

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Dibromido(4,7-diazadecane-1,10-diamine)copper(II)

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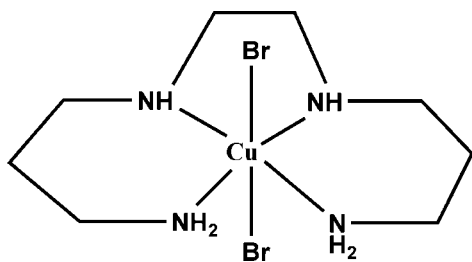
Received 8 August 2011; accepted 10 August 2011

Key indicators: single-crystal X-ray study; $T = 110$ K; mean $\sigma(\text{C}-\text{C}) = 0.014$ Å; R factor = 0.071; wR factor = 0.183; data-to-parameter ratio = 20.3.

In the title compound, $[\text{CuBr}_2(\text{C}_8\text{H}_{22}\text{N}_4)]$, the Cu^{II} atom is six-coordinate forming a distorted octahedral complex and is bonded to two axial bromide anions and four equatorial nitrogen donors. The equatorial $\text{Cu}-\text{N}$ bond distances range from 2.005 (8) to 2.046 (8) Å while the axial $\text{Cu}-\text{Br}$ distances are 2.8616 (17) and 2.9402 (17) Å, thus the six-coordinate Cu complex shows the usual Jahn–Teller distortion. All amine hydrogen atoms participate in either inter- or intramolecular hydrogen bonding to the Br anions.

Related literature

For related structures, see: Lee *et al.* (1986). For other related literature, see: Jahn & Teller (1937).



Experimental

Crystal data

 $[\text{CuBr}_2(\text{C}_8\text{H}_{22}\text{N}_4)]$
 $M_r = 397.66$

 Orthorhombic, $P2_12_12_1$
 $a = 6.9666$ (4) Å

 $b = 8.4146$ (6) Å
 $c = 24.0261$ (15) Å
 $V = 1408.45$ (15) Å³
 $Z = 4$

 Mo $K\alpha$ radiation
 $\mu = 7.20$ mm⁻¹
 $T = 110$ K
 $0.47 \times 0.31 \times 0.22$ mm

Data collection

 Oxford Diffraction Xcalibur diffractometer with a Ruby detector
 Absorption correction: analytical (*CrysAlis PRO*; Oxford)

 Diffraction, 2007
 $T_{\text{min}} = 0.157$, $T_{\text{max}} = 0.282$
 9561 measured reflections
 2758 independent reflections
 2262 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.072$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.071$
 $wR(F^2) = 0.183$
 $S = 1.07$
 2758 reflections
 136 parameters

 24 restraints
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 2.51$ e Å⁻³
 $\Delta\rho_{\text{min}} = -1.98$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1C}\cdots\text{Br2}^{\text{i}}$	0.92	2.66	3.466 (9)	147
$\text{N1}-\text{H1D}\cdots\text{Br2}$	0.92	2.80	3.339 (8)	119
$\text{N2}-\text{H2C}\cdots\text{Br1}^{\text{ii}}$	0.93	2.66	3.407 (8)	138
$\text{N2}-\text{H2C}\cdots\text{Br2}$	0.93	3.01	3.519 (7)	116
$\text{N3}-\text{H3C}\cdots\text{Br1}$	0.93	2.90	3.409 (8)	116
$\text{N4}-\text{H4C}\cdots\text{Br2}^{\text{i}}$	0.92	2.60	3.515 (8)	171
$\text{N4}-\text{H4D}\cdots\text{Br2}^{\text{iii}}$	0.92	2.69	3.425 (8)	138
$\text{N4}-\text{H4D}\cdots\text{Br1}$	0.92	2.94	3.433 (8)	115

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

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2007); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis RED* (Oxford Diffraction, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

RJB wishes to acknowledge the NSF–MRI program (grant No. CHE-0619278) for funds to purchase the diffractometer.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2444).

