



# The crystal structure of tetrameric copper(II) complexes, Hirshfeld surface analysis, and vector analyses of $\text{Cu}_4\text{OCl}_6\text{L}_4$ complexes with N-donor ligands

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## Abstract

It is known, that in the  $\text{Cu}_4\text{OX}_6\text{L}_4$  ( $X = \text{Cl}, \text{Br}$ ) complexes can be present many different ligands L, including bioligands. The synthesis and characterization of  $\text{Cu}_4\text{OBrCl}_6(\text{ron})_4$  (**1**) and  $\text{Cu}_4\text{OCl}_6(3\text{-Mepy})_4$  (**2**) (where ron is ronicol or 3-methanolpyridine and 3-Mepy is 3-methylpyridine) are reported. The complexes under study were X-ray structure analysis and Hirshfeld surface analysis. Tetranuclear  $\text{Cu}_4\text{OX}_6\text{L}_4$  complexes with molecular structure (Fig. 1) can help to better understand the role of donor–acceptor and electron-transfer properties in copper proteins. The coordination sphere about each copper(II) atom is trigonal bipyramidal with three chlorine atoms in the equatorial plane. The apical positions are occupied by the central oxygen atom and the nitrogen atom of the respective ligand ( $\text{CuCl}_3\text{ON}$ ). Here are studied chloridocomplexes of some N-donor ligands,  $L = \text{chloro-promazine}$ , ronicol (3-pyridylmethanol), 2-ethylpyrazine, seven derivatives of pyrazol and for comparisons 3-methylpyridine. The  $\text{Cu}_4\text{OCl}_6\text{L}_4$  molecule is regarded as a supramolecular model of interactions between bioligand L and hypothetical “round-shaped” coordination tetra-receptor  $\text{Cu}_4\text{OCl}_6$ . Vector calculations applied usually to mechanical and electrical macroconstructions are here applied to microconstructions represented by structures of  $\text{Cu}_4\text{OX}_6\text{L}_4$  molecules. For vector calculations each  $\text{Cu}_4\text{OX}_6\text{L}_4$  structure is placed (Fig. 1) into the three-dimensional Cartesian coordinate system with the central oxygen atom O1 placed in its origin 0. Studied bioligands are compared and described by molecular structural dynamics and corresponding shifts of electron densities by means of bond lengths ( $\text{O1-Cu}$ ,  $\text{Cu-L}$ ,  $\text{Cu-X}$ ) and structural distances ( $\text{O1}\cdots\text{X}$ ,  $\text{O1}\cdots\text{L}$ ).

**Keywords** Crystal structure · Tetramers · Vector analyses · Copper(II) complexes · Hirshfeld surface

## Introduction

Over 40 crystal structures tetranuclear copper(II) complexes of type  $\text{Cu}_4\text{OX}_6\text{L}_4$ , which various ligand L are known. Selected structural parameters for  $\text{Cu}_4\text{OCl}_6(\text{NL})_4$  complexes [NL = ligand with nitrogen donor atom, such as derivatives of amine (Becker et al. 2015; Bowmaker et al. 2011; Löw et al. 2013), pyridine (El-Toukhy et al. 1984; Gill and Sterns 1970; van Albada et al. 2011; Zhang et al. 2014), pyrazine (Näther and Jess 2002), pyrazole (Kashyap et al. 2013; Keij et al. 1991; Vafazadeh et al. 2015; Vafazadeh and Willis 2016), imidazole (Atria et al. 1999, Betanzos-Lara et al. 2012, Cortes et al. 2006, Jian et al. 2004, Lobana et al. 2011, Tosik et al. 2009), triazole (Richardson and Steel 2003; Skorda et al. 2005; Voitekhovich et al. 2009), and others (Kariuki and Newman 2018)], are summarized in the review by Melník and co-authors (Melník et al. 2011). The coordination sphere about each copper(II) atom is trigonal

