



Crystal structure of bis(2-[(3-bromopropyl)imino]methyl]phenolato- κ^2N,O)-copper(II)

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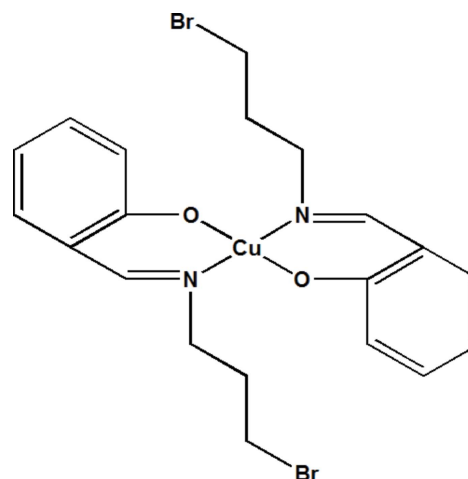
In the title compound, $[\text{Cu}(\text{C}_{10}\text{H}_{11}\text{BrNO})_2]$, the asymmetric unit consists of one-half of the molecule, the other half being generated by an inversion centre. Hence the Cu^{II} cation is symmetrically coordinated by two bidentate Schiff base anions in a slightly distorted square-planar environment with $\text{Cu}-\text{O}$ and $\text{Cu}-\text{N}$ bond lengths of 1.8786 (19) and 2.009 (2) Å, respectively. In the crystal, individual molecules are packed in alternating zigzag layers parallel to (001). Weak $\text{C}-\text{H}\cdots\pi$ interactions exist between the molecules.

Keywords: crystal structure; copper(II) complex; $\text{C}-\text{H}\cdots\pi$ interactions.

CCDC reference: 1044698

1. Related literature

For synthesis and applications of similar complexes derived from salicylaldehyde, see: Ghelenji *et al.* (2011); Kia *et al.* (2010); Zhang *et al.* (2013). For the importance of copper in biological systems, see: Siegel (1973); Mohan *et al.* (1998). For isotopic structures, see: Floyd *et al.* (2005); Ourari *et al.* (2015).



2. Experimental

2.1. Crystal data

$[\text{Cu}(\text{C}_{10}\text{H}_{11}\text{BrNO})_2]$
 $M_r = 545.75$
Monoclinic, $P2_1/n$
 $a = 10.6478$ (4) Å
 $b = 7.1990$ (3) Å
 $c = 13.9283$ (5) Å
 $\beta = 104.900$ (2)°
 $V = 1031.75$ (7) Å³
 $Z = 2$
Mo $K\alpha$ radiation
 $\mu = 4.95$ mm⁻¹
 $T = 295$ K
 $0.19 \times 0.18 \times 0.15$ mm

2.2. Data collection

Bruker APEXII diffractometer
Absorption correction: multi-scan
(SADABS; Bruker, 2011)
 $T_{\text{min}} = 0.677$, $T_{\text{max}} = 0.796$
8212 measured reflections
2594 independent reflections
2088 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$

2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.089$
 $S = 1.04$
2594 reflections
124 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.62$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.58$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the C5–C10 ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C1}-\text{H1A}\cdots\text{Cg1}^{\text{i}}$	0.97	2.74	3.645 (4)	155
$\text{C4}-\text{H4}\cdots\text{Cg1}^{\text{ii}}$	0.93	2.90	3.805 (3)	164

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

Data collection: APEX2 (Bruker, 2011); cell refinement: SAINT (Bruker, 2011); data reduction: SAINT; program(s) used to solve structure: SIR2002 (Burla *et al.*, 2005); program(s) used to refine structure: SHELXL97 (Sheldrick, 2015); molecular graphics: ORTEP-3 for Windows (Farrugia, 2012) and DIAMOND (Brandenburg, 2001); software used to prepare material for publication: WinGX (Farrugia, 2012).

Acknowledgements

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Supporting information for this paper is available from the IUCr electronic archives (Reference: WM5116).

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supporting information

Acta Cryst. (2015). E71, m33–m34 [doi:10.1107/S2056989015001309]

Crystal structure of bis(2-[(3-bromopropyl)imino]methyl}phenolato- κ^2N,O)copper(II)

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S1. Experimental

Ligand (HL) synthesis: 331.5 mg (1.5 mmol) of 2-bromopropyl ammonium hydrobromide were dissolved in absolute ethanol (15 ml). First, 756 mg (1.5 mmol; excess of 5%) and then 183 mg salicylaldehyde, each dissolved in 10 ml of absolute ethanol, were added and the resulting solution was refluxed under nitrogen atmosphere for 2 h at 333 K. The solvent was removed under reduced pressure and 15 ml of dichloromethane were added to the residue obtained. The mixture was stirred for 15 min, filtered and the solvent evaporated, resulting in a yellow viscous oil (yield: 82%).

Synthesis of the copper complex (I): 215 mg ligand HL (1 mmol) were placed in 10 ml of absolute ethanol. 99.8 mg of copper acetate monohydrate (0.5 mmol), dissolved in 5 ml of absolute ethanol, were added to this solution. The content of the flask was refluxed under stirring and nitrogen atmosphere for 2 h at 333 K. The precipitate obtained was filtered, washed with ethanol and then dried in an oven at moderate temperature (yield 70%; m. p. 393 K). Suitable single crystals were obtained from acetone solution by slow evaporation yielding green single crystals.

S2. Refinement

H atoms were localized in Fourier maps but introduced in calculated positions and treated as riding on their parent atom, with C—H = 0.97 Å (methylene) or 0.93 Å (aromatic) and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}$. Reflection $\bar{1}01$ was obstructed from the beam stop and was omitted from the refinement.

