (5)	142 (6)
O1w—H1w2 \cdots O3 ⁱⁱ	0.86 (1)	2.25 (4)	3.018 (6)	149 (6)

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

The C-bound H atoms were positioned geometrically ($C-H = 0.93 \text{ \AA}$) and were included in the refinement in the riding-model approximation, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The water H atoms were located in a difference Fourier map, and were refined with distance restraints of $O-H = 0.85 (1) \text{ \AA}$ and $H\cdots H = 1.39 (1) \text{ \AA}$.

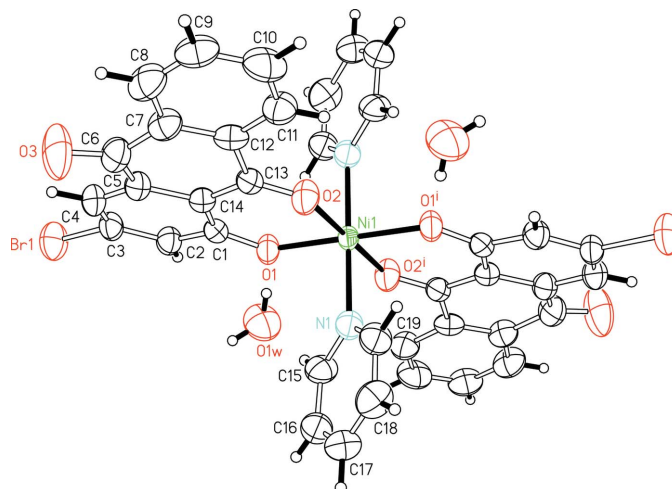


Figure 1

ORTEP plot (Johnson, 1976) of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as spheres of arbitrary radii. The Cu atom lies at the center of inversion ($\frac{1}{2}, \frac{1}{2}, \frac{1}{2}$) [symmetry code: (i) $1 - x, 1 - y, 1 - z$].

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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