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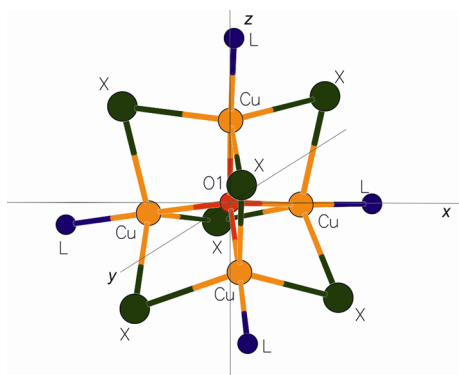
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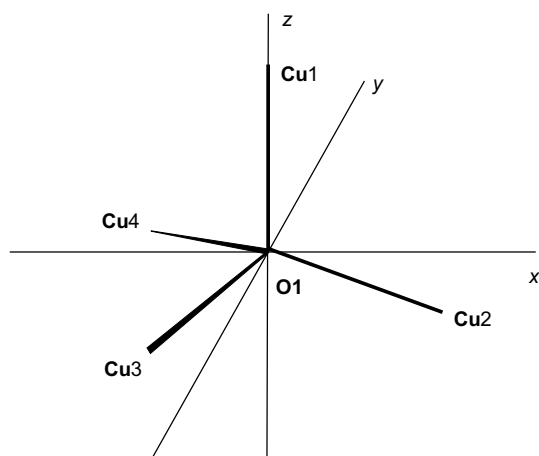
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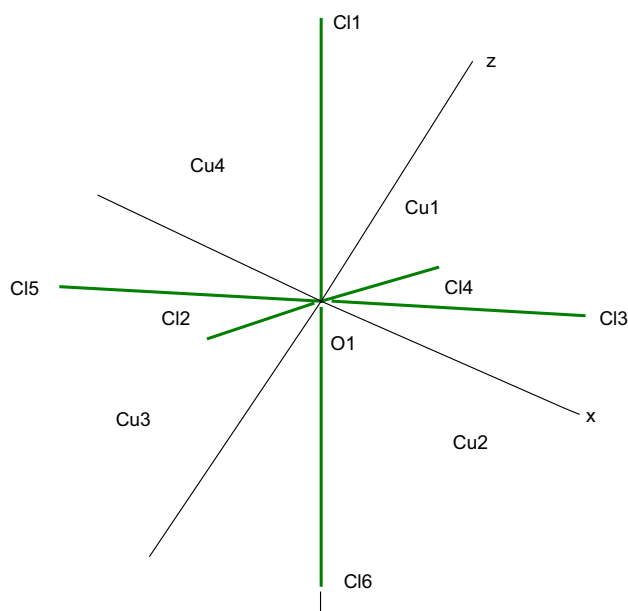
**Fig. 1** Structure of the  $\text{Cu}_4\text{OX}_6\text{L}_4$  molecule

bipyramidal with three chlorine atoms in the equatorial plane. The apical positions are occupied by the central oxygen atom and the nitrogen atom of the respective ligand ( $\text{CuCl}_3\text{ON}$ ). The equatorial plane is much less crowded than the apical sides, with mean  $\text{Cu}-\text{Cl}_{\text{eq}}$  bond distances of 2.41 (range 2.325–2.46) Å and the mean  $\text{Cu}-\text{O}_{\text{ap}}$  and  $\text{Cu}-\text{N}_{\text{ap}}$  bond distances of 1.905 (range 1.89–1.92) Å and 1.97 (range 1.930–2.025) Å. The mean  $\text{Cu}\cdots\text{Cu}$  separation of 3.110 Å (range from 3.090 to 3.133 Å) excludes a direct metal–metal bond. The deviation of the mean tetrahedral  $\text{Cu}-\mu\text{O}-\text{Cu}$  bond angle is from the ideal value of  $109.5^\circ$  (range  $0.3\text{--}6.4^\circ$ ). The  $\text{Cu}-\mu\text{Cl}-\text{Cu}$  bond angle ranges from  $76.7$  to  $82.7^\circ$  (mean  $80.4^\circ$ ).

The vector analysis never been applied to chemical objects represented by molecular structures before (Červeň 2015; Cikunov 1973). Figures 2 and 3 clearly demonstrate positions of bonds  $\text{Cu}-\text{O}$ ,  $\text{Cu}-\text{N}$  and  $\text{Cu}-\text{Cl}$  in the structures. These bonds form polyhedra: tetrahedron  $\text{OCu}_4$  and four trigonal bipyramids  $\text{OCuCl}_3\text{N}$ . The bonds  $\text{O}-\text{Cu}$  participate in both types of polyhedra. The central oxygen atom O of



**Fig. 2** The bond equivalent tetrahedron  $\text{O1}-\text{Cu}_4$  is represented by four bonding vectors



**Fig. 3** The non-bonding distance equivalent octahedron  $\text{O1}\cdots\text{X}_6$  is represented by six distance vectors  $\text{O1}\cdots\text{X}$  giving total octahