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Crystal structure of di- μ -chlorido-bis[chlorido- bis(1,2-dimethyl-5-nitro-1H-imidazole- κN^3)- copper(II)] acetonitrile disolvate

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1,2-Dimethyl-5-nitroimidazole (dimetridazole, dimet) is a compound that belongs to a class of nitroimidazole drugs that are effective at inhibiting the activity of certain parasites and bacteria. However, there are few reports that describe structures of compounds that feature metals complexed by dimet. Therefore, we report here that dimet reacts with CuCl₂·H₂O to yield a chloride-bridged copper(II) dimer, [Cu₂Cl₄(C₅H₇N₃O₂)₄] or [Cu(μ -Cl)Cl(dimet)₂]₂. In this molecule, the Cu^{II} ions are coordinated in an approximately trigonal-bipyramidal manner, and the molecule lies across an inversion center. The dihedral angle between the imidazole rings in the asymmetric unit is 4.28 (7) $^\circ$. Compared to metronidazole, dimetridazole lacks the hydroxyethyl group, and thus cannot form intermolecular O \cdots H hydrogen-bonding interactions. Instead, [Cu(μ -Cl)Cl(dimet)₂]₂ exhibits weak intermolecular interactions between the hydrogen atoms of C–H groups and (i) oxygen in the nitro groups, and (ii) the terminal and bridging chloride ligands. The unit cell contains four disordered acetonitrile molecules. These were modeled as providing a diffuse contribution to the overall scattering by SQUEEZE [Spek (2015). *Acta Cryst. C71*, 9–18], which identified two voids, each with a volume of 163 Å³ and a count of 46 electrons, indicative of a total of four acetonitrile molecules. These acetonitrile molecules are included in the chemical formula to give the expected calculated density and *F*(000).

1. Chemical context

1,2-Dimethyl-5-nitroimidazole, also known as dimetridazole (dimet), is structurally related to metronidazole [2-(2-methyl-5-nitro-1H-imidazol-1-yl)ethanol, MET]. Thus, both compounds contain a 2-methyl-5-nitroimidazole core and are only differentiated according to whether one of the nitrogen atoms possesses a methyl substituent (as in dimet) or a hydroxyethyl substituent (as in MET), as illustrated in Fig. 1. Both MET and dimet are used to treat microbial infections,

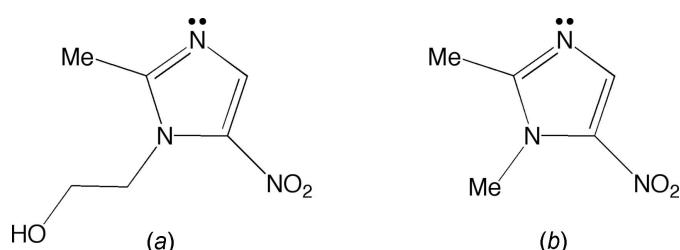
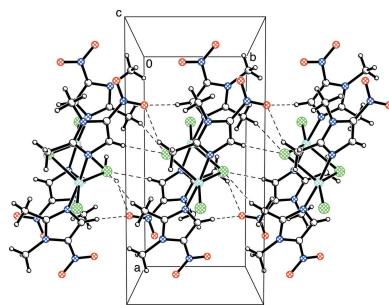
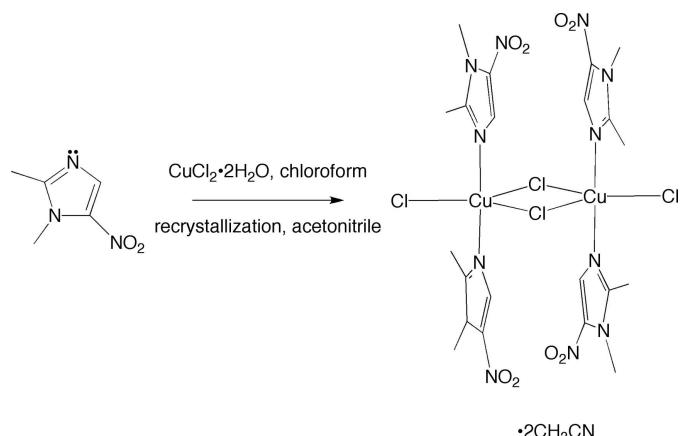


Figure 1

A comparison of the structures of (a) metronidazole (MET) and (b) dimetridazole (dimet).