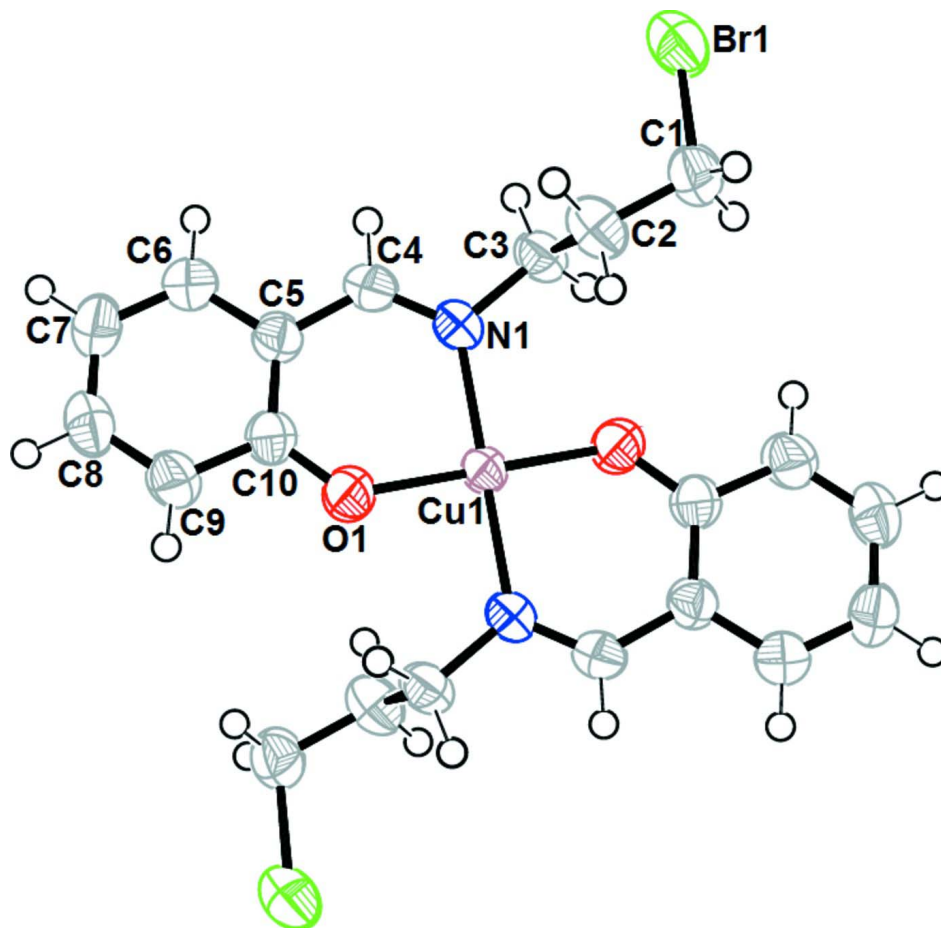
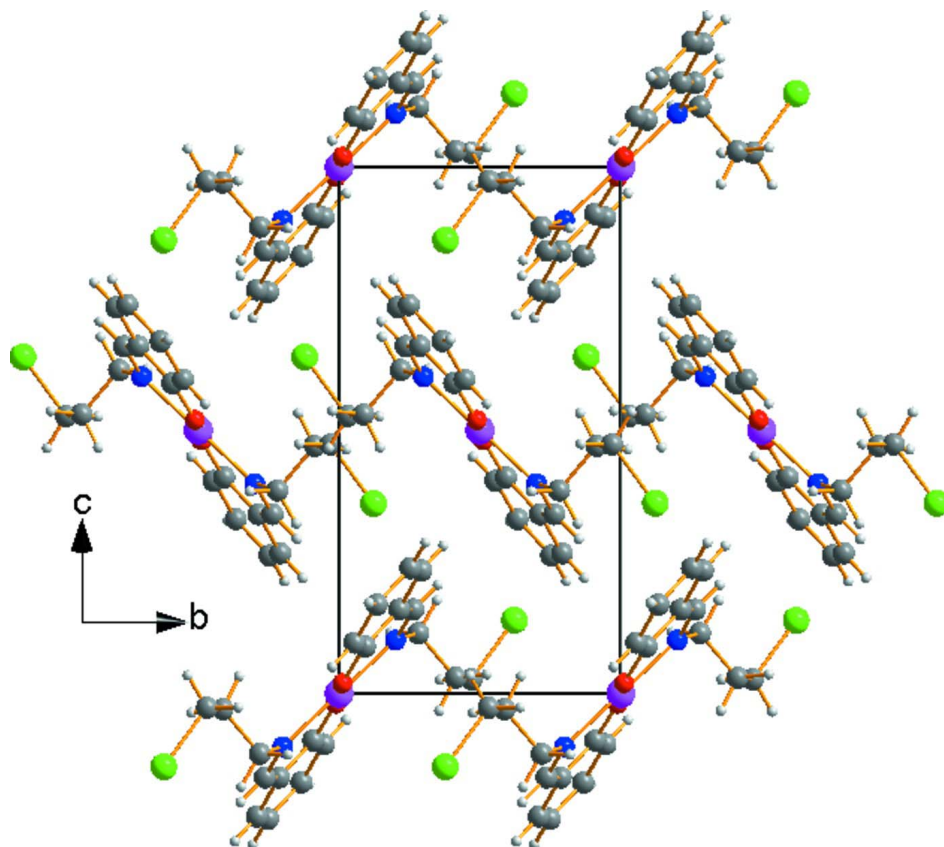
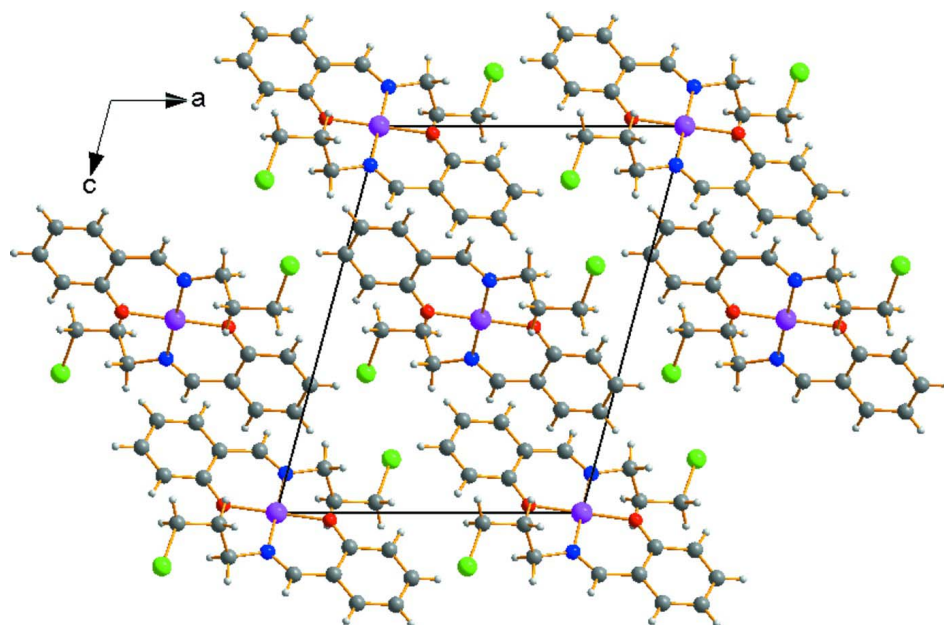