References

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supporting information

Acta Cryst. (2011). E67, m1266 [doi:10.1107/S160053681103251X]

Dibromido(4,7-diazadecane-1,10-diamine)copper(II)

Gervas E. Assey, Ray J. Butcher and Yilma Gultneh

S1. Comment

In this study, the title compound was prepared and its structure determined by X-ray analysis. Owing to the Jahn-Teller distortion (Jahn & Teller, 1937), the Cu(II) center adopts an axially distorted octahedral CuN_4Br_2 conformation with the axial positions are occupied by the bromide anions. The equatorial positions are occupied by the N_4 set of donor nitrogen atoms and the Cu1 lies in the N_4 plane; maximum deviation of any atom from the mean-plane formed by CuN_4 fragment being 0.042 (4) for N3. The structure of a related compound containing the same linear tetramine, has been reported (Lee *et al.* 1986) and its structural features compared with those of other linear Cu(II) aliphatic tetraamines of the type $\text{H}_2\text{N}(\text{CH}_2)_l\text{NH}-(\text{CH}_2)_m\text{NH}(\text{CH}_2)_n\text{NH}_2$ where l , m and n are 2 or 3. From this it can be seen that in the title complex, the equatorial Cu—N bond distances range from 2.005 (8) to 2.046 (8) Å and are in the normal range for such bonds. However, the axial Cu—Br distances are elongated at 2.8616 (17) and 2.9402 (17) Å, thus the 6-coordinate Cu complex shows the usual Jahn-Teller distortion. All amine H's participate in either inter or intramolecular hydrogen bonding to the Br anions.

S2. Experimental

The title compound was obtained as a byproduct of an attempt to prepare copper complexes of ethylenediamine *N,N*-bis(propylsalicylalimine). A solution of *N,N*-bis(3-aminopropylethylene)diamine (5 g, 30.52 mmol) in methanol (20 ml) was added dropwise to a solution of salicylaldehyde (7.45 g, 61.04 mmol) in methanol (20 ml). The mixture was refluxed overnight while stirring with magnetic stirrer. Then the reaction mixture was evaporated under reduced pressure. An oily orange product was obtained which later solidified into a yellow compound, [2-(3-amino-propylamino)-ethyl]-propane-1,3-diamine-bis(salicyladimine), used as a ligand (H_2L_4) in the subsequent reaction. The synthesis of the title complex was achieved by the reaction of CuBr (1.5 g, 10.5 mmol) in methanol (20 ml) with of the ligand H_2L_4 (2 g, 5.2 mmol) dissolved in CH_2Cl_2 (25 ml). The ligand solution was added drop-wise to the solution of the metal salt and stirred at room temperature for 24 h. The mixture was then concentrated by evaporation under reduced pressure to afford a thick greenish liquid. Part of the complex was dissolved in dimethyl formamide (DMF), filtered and layered with diethyl ether for slow diffusion and X-ray quality crystals were obtained.

S3. Refinement

H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms with a C—H distance of 0.99 Å and N—H distances of 0.92 (primary amine) and 0.93 (secondary amine) with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$. Even though a face-indexed absorption correction was carried out, the thermal parameters for C3, C6, C7, and N4 atoms did not behave well and thus were restrained using ISOR command in SHELXL. The crystal was originally refined as a racemic twin with components 0.87 (3):0.13 (3). However, as the absolute configuration was not established unambiguously, the data were merged. In addition, the highest peak ($2.50 \text{ e}^-/\text{Å}^3$, 0.70 Å from Cu) and deepest hole (-1.98

$e^{-}/\text{\AA}^3$, 0.54 \AA from Br2) are indicative of the problems with both the racemic twinning and absorption effects.