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but dimet has specifically been used in animals for the treatment of, for instance, bovine trichomoniasis (McLoughlin, 1968), giardiasis in birds (Panigrahy *et al.*, 1978) and swine dysentery (Messier *et al.*, 1990). In order to control outbreaks of infection, a previous common practice was to incorporate dimet as a feed additive given, for example, to poultry and pigs (Buizer & Severijnen, 1975). However, concerns about the mutagenic properties displayed by this class of drug (Voogd *et al.*, 1974), and the fact that trace amounts can be detected in certain animal products intended for human consumption (Arias *et al.*, 2016), have led to a discontinuation of this practice (*EC bans use of dimetridazole in food animals*, 1995). Reports of structures of metal compounds involving the coordination by dimetridazole are scarce. Herein, we describe the structure of the copper compound $[\text{Cu}(\mu\text{-Cl})\text{Cl}(\text{dimet})_2]_2$, which is obtained by the reaction of dimet with $\text{CuCl}_2 \cdot 2\text{H}_2\text{O}$ (see Scheme).



2. Structural commentary

Crystals of composition $[\text{Cu}(\mu\text{-Cl})\text{Cl}(\text{dimet})_2]_2$ were obtained by addition of dimet to $\text{CuCl}_2 \cdot 2\text{H}_2\text{O}$ in chloroform, followed by recrystallization of the blue precipitate from acetonitrile. The molecular structure, as illustrated in Fig. 2, shows a centrosymmetric chlorido-bridged dimer. The coordination geometry around each copper atom is a slightly distorted trigonal-bipyramidal with two axial dimet ligands, and three chlorine ligands in the equatorial plane, two of which bridge to the adjacent copper. This structure is analogous to a previously reported copper(II) dimer containing MET, instead of dimet, $[\text{Cu}(\text{MET})_2(\mu\text{-Cl})\text{Cl}]_2$ (Barba-Behrens *et al.*, 1991), and a comparison of the two structures is shown in Fig. 3. Other recent examples of metal compounds containing MET include: $\text{Cu}(\text{MET})_2\text{Cl}_2$, $[\text{Ag}(\text{MET})_2](\text{BF}_4)$, and $[\text{H}(\text{MET})][\text{AuCl}_4]$ (Palmer *et al.*, 2015; Palmer & Upmacis, 2015; Quinlivan *et al.*, 2015).

Examination of the structure of the $[\text{Cu}(\mu\text{-Cl})\text{Cl}(\text{dimet})_2]_2$ complex demonstrates that, interestingly, the chlorine atoms bridge in an asymmetric manner, with $\text{Cu}-\text{Cl}_{\text{bridge}}$ bond lengths of 2.3811 (3) and 2.6024 (3) Å, both of which are longer than the terminal $\text{Cu}-\text{Cl}$ bond length of 2.2822 (3) Å. Of note, these features are also observed for the MET analog,

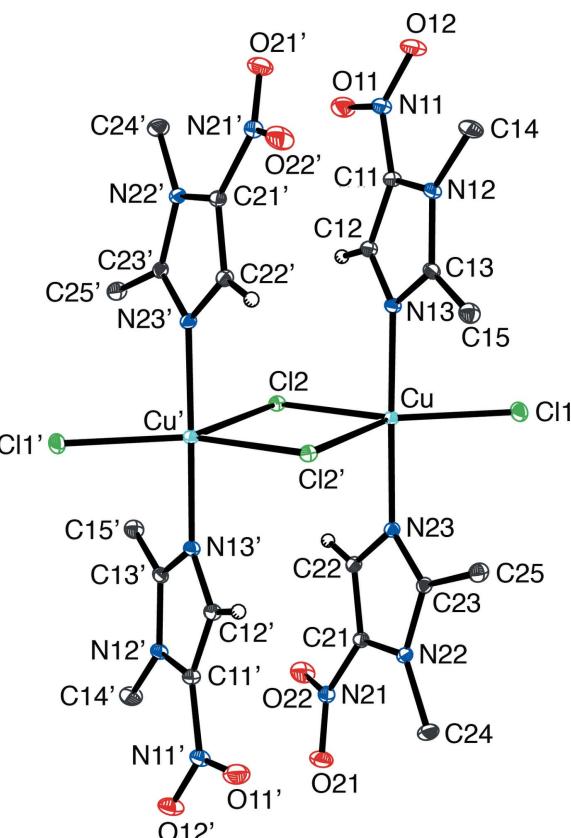


Figure 2

The molecular structure of $[\text{Cu}(\mu\text{-Cl})\text{Cl}(\text{dimet})_2]_2$, with displacement ellipsoids depicted at the 30% probability level. H atoms associated with methyl groups are not shown [symmetry code ('): $-x, -y + 1, -z + 1$].