Figure 1

The molecular structure of the title compound with the atomic labelling scheme. Displacement are drawn at the 50% probability level. Non-labelled atoms are generated by symmetry code $-x+2, -y, -z+2$.

**Figure 2**

Formation of alternating zigzag layers parallel to (001).

**Figure 3**

A view of the layers along [010].

Bis(2-[(3-bromopropyl)imino]methyl)phenolato- κ^2N,O)copper(II)*Crystal data*

[Cu(C₁₀H₁₁BrNO)₂]
M_r = 545.75
 Monoclinic, *P*2₁/*n*
a = 10.6478 (4) Å
b = 7.1990 (3) Å
c = 13.9283 (5) Å
 β = 104.900 (2)°
V = 1031.75 (7) Å³
Z = 2

F(000) = 542.0
D_x = 1.757 Mg m⁻³
 Mo *K* α radiation, λ = 0.71073 Å
 Cell parameters from 3645 reflections
 θ = 3.2–27.7°
 μ = 4.95 mm⁻¹
T = 295 K
 Prism, green
 0.19 × 0.18 × 0.15 mm

Data collection

Bruker APEXII
 diffractometer
 Graphite monochromator
 CCD rotation images, thin slices scans
 Absorption correction: multi-scan
 (SADABS; Bruker, 2011)
T_{min} = 0.677, *T_{max}* = 0.796
 8212 measured reflections

2594 independent reflections
 2088 reflections with *I* > 2σ(*I*)
R_{int} = 0.020
 θ_{\max} = 28.5°, θ_{\min} = 2.8°
h = -12→14
k = -9→6
l = -18→18

Refinement

Refinement on *F*²
 Least-squares matrix: full
R[*F*² > 2σ(*F*²)] = 0.032
wR(*F*²) = 0.089
S = 1.04
 2594 reflections
 124 parameters
 0 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
 $w = 1/[\sigma^2(F_o^2) + (0.0419P)^2 + 0.8205P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.62 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.58 \text{ e \AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'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 > σ(*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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{iso} [*] / <i>U</i> _{eq}
Cu1	1	0	1	0.03472 (12)
Br1	1.32859 (3)	-0.62235 (6)	0.86101 (3)	0.06803 (14)
O1	0.81891 (18)	-0.0114 (3)	0.98050 (16)	0.0559 (6)
N1	0.99577 (19)	-0.2004 (3)	0.89887 (15)	0.0350 (4)
C5	0.7594 (2)	-0.2139 (3)	0.84019 (19)	0.0368 (5)