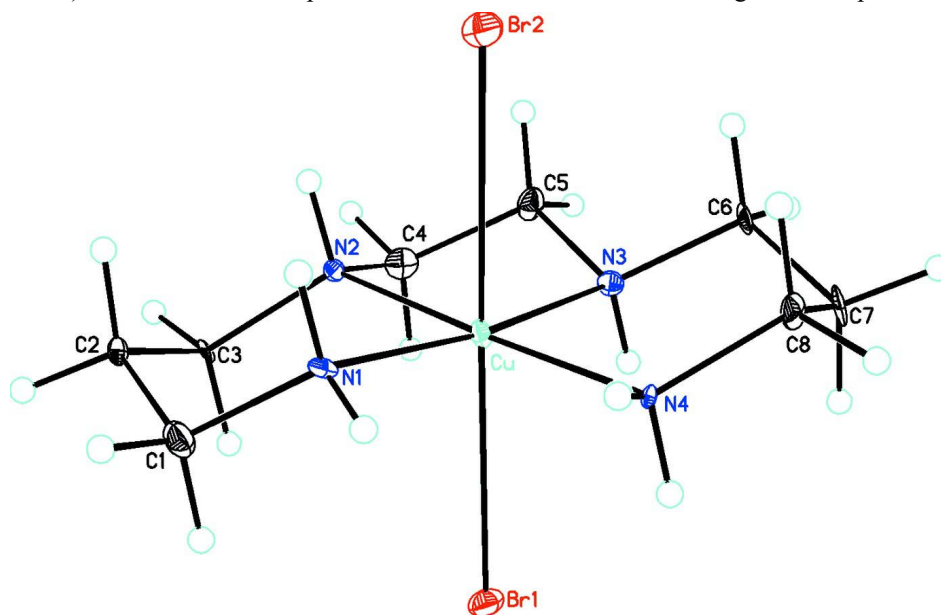
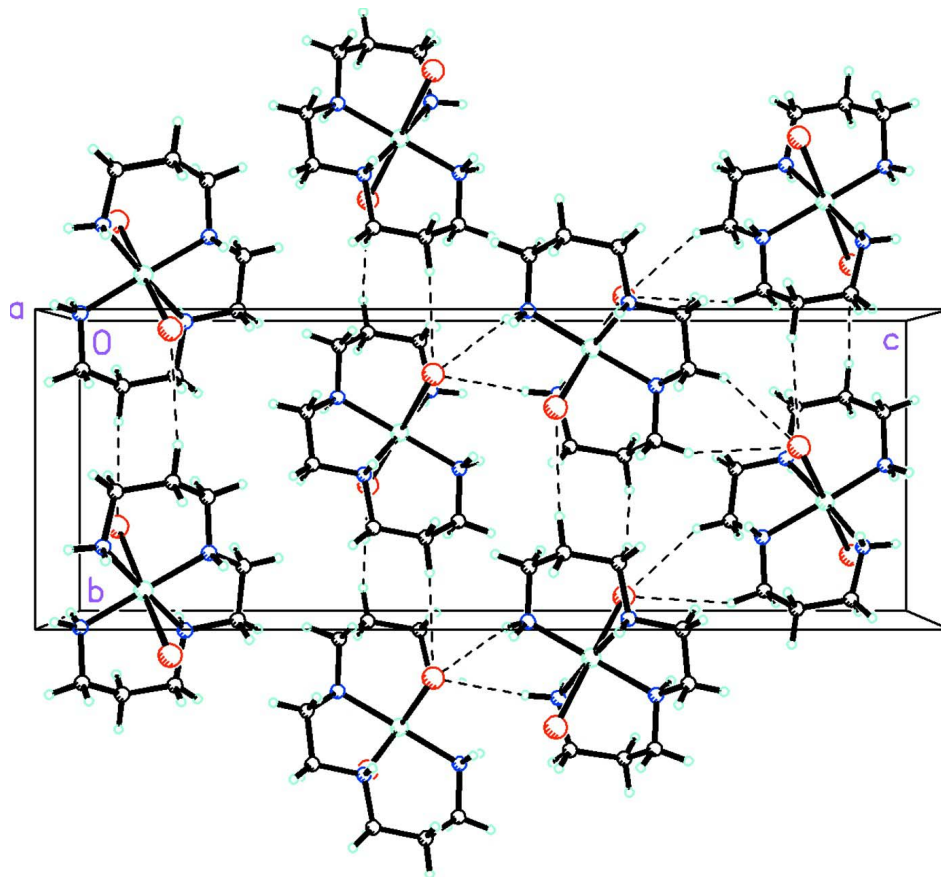


Figure 1

An ORTEP drawing of the title complex showing atom labeling. Thermal ellipsoids are drawn at the 30% probability level.

**Figure 2**

The molecular packing for the title compound viewed down the *a* axis. Hydrogen bonds are shown by dashed lines.

Dibromido(4,7-diazadecane-1,10-diamine)copper(II)

Crystal data

[CuBr₂(C₈H₂₂N₄)]

M_r = 397.66

Orthorhombic, *P*2₁2₁2₁

Hall symbol: P 2ac 2ab

a = 6.9666 (4) Å

b = 8.4146 (6) Å

c = 24.0261 (15) Å

V = 1408.45 (15) Å³

Z = 4

F(000) = 788

D_x = 1.875 Mg m⁻³

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 4805 reflections

θ = 4.6–32.8°

μ = 7.20 mm⁻¹

T = 110 K

Prism, dark blue

0.47 × 0.31 × 0.22 mm

Data collection

Goniometer Xcalibur, detector Ruby (Gemini Mo)

diffractometer

Radiation source: Enhance (Mo) X-ray Source

Graphite monochromator

Detector resolution: 10.5081 pixels mm⁻¹

ω scans

Absorption correction: analytical

(*CrysAlis PRO*; Oxford Diffraction, 2007)