$[\text{Cu}(\text{MET})_2(\mu\text{-Cl})\text{Cl}]_2$, which possesses bridging $\text{Cu}-\text{Cl}$ distances of 2.418 (1) and 2.619 (1) Å, and a terminal bond length of 2.297 (2) Å (Barba-Behrens *et al.*, 1991). Furthermore, the $\text{Cu}-\text{N}$ bond lengths [2.0009 (10) and 1.9914 (9) Å] are also similar to the $\text{Cu}-\text{N}$ bond lengths reported for the MET analog [2.002 (4) and 1.993 (4) Å]. In terms of the bond angles, the $\text{N}-\text{Cu}-\text{Cl}_{\text{term}}$ and $\text{N}-\text{Cu}-\text{Cl}_{\text{bridge}}$ angles are all close to 90° [ranging from 88.95 (3)° for $\text{N}13-\text{Cu}-\text{Cl}1$ to

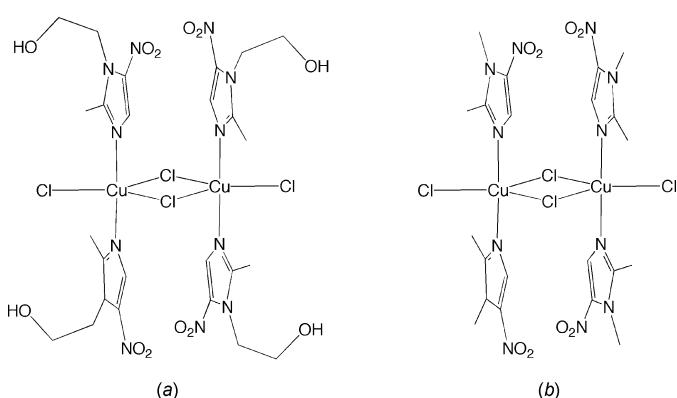


Figure 3

A comparison of the structures of the dinuclear Cu complexes which are derived from (a) metronidazole (MET) and (b) dimetridazole (dimet).

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C25—H25C \cdots O22 ⁱ	0.98	2.39	3.2804 (17)	150
C15—H15B \cdots Cl2 ⁱ	0.98	2.73	3.6737 (13)	163
C12—H12A \cdots O22 ⁱⁱ	0.95	2.51	3.4029 (16)	156
C22—H22A \cdots Cl2 ⁱⁱ	0.95	2.75	3.6828 (12)	167
C24—H24C \cdots Cl1 ⁱⁱⁱ	0.98	2.84	3.7555 (13)	156

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

91.85 (3)° for N23—Cu—Cl2], with the exception of N23—Cu—Cl2_{bridge} which is 85.42 (3) Å.

3. Supramolecular features

The crystal structure displays a number of weak intermolecular interactions between hydrogen atoms of CH groups and the more electronegative atoms on adjacent molecules, such as the oxygen atoms in the nitro groups of the dimet ligand and also the terminal and bridging chlorine atoms (see Table 1 and Fig. 4). In this regard, one of the oxygen atoms of the nitro group participates in intermolecular hydrogen-bonding interactions with CH₃ and CH groups of an adjacent molecule. For reference, intermolecular and intramolecular C—H \cdots O hydrogen bonds involving an O atom from a nitro group (or other O-containing groups) have been reported (Desiraju, 1991; Sharma & Desiraju, 1994; Forlani, 2009). As an illustration, intermolecular C—H \cdots O interactions (involving C—H motifs from an NMe₂ substituent and the O atoms of a nitro group) are reported at 2.71 (3) Å, with C \cdots O distances of 3.658 (4) and 3.725 (4) Å (Sharma & Desiraju, 1994). The results of our structure analysis are also comparable to the average values that have been reported for hydrogen-bonding interactions of (N,C)Csp³—H (2.48 and

3.47 Å) and Csp³—CH₃ (2.63 and 3.61 Å) groups with a water O atom (Steiner, 2002). For comparison, intramolecular N—H \cdots O interactions to an O atom of a nitro substituent form shorter contacts, e.g. 1.927 (15) Å for *N*-(2-nitrophenyl)benzamide (Saeed & Simpson, 2009) and 2.11 Å for 2-iodo-*N*-(2-nitrophenyl)benzamide (Wardell *et al.*, 2005), which is in accord with the reports that C—H \cdots O bonds are weaker than N—H \cdots O bonds (Desiraju, 1991).