C8	0.5014 (3)	-0.1335 (4)	0.8310 (2)	0.0476 (7)
H8	0.4152	-0.1059	0.8282	0.057*
C4	0.8909 (2)	-0.2635 (4)	0.84046 (19)	0.0382 (5)
H4	0.9006	-0.3503	0.7934	0.046*
C9	0.5984 (3)	-0.0546 (4)	0.9037 (2)	0.0468 (6)
H9	0.5767	0.0251	0.9494	0.056*
C6	0.6576 (3)	-0.2913 (4)	0.7663 (2)	0.0492 (7)
H6	0.677	-0.3702	0.7192	0.059*
C3	1.1185 (2)	-0.2808 (4)	0.88563 (19)	0.0394 (5)
H3A	1.1795	-0.1817	0.8837	0.047*
H3B	1.1011	-0.3469	0.8229	0.047*
C10	0.7306 (2)	-0.0917 (4)	0.91054 (19)	0.0392 (5)
C2	1.1783 (3)	-0.4136 (4)	0.9701 (2)	0.0471 (6)
H2A	1.1776	-0.3551	1.0326	0.056*
H2B	1.1249	-0.5245	0.9633	0.056*
C7	0.5294 (3)	-0.2529 (5)	0.7621 (2)	0.0522 (7)
H7	0.4628	-0.3068	0.7134	0.063*
C1	1.3158 (3)	-0.4696 (4)	0.9730 (2)	0.0493 (7)
H1A	1.3521	-0.5375	1.034	0.059*
H1B	1.3676	-0.3584	0.9742	0.059*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cu1	0.0302 (2)	0.0403 (2)	0.0355 (2)	-0.00058 (17)	0.01176 (16)	-0.00451 (17)
Br1	0.0490 (2)	0.0798 (3)	0.0742 (2)	0.01763 (16)	0.01380 (16)	-0.01284 (18)
O1	0.0347 (9)	0.0754 (15)	0.0616 (12)	-0.0084 (9)	0.0195 (9)	-0.0318 (11)
N1	0.0324 (10)	0.0359 (11)	0.0380 (10)	0.0046 (8)	0.0115 (8)	0.0000 (8)
C5	0.0342 (12)	0.0326 (12)	0.0428 (13)	0.0009 (10)	0.0083 (10)	0.0000 (10)
C8	0.0324 (12)	0.0529 (17)	0.0582 (16)	0.0005 (12)	0.0129 (12)	0.0067 (13)
C4	0.0403 (12)	0.0333 (12)	0.0416 (13)	0.0047 (10)	0.0118 (10)	-0.0034 (10)
C9	0.0382 (13)	0.0504 (15)	0.0570 (16)	-0.0021 (12)	0.0213 (12)	-0.0063 (13)
C6	0.0428 (14)	0.0445 (15)	0.0576 (17)	0.0020 (12)	0.0082 (12)	-0.0117 (13)
C3	0.0372 (12)	0.0427 (14)	0.0405 (13)	0.0073 (11)	0.0141 (10)	-0.0014 (11)
C10	0.0344 (12)	0.0411 (14)	0.0443 (13)	-0.0041 (10)	0.0140 (10)	-0.0024 (11)
C2	0.0452 (15)	0.0470 (16)	0.0515 (15)	0.0099 (12)	0.0172 (12)	0.0093 (13)
C7	0.0359 (13)	0.0553 (17)	0.0590 (17)	-0.0034 (13)	0.0008 (12)	-0.0053 (14)
C1	0.0411 (14)	0.0502 (17)	0.0534 (16)	0.0057 (12)	0.0062 (12)	0.0030 (13)

Geometric parameters (Å, °)

Cu1—O1 ⁱ	1.8786 (19)	C4—H4	0.93
Cu1—O1	1.8786 (19)	C9—C10	1.412 (4)
Cu1—N1	2.009 (2)	C9—H9	0.93
Cu1—N1 ⁱ	2.009 (2)	C6—C7	1.380 (4)
Br1—C1	1.942 (3)	C6—H6	0.93
O1—C10	1.302 (3)	C3—C2	1.522 (4)
N1—C4	1.284 (3)	C3—H3A	0.97