T_{min} = 0.157, *T_{max}* = 0.282

9561 measured reflections

2758 independent reflections

2262 reflections with *I* > 2σ(*I*)

R_{int} = 0.072

θ_{max} = 32.8°, θ_{min} = 4.6°

h = -10→9

k = -12→11

l = -36→35

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.071$
 $wR(F^2) = 0.183$
 $S = 1.07$
 2758 reflections
 136 parameters
 24 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.103P)^2 + 8.8289P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 2.51 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -1.98 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Cu	0.84387 (19)	0.88632 (13)	0.11140 (4)	0.0099 (2)
Br1	1.15943 (15)	1.06828 (12)	0.15326 (4)	0.0187 (2)
Br2	0.52369 (16)	0.69368 (14)	0.06892 (5)	0.0225 (3)
N1	0.8009 (12)	1.0062 (10)	0.0394 (3)	0.0132 (16)
H1C	0.9013	0.9823	0.0160	0.016*
H1D	0.6914	0.9661	0.0233	0.016*
N2	0.6456 (12)	1.0252 (8)	0.1509 (3)	0.0106 (13)
H2C	0.5256	0.9901	0.1392	0.013*
N3	0.8538 (13)	0.7653 (9)	0.1852 (3)	0.0125 (14)
H3C	0.9601	0.8035	0.2044	0.015*
N4	1.0360 (12)	0.7415 (9)	0.0756 (3)	0.0110 (14)
H4C	1.0215	0.7508	0.0376	0.013*
H4D	1.1562	0.7796	0.0841	0.013*
C1	0.7815 (17)	1.1841 (12)	0.0409 (4)	0.0178 (19)
H1A	0.9031	1.2316	0.0542	0.021*
H1B	0.7571	1.2240	0.0028	0.021*
C2	0.6201 (14)	1.2343 (11)	0.0785 (4)	0.0143 (18)
H2A	0.5018	1.1786	0.0666	0.017*
H2B	0.5983	1.3496	0.0734	0.017*
C3	0.6518 (15)	1.2020 (11)	0.1399 (4)	0.0123 (15)
H3A	0.7780	1.2451	0.1514	0.015*
H3B	0.5511	1.2559	0.1620	0.015*
C4	0.6583 (17)	0.9909 (12)	0.2110 (4)	0.0160 (17)
H4A	0.5407	1.0279	0.2302	0.019*
H4B	0.7700	1.0467	0.2274	0.019*

C5	0.6801 (16)	0.8160 (13)	0.2179 (4)	0.0179 (19)
H5A	0.5644	0.7606	0.2039	0.021*
H5B	0.6970	0.7891	0.2577	0.021*
C6	0.8734 (14)	0.5888 (11)	0.1836 (4)	0.0138 (18)
H6A	0.7584	0.5421	0.1658	0.017*
H6B	0.8813	0.5473	0.2221	0.017*
C7	1.0503 (15)	0.5394 (11)	0.1517 (4)	0.0158 (18)
H7A	1.0739	0.4248	0.1581	0.019*
H7B	1.1623	0.5986	0.1663	0.019*
C8	1.0339 (15)	0.5691 (11)	0.0892 (4)	0.0142 (17)
H8A	1.1419	0.5161	0.0700	0.017*
H8B	0.9130	0.5216	0.0754	0.017*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cu	0.0119 (5)	0.0064 (4)	0.0113 (4)	0.0022 (4)	0.0026 (4)	0.0007 (4)
Br1	0.0107 (4)	0.0185 (5)	0.0270 (5)	-0.0016 (4)	-0.0007 (4)	-0.0062 (4)
Br2	0.0169 (5)	0.0213 (5)	0.0293 (5)	-0.0013 (4)	-0.0023 (4)	-0.0026 (4)
N1	0.011 (4)	0.011 (4)	0.017 (4)	-0.004 (3)	-0.001 (3)	0.002 (3)
N2	0.010 (3)	0.005 (3)	0.017 (3)	0.001 (3)	-0.001 (3)	-0.001 (3)
N3	0.011 (3)	0.012 (3)	0.015 (3)	-0.001 (3)	0.003 (3)	0.002 (3)
N4	0.008 (3)	0.009 (3)	0.016 (3)	0.002 (2)	0.002 (2)	0.000 (2)
C1	0.023 (5)	0.012 (4)	0.018 (4)	-0.002 (4)	0.003 (4)	0.005 (4)
C2	0.012 (4)	0.005 (3)	0.026 (5)	0.004 (3)	0.001 (3)	0.002 (3)
C3	0.012 (3)	0.006 (3)	0.019 (3)	0.001 (3)	0.001 (3)	0.002 (3)
C4	0.017 (4)	0.021 (4)	0.010 (3)	0.002 (4)	0.004 (4)	-0.002 (3)
C5	0.019 (5)	0.020 (4)	0.015 (4)	0.006 (4)	0.006 (4)	0.003 (4)
C6	0.016 (4)	0.009 (3)	0.017 (3)	0.003 (3)	0.002 (3)	0.004 (3)
C7	0.020 (4)	0.009 (3)	0.018 (3)	0.005 (3)	0.001 (3)	0.003 (3)
C8	0.016 (4)	0.009 (4)	0.017 (4)	0.004 (4)	0.004 (3)	-0.003 (3)