The bridging chlorine atoms also form weak intermolecular interactions with CH₃ and CH groups of an adjacent molecule. In addition, the terminal chlorine atom participates in a hydrogen-bonding interaction with a CH₃ group of an adjacent molecule.

While C—H \cdots O interactions are widely accepted (Desiraju, 1991), C—H \cdots Cl interactions are considered more controversial, but a survey of the literature reveals that they also represent a common phenomenon (Aakeröy *et al.*, 1999). For example, hydrogen-bonding interactions of sp² (N,C)C—H with Cl[−] are reported at 2.64 Å (Kovacs & Varga, 2006). However, when Cl is bonded to a metal, the average C—H \cdots Cl—M hydrogen-bonding distance is 2.974 Å (Thallapally & Nangia, 2001).

Fig. 4 illustrates some of these intermolecular interactions. An important difference between this structure and the MET analog is that the dimet compound lacks the hydroxyethyl group, which is involved in classical intermolecular hydrogen-bonding interactions for the MET derivative (Barba-Behrens *et al.*, 1991).

4. Database survey

There is only one structurally characterized metal compound containing dimet listed in the Cambridge Database (CSD Version 5.37; Groom *et al.*, 2016), namely, a mononuclear cobalt complex, [Co(dimet)₂Cl₂], in which the cobalt(II) atom is surrounded by two dimet and two chlorido ligands in a distorted tetrahedron (Rosu *et al.*, 1997; Idešicová *et al.*, 2012). The Co—N distances are reported to be 2.228 (2) and 2.035 (4) Å (Rosu *et al.*, 1997).

5. Synthesis and crystallization

CuCl₂·H₂O (3 mg, 0.018 mmol) was added to a solution of dimet (6 mg, 0.043 mmol) in chloroform (0.7 mL), resulting in the precipitation of a blue solid over the course of 1 h at room temperature. The blue solid was isolated by decantation and crystals of [Cu(μ -Cl)Cl(dimet)₂]₂, suitable for X-ray diffraction, were obtained by slow evaporation from a solution in acetonitrile.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. Hydrogen atoms on carbon were placed in calculated positions (C—H = 0.95–1.00 Å) and included as riding contributions with isotropic displacement parameters $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{Csp}^3)$ or $1.5U_{\text{eq}}(\text{Csp}^3)$. The unit

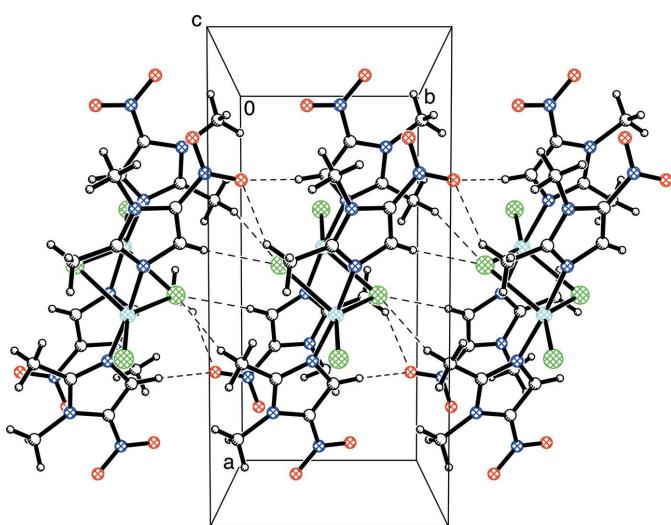


Figure 4
Weak intermolecular hydrogen-bonding interactions (shown as dashed lines) for [Cu(μ -Cl)Cl(dimet)₂]₂.

Table 2
Experimental details.

Crystal data	
Chemical formula	[Cu ₂ Cl ₄ (C ₅ H ₇ N ₃ O ₂) ₄]·2C ₂ H ₃ N
M _r	915.53
Crystal system, space group	Monoclinic, P2 ₁ /c
Temperature (K)	130
a, b, c (Å)	13.9545 (8), 6.7004 (4), 19.5031 (11)
β (°)	96.424 (1)
V (Å ³)	1812.10 (18)
Z	2
Radiation type	Mo Kα
μ (mm ⁻¹)	1.54
Crystal size (mm)	0.35 × 0.17 × 0.10
Data collection	
Diffractometer	Bruker APEXII CCD
Absorption correction	Multi-scan (SADABS; Bruker, 2010)
T _{min} , T _{max}	0.637, 0.746
No. of measured, independent and observed [I > 2σ(I)] reflections	28822, 5564, 5034
R _{int}	0.033
(sin θ/λ) _{max} (Å ⁻¹)	0.716
Refinement	
R[F ² > 2σ(F ²)], wR(F ²), S	0.024, 0.071, 1.26
No. of reflections	5564
No. of parameters	212
H-atom treatment	H-atom parameters constrained
Δρ _{max} , Δρ _{min} (e Å ⁻³)	0.51, -0.51