N1—C3	1.484 (3)	C3—H3B	0.97
C5—C6	1.404 (4)	C2—C1	1.509 (4)
C5—C10	1.408 (4)	C2—H2A	0.97
C5—C4	1.444 (3)	C2—H2B	0.97
C8—C9	1.369 (4)	C7—H7	0.93
C8—C7	1.377 (4)	C1—H1A	0.97
C8—H8	0.93	C1—H1B	0.97
O1 ⁱ —Cu1—O1	180.0000 (10)	N1—C3—C2	110.9 (2)
O1 ⁱ —Cu1—N1	88.28 (8)	N1—C3—H3A	109.5
O1—Cu1—N1	91.72 (8)	C2—C3—H3A	109.5
O1 ⁱ —Cu1—N1 ⁱ	91.72 (8)	N1—C3—H3B	109.5
O1—Cu1—N1 ⁱ	88.28 (8)	C2—C3—H3B	109.5
N1—Cu1—N1 ⁱ	180.0000 (10)	H3A—C3—H3B	108.1
C10—O1—Cu1	130.10 (17)	O1—C10—C5	123.6 (2)
C4—N1—C3	115.7 (2)	O1—C10—C9	118.8 (2)
C4—N1—Cu1	123.89 (16)	C5—C10—C9	117.6 (2)
C3—N1—Cu1	120.39 (16)	C1—C2—C3	113.5 (2)
C6—C5—C10	119.5 (2)	C1—C2—H2A	108.9
C6—C5—C4	118.0 (2)	C3—C2—H2A	108.9
C10—C5—C4	122.5 (2)	C1—C2—H2B	108.9
C9—C8—C7	121.1 (3)	C3—C2—H2B	108.9
C9—C8—H8	119.4	H2A—C2—H2B	107.7
C7—C8—H8	119.4	C8—C7—C6	118.9 (3)
N1—C4—C5	126.8 (2)	C8—C7—H7	120.5
N1—C4—H4	116.6	C6—C7—H7	120.5
C5—C4—H4	116.6	C2—C1—Br1	113.4 (2)
C8—C9—C10	121.4 (3)	C2—C1—H1A	108.9
C8—C9—H9	119.3	Br1—C1—H1A	108.9
C10—C9—H9	119.3	C2—C1—H1B	108.9
C7—C6—C5	121.4 (3)	Br1—C1—H1B	108.9
C7—C6—H6	119.3	H1A—C1—H1B	107.7
C5—C6—H6	119.3		
N1—Cu1—O1—C10	-13.1 (3)	Cu1—N1—C3—C2	75.6 (3)
N1 ⁱ —Cu1—O1—C10	166.9 (3)	Cu1—O1—C10—C5	9.8 (4)
O1 ⁱ —Cu1—N1—C4	-170.0 (2)	Cu1—O1—C10—C9	-169.8 (2)
O1—Cu1—N1—C4	10.0 (2)	C6—C5—C10—O1	-178.9 (3)
O1 ⁱ —Cu1—N1—C3	7.98 (19)	C4—C5—C10—O1	1.0 (4)
O1—Cu1—N1—C3	-172.02 (19)	C6—C5—C10—C9	0.8 (4)
C3—N1—C4—C5	177.6 (2)	C4—C5—C10—C9	-179.4 (3)
Cu1—N1—C4—C5	-4.3 (4)	C8—C9—C10—O1	179.3 (3)
C6—C5—C4—N1	176.5 (3)	C8—C9—C10—C5	-0.3 (4)
C10—C5—C4—N1	-3.3 (4)	N1—C3—C2—C1	-168.0 (2)
C7—C8—C9—C10	0.3 (5)	C9—C8—C7—C6	-0.7 (5)
C10—C5—C6—C7	-1.2 (4)	C5—C6—C7—C8	1.1 (5)

C4—C5—C6—C7	179.0 (3)	C3—C2—C1—Br1	-67.5 (3)
C4—N1—C3—C2	-106.3 (3)		

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

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

Cg1 is the centroid of the C5–C10 ring.

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C1—H1A \cdots Cg1 ⁱⁱ	0.97	2.74	3.645 (4)	155
C4—H4 \cdots Cg1 ⁱⁱⁱ	0.93	2.90	3.805 (3)	164

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