Geometric parameters (Å, °)

Cu—N4	2.005 (8)	C1—H1B	0.9900
Cu—N1	2.025 (8)	C2—C3	1.517 (13)
Cu—N2	2.043 (8)	C2—H2A	0.9900
Cu—N3	2.046 (8)	C2—H2B	0.9900
Cu—Br1	2.8616 (17)	C3—H3A	0.9900
Cu—Br2	2.9402 (17)	C3—H3B	0.9900
N1—C1	1.503 (13)	C4—C5	1.489 (15)
N1—H1C	0.9200	C4—H4A	0.9900
N1—H1D	0.9200	C4—H4B	0.9900
N2—C4	1.476 (12)	C5—H5A	0.9900
N2—C3	1.512 (11)	C5—H5B	0.9900
N2—H2C	0.9300	C6—C7	1.510 (14)
N3—C6	1.492 (11)	C6—H6A	0.9900
N3—C5	1.504 (13)	C6—H6B	0.9900

N3—H3C	0.9300	C7—C8	1.526 (14)
N4—C8	1.487 (12)	C7—H7A	0.9900
N4—H4C	0.9200	C7—H7B	0.9900
N4—H4D	0.9200	C8—H8A	0.9900
C1—C2	1.503 (14)	C8—H8B	0.9900
C1—H1A	0.9900		
N4—Cu—N1	92.0 (3)	H1A—C1—H1B	108.0
N4—Cu—N2	177.1 (3)	C1—C2—C3	115.2 (8)
N1—Cu—N2	90.7 (3)	C1—C2—H2A	108.5
N4—Cu—N3	92.7 (3)	C3—C2—H2A	108.5
N1—Cu—N3	173.4 (4)	C1—C2—H2B	108.5
N2—Cu—N3	84.6 (3)	C3—C2—H2B	108.5
N4—Cu—Br1	87.9 (2)	H2A—C2—H2B	107.5
N1—Cu—Br1	98.5 (2)	N2—C3—C2	110.0 (7)
N2—Cu—Br1	92.9 (2)	N2—C3—H3A	109.7
N3—Cu—Br1	86.3 (3)	C2—C3—H3A	109.7
N4—Cu—Br2	91.3 (2)	N2—C3—H3B	109.7
N1—Cu—Br2	82.3 (2)	C2—C3—H3B	109.7
N2—Cu—Br2	87.9 (2)	H3A—C3—H3B	108.2
N3—Cu—Br2	93.0 (3)	N2—C4—C5	107.9 (8)
Br1—Cu—Br2	178.89 (6)	N2—C4—H4A	110.1
C1—N1—Cu	119.3 (7)	C5—C4—H4A	110.1
C1—N1—H1C	107.5	N2—C4—H4B	110.1
Cu—N1—H1C	107.5	C5—C4—H4B	110.1
C1—N1—H1D	107.5	H4A—C4—H4B	108.4
Cu—N1—H1D	107.5	C4—C5—N3	107.7 (9)
H1C—N1—H1D	107.0	C4—C5—H5A	110.2
C4—N2—C3	111.2 (7)	N3—C5—H5A	110.2
C4—N2—Cu	107.6 (6)	C4—C5—H5B	110.2
C3—N2—Cu	117.6 (6)	N3—C5—H5B	110.2
C4—N2—H2C	106.6	H5A—C5—H5B	108.5
C3—N2—H2C	106.6	N3—C6—C7	111.2 (8)
Cu—N2—H2C	106.6	N3—C6—H6A	109.4
C6—N3—C5	111.7 (8)	C7—C6—H6A	109.4
C6—N3—Cu	118.5 (6)	N3—C6—H6B	109.4
C5—N3—Cu	106.5 (6)	C7—C6—H6B	109.4
C6—N3—H3C	106.5	H6A—C6—H6B	108.0
C5—N3—H3C	106.5	C6—C7—C8	113.1 (8)
Cu—N3—H3C	106.5	C6—C7—H7A	109.0
C8—N4—Cu	119.4 (6)	C8—C7—H7A	109.0
C8—N4—H4C	107.5	C6—C7—H7B	109.0
Cu—N4—H4C	107.5	C8—C7—H7B	109.0
C8—N4—H4D	107.5	H7A—C7—H7B	107.8
Cu—N4—H4D	107.5	N4—C8—C7	112.1 (8)
H4C—N4—H4D	107.0	N4—C8—H8A	109.2
C2—C1—N1	111.2 (8)	C7—C8—H8A	109.2
C2—C1—H1A	109.4	N4—C8—H8B	109.2