Computer programs: APEX2 and SAINT (Bruker, 2010), SHELXS97 and SHELXTL (Sheldrick, 2008) and SHELXL2014 (Sheldrick, 2015).

cell contains four disordered acetonitrile molecules. In view of the disorder, the acetonitrile molecules were modeled as providing a diffuse contribution to the overall scattering by SQUEEZE (Spek, 2015), which identified two voids, each with a volume of 163 Å³ and a count of 46 electrons, indicative of a total of four acetonitrile molecules.

Acknowledgements

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Acta Cryst. (2016). E72, 1633-1636 [https://doi.org/10.1107/S2056989016015413]

Crystal structure of di- μ -chlorido-bis[chloridobis(1,2-dimethyl-5-nitro-1*H*-imidazole- κN^3)copper(II)] acetonitrile disolvate

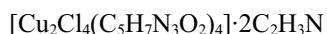
Patrick J. Quinlivan and Rita K. Upmacis

Computing details

Data collection: *APEX2* (Bruker, 2010); cell refinement: *SAINT* (Bruker, 2010); data reduction: *SAINT* (Bruker, 2010); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

Di- μ -chlorido-bis[chloridobis(1,2-dimethyl-5-nitro-1*H*-imidazole- κN^3)copper(II)] acetonitrile disolvate

Crystal data



$$M_r = 915.53$$

Monoclinic, $P2_1/c$

$$a = 13.9545 (8) \text{ \AA}$$

$$b = 6.7004 (4) \text{ \AA}$$

$$c = 19.5031 (11) \text{ \AA}$$

$$\beta = 96.424 (1)^\circ$$

$$V = 1812.10 (18) \text{ \AA}^3$$

$$Z = 2$$

$$F(000) = 932$$

$$D_x = 1.678 \text{ Mg m}^{-3}$$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 9452 reflections

$$\theta = 2.9\text{--}30.6^\circ$$

$$\mu = 1.54 \text{ mm}^{-1}$$

$$T = 130 \text{ K}$$

Block, blue

$$0.35 \times 0.17 \times 0.10 \text{ mm}$$

Data collection

Bruker APEXII CCD

 diffractometer

φ and ω scans

Absorption correction: multi-scan
 (SADABS; Bruker, 2010)

$$T_{\min} = 0.637, T_{\max} = 0.746$$

28822 measured reflections

5564 independent reflections

5034 reflections with $I > 2\sigma(I)$

$$R_{\text{int}} = 0.033$$

$$\theta_{\max} = 30.6^\circ, \theta_{\min} = 1.5^\circ$$

$$h = -19 \rightarrow 19$$

$$k = -9 \rightarrow 9$$

$$l = -27 \rightarrow 27$$

Refinement

Refinement on F^2

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.024$$

$$wR(F^2) = 0.071$$

$$S = 1.26$$

5564 reflections

212 parameters

0 restraints

Hydrogen site location: inferred from
 neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0371P)^2]$$

where $P = (F_o^2 + 2F_c^2)/3$

$$(\Delta/\sigma)_{\max} = 0.033$$

$$\Delta\rho_{\max} = 0.51 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.51 \text{ e \AA}^{-3}$$

Special details

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

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cu	0.41957 (2)	0.46806 (2)	0.42320 (2)	0.01249 (5)
Cl1	0.32308 (2)	0.45734 (5)	0.32075 (2)	0.02105 (7)
Cl2	0.46436 (2)	0.25939 (4)	0.52033 (2)	0.01295 (6)
O11	0.10021 (7)	0.39434 (18)	0.58607 (6)	0.0303 (2)
O12	0.02709 (7)	0.66914 (16)	0.55203 (5)	0.0280 (2)
O21	0.81805 (7)	0.13329 (16)	0.33794 (5)	0.0263 (2)
O22	0.72749 (7)	-0.08394 (15)	0.38274 (6)	0.0276 (2)
N11	0.09544 (8)	0.55179 (17)	0.55458 (6)	0.0209 (2)
N12	0.18478 (7)	0.77681 (16)	0.48250 (5)	0.01611 (19)
N13	0.30915 (7)	0.58276 (15)	0.46736 (5)	0.01353 (18)
N21	0.74156 (7)	0.07990 (16)	0.35776 (5)	0.0175 (2)
N22	0.65940 (7)	0.39673 (15)	0.31873 (5)	0.01363 (18)
N23	0.53211 (7)	0.37776 (14)	0.37688 (5)	0.01264 (18)
C11	0.17488 (8)	0.59978 (19)	0.51720 (6)	0.0164 (2)
C12	0.25207 (9)	0.48130 (18)	0.50794 (6)	0.0152 (2)
H12A	0.2639	0.3513	0.5264	0.018*
C13	0.26757 (8)	0.76046 (17)	0.45264 (6)	0.0146 (2)
C14	0.12067 (10)	0.9527 (2)	0.47787 (8)	0.0247 (3)
H14A	0.1480	1.0572	0.4508	0.037*
H14B	0.1145	1.0030	0.5244	0.037*
H14C	0.0569	0.9147	0.4554	0.037*
C15	0.30624 (9)	0.92050 (19)	0.41131 (7)	0.0189 (2)
H15A	0.3555	0.8650	0.3847	0.028*
H15B	0.3350	1.0252	0.4422	0.028*
H15C	0.2538	0.9774	0.3797	0.028*
C21	0.66334 (8)	0.21847 (17)	0.35462 (6)	0.0138 (2)
C22	0.58485 (8)	0.20765 (17)	0.39040 (6)	0.0135 (2)
H22A	0.5696	0.1011	0.4194	0.016*
C23	0.57767 (8)	0.48769 (17)	0.33275 (6)	0.0131 (2)
C24	0.72860 (10)	0.4744 (2)	0.27397 (7)	0.0218 (3)
H24A	0.7509	0.3653	0.2463	0.033*
H24B	0.7838	0.5330	0.3024	0.033*
H24C	0.6973	0.5768	0.2433	0.033*
C25	0.54213 (9)	0.68022 (18)	0.30190 (6)	0.0181 (2)
H25A	0.4740	0.6973	0.3085	0.027*
H25B	0.5491	0.6802	0.2525	0.027*
H25C	0.5798	0.7902	0.3244	0.027*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cu	0.01138 (8)	0.01446 (8)	0.01188 (8)	0.00362 (5)	0.00239 (5)	-0.00060 (5)
Cl1	0.01728 (14)	0.02815 (16)	0.01670 (14)	0.00367 (11)	-0.00253 (11)	-0.00714 (11)
Cl2	0.01421 (12)	0.01058 (12)	0.01442 (12)	0.00041 (9)	0.00312 (9)	0.00093 (9)
O11	0.0229 (5)	0.0378 (6)	0.0318 (6)	0.0017 (4)	0.0102 (4)	0.0092 (5)
O12	0.0167 (4)	0.0344 (6)	0.0343 (6)	0.0063 (4)	0.0090 (4)	-0.0021 (4)
O21	0.0165 (4)	0.0332 (5)	0.0304 (5)	0.0058 (4)	0.0078 (4)	0.0035 (4)
O22	0.0259 (5)	0.0198 (4)	0.0379 (6)	0.0084 (4)	0.0066 (4)	0.0071 (4)
N11	0.0131 (5)	0.0292 (6)	0.0208 (5)	0.0021 (4)	0.0034 (4)	-0.0032 (4)
N12	0.0126 (4)	0.0188 (5)	0.0171 (5)	0.0049 (4)	0.0020 (4)	-0.0022 (4)
N13	0.0116 (4)	0.0151 (4)	0.0140 (4)	0.0015 (3)	0.0020 (3)	-0.0017 (3)
N21	0.0157 (5)	0.0211 (5)	0.0157 (5)	0.0038 (4)	0.0018 (4)	-0.0016 (4)
N22	0.0139 (4)	0.0151 (4)	0.0122 (4)	0.0009 (4)	0.0029 (3)	0.0007 (3)
N23	0.0135 (4)	0.0125 (4)	0.0122 (4)	0.0023 (3)	0.0026 (3)	0.0003 (3)
C11	0.0131 (5)	0.0211 (6)	0.0155 (5)	0.0017 (4)	0.0035 (4)	-0.0016 (4)
C12	0.0132 (5)	0.0173 (5)	0.0153 (5)	0.0001 (4)	0.0022 (4)	-0.0009 (4)
C13	0.0116 (5)	0.0171 (5)	0.0148 (5)	0.0015 (4)	0.0001 (4)	-0.0034 (4)
C14	0.0214 (6)	0.0248 (7)	0.0287 (7)	0.0140 (5)	0.0061 (5)	0.0000 (5)
C15	0.0190 (6)	0.0162 (5)	0.0217 (6)	0.0024 (4)	0.0030 (5)	0.0019 (4)
C21	0.0139 (5)	0.0143 (5)	0.0135 (5)	0.0031 (4)	0.0020 (4)	-0.0008 (4)
C22	0.0154 (5)	0.0122 (5)	0.0130 (5)	0.0023 (4)	0.0025 (4)	0.0005 (4)
C23	0.0142 (5)	0.0133 (5)	0.0118 (5)	0.0007 (4)	0.0015 (4)	-0.0009 (4)
C24	0.0190 (6)	0.0278 (7)	0.0202 (6)	0.0008 (5)	0.0087 (5)	0.0068 (5)
C25	0.0207 (6)	0.0158 (5)	0.0186 (6)	0.0031 (4)	0.0049 (4)	0.0047 (4)