N1—C1—H1A	109.4	C7—C8—H8B	109.2
C2—C1—H1B	109.4	H8A—C8—H8B	107.9
N1—C1—H1B	109.4		
N4—Cu—N1—C1	-139.1 (8)	N1—Cu—N4—C8	-138.9 (7)
N2—Cu—N1—C1	42.1 (8)	N3—Cu—N4—C8	36.5 (7)
Br1—Cu—N1—C1	-50.9 (8)	Br1—Cu—N4—C8	122.7 (7)
Br2—Cu—N1—C1	129.9 (8)	Br2—Cu—N4—C8	-56.6 (7)
N1—Cu—N2—C4	-170.1 (6)	Cu—N1—C1—C2	-57.3 (11)
N3—Cu—N2—C4	14.4 (6)	N1—C1—C2—C3	67.5 (11)
Br1—Cu—N2—C4	-71.6 (6)	C4—N2—C3—C2	-174.8 (8)
Br2—Cu—N2—C4	107.6 (6)	Cu—N2—C3—C2	60.6 (10)
N1—Cu—N2—C3	-43.7 (7)	C1—C2—C3—N2	-69.7 (11)
N3—Cu—N2—C3	140.8 (7)	C3—N2—C4—C5	-171.9 (9)
Br1—Cu—N2—C3	54.8 (6)	Cu—N2—C4—C5	-41.8 (10)
Br2—Cu—N2—C3	-126.0 (6)	N2—C4—C5—N3	56.4 (11)
N4—Cu—N3—C6	-37.2 (8)	C6—N3—C5—C4	-172.8 (8)
N2—Cu—N3—C6	141.9 (8)	Cu—N3—C5—C4	-42.0 (9)
Br1—Cu—N3—C6	-124.9 (7)	C5—N3—C6—C7	-178.8 (8)
Br2—Cu—N3—C6	54.3 (7)	Cu—N3—C6—C7	56.9 (10)
N4—Cu—N3—C5	-164.0 (7)	N3—C6—C7—C8	-70.4 (11)
N2—Cu—N3—C5	15.1 (6)	Cu—N4—C8—C7	-55.5 (10)
Br1—Cu—N3—C5	108.3 (6)	C6—C7—C8—N4	69.7 (11)
Br2—Cu—N3—C5	-72.5 (6)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1C...Br2 ⁱ	0.92	2.66	3.466 (9)	147
N1—H1D...Br2	0.92	2.80	3.339 (8)	119
N2—H2C...Br1 ⁱⁱ	0.93	2.66	3.407 (8)	138
N2—H2C...Br2	0.93	3.01	3.519 (7)	116
N3—H3C...Br1	0.93	2.90	3.409 (8)	116
N4—H4C...Br2 ⁱ	0.92	2.60	3.515 (8)	171
N4—H4D...Br2 ⁱⁱⁱ	0.92	2.69	3.425 (8)	138
N4—H4D...Br1	0.92	2.94	3.433 (8)	115

Symmetry codes: (i) $x+1/2, -y+3/2, -z$; (ii) $x-1, y, z$; (iii) $x+1, y, z$.