Geometric parameters (\AA , $^\circ$)

Cu—N23	1.9914 (9)	N22—C24	1.4678 (16)
Cu—N13	2.0009 (10)	N23—C23	1.3455 (15)
Cu—Cl1	2.2822 (3)	N23—C22	1.3663 (14)
Cu—Cl2	2.3811 (3)	C11—C12	1.3662 (16)
Cu—Cl2 ⁱ	2.6024 (3)	C12—H12A	0.9500
Cl2—Cu ⁱ	2.6024 (3)	C13—C15	1.4792 (17)
O11—N11	1.2188 (16)	C14—H14A	0.9800
O12—N11	1.2329 (14)	C14—H14B	0.9800
O12—N11 ⁱⁱ	2.9399 (15)	C14—H14C	0.9800
O21—N21	1.2285 (14)	C15—H15A	0.9800
O22—N21	1.2258 (15)	C15—H15B	0.9800
N11—C11	1.4299 (16)	C15—H15C	0.9800
N11—O12 ⁱⁱ	2.9399 (15)	C21—C22	1.3649 (16)
N12—C13	1.3552 (15)	C22—H22A	0.9500
N12—C11	1.3802 (16)	C23—C25	1.4850 (16)
N12—C14	1.4764 (15)	C24—H24A	0.9800
N13—C13	1.3414 (15)	C24—H24B	0.9800
N13—C12	1.3653 (15)	C24—H24C	0.9800
N21—C21	1.4291 (15)	C25—H25A	0.9800

N22—C23	1.3478 (15)	C25—H25B	0.9800
N22—C21	1.3823 (15)	C25—H25C	0.9800
N23—Cu—N13	175.06 (4)	N13—C12—H12A	126.1
N23—Cu—Cl1	90.65 (3)	N13—C13—N12	110.42 (11)
N13—Cu—Cl1	88.95 (3)	N13—C13—C15	125.78 (11)
N23—Cu—Cl2	91.85 (3)	N12—C13—C15	123.78 (11)
N13—Cu—Cl2	91.68 (3)	N12—C14—H14A	109.5
Cl1—Cu—Cl2	139.002 (12)	N12—C14—H14B	109.5
N23—Cu—Cl2 ⁱ	85.42 (3)	H14A—C14—H14B	109.5
N13—Cu—Cl2 ⁱ	91.21 (3)	N12—C14—H14C	109.5
Cl1—Cu—Cl2 ⁱ	132.139 (12)	H14A—C14—H14C	109.5
Cl2—Cu—Cl2 ⁱ	88.842 (10)	H14B—C14—H14C	109.5
Cu—Cl2—Cu ⁱ	91.158 (10)	C13—C15—H15A	109.5
N11—O12—N11 ⁱⁱ	95.37 (8)	C13—C15—H15B	109.5
O11—N11—O12	124.80 (11)	H15A—C15—H15B	109.5
O11—N11—C11	116.73 (10)	C13—C15—H15C	109.5
O12—N11—C11	118.46 (11)	H15A—C15—H15C	109.5
O11—N11—O12 ⁱⁱ	85.14 (8)	H15B—C15—H15C	109.5
O12—N11—O12 ⁱⁱ	84.63 (8)	C22—C21—N22	108.42 (10)
C11—N11—O12 ⁱⁱ	100.10 (7)	C22—C21—N21	126.61 (11)
C13—N12—C11	106.15 (10)	N22—C21—N21	124.75 (10)
C13—N12—C14	125.32 (11)	C21—C22—N23	107.62 (10)
C11—N12—C14	128.53 (10)	C21—C22—H22A	126.2
C13—N13—C12	107.43 (10)	N23—C22—H22A	126.2
C13—N13—Cu	125.75 (8)	N23—C23—N22	110.69 (10)
C12—N13—Cu	125.84 (8)	N23—C23—C25	125.06 (10)
O21—N21—O22	124.71 (11)	N22—C23—C25	124.23 (11)
O21—N21—C21	118.93 (11)	N22—C24—H24A	109.5
O22—N21—C21	116.32 (10)	N22—C24—H24B	109.5
C23—N22—C21	105.94 (9)	H24A—C24—H24B	109.5
C23—N22—C24	126.03 (10)	N22—C24—H24C	109.5
C21—N22—C24	128.03 (10)	H24A—C24—H24C	109.5
C23—N23—C22	107.29 (9)	H24B—C24—H24C	109.5
C23—N23—Cu	125.24 (8)	C23—C25—H25A	109.5
C22—N23—Cu	126.94 (8)	C23—C25—H25B	109.5
C12—C11—N12	108.11 (10)	H25A—C25—H25B	109.5
C12—C11—N11	127.19 (12)	C23—C25—H25C	109.5
N12—C11—N11	124.69 (10)	H25A—C25—H25C	109.5
C11—C12—N13	107.88 (11)	H25B—C25—H25C	109.5
C11—C12—H12A	126.1		
N11 ⁱⁱ —O12—N11—O11	-80.27 (13)	C11—N12—C13—C15	-178.76 (11)
N11 ⁱⁱ —O12—N11—C11	98.59 (11)	C14—N12—C13—C15	0.78 (19)
N11 ⁱⁱ —O12—N11—O12 ⁱⁱ	0.0	C23—N22—C21—C22	-1.06 (13)
C13—N12—C11—C12	0.40 (13)	C24—N22—C21—C22	179.19 (11)
C14—N12—C11—C12	-179.12 (12)	C23—N22—C21—N21	-175.98 (11)
C13—N12—C11—N11	-178.29 (11)	C24—N22—C21—N21	4.28 (19)

C14—N12—C11—N11	2.2 (2)	O21—N21—C21—C22	−160.72 (12)
O11—N11—C11—C12	6.01 (19)	O22—N21—C21—C22	16.86 (18)
O12—N11—C11—C12	−172.94 (12)	O21—N21—C21—N22	13.27 (17)
O12 ⁱⁱ —N11—C11—C12	−83.60 (13)	O22—N21—C21—N22	−169.15 (11)
O11—N11—C11—N12	−175.55 (12)	N22—C21—C22—N23	0.11 (13)
O12—N11—C11—N12	5.50 (18)	N21—C21—C22—N23	174.90 (11)
O12 ⁱⁱ —N11—C11—N12	94.84 (12)	C23—N23—C22—C21	0.90 (13)
N12—C11—C12—N13	−0.50 (13)	Cu—N23—C22—C21	−171.07 (8)
N11—C11—C12—N13	178.15 (11)	C22—N23—C23—N22	−1.62 (13)
C13—N13—C12—C11	0.40 (13)	Cu—N23—C23—N22	170.52 (7)
Cu—N13—C12—C11	−168.81 (8)	C22—N23—C23—C25	176.82 (11)
C12—N13—C13—N12	−0.15 (13)	Cu—N23—C23—C25	−11.04 (16)
Cu—N13—C13—N12	169.07 (8)	C21—N22—C23—N23	1.66 (13)
C12—N13—C13—C15	178.42 (11)	C24—N22—C23—N23	−178.59 (11)
Cu—N13—C13—C15	−12.36 (17)	C21—N22—C23—C25	−176.79 (11)
C11—N12—C13—N13	−0.16 (13)	C24—N22—C23—C25	2.96 (19)
C14—N12—C13—N13	179.38 (11)		

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

Hydrogen-bond geometry (\AA , °)

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
C25—H25C \cdots O22 ⁱⁱⁱ	0.98	2.39	3.2804 (17)	150
C15—H15B \cdots Cl2 ⁱⁱⁱ	0.98	2.73	3.6737 (13)	163
C12—H12A \cdots O22 ^{iv}	0.95	2.51	3.4029 (16)	156
C22—H22A \cdots Cl2 ^{iv}	0.95	2.75	3.6828 (12)	167
C24—H24C \cdots C11 ^v	0.98	2.84	3.7555 (13)	156

Symmetry codes: (iii) $x, y+1, z$; (iv) $-x+1, -y, -z+1$; (v) $-x+1, y+1/2, -z+1/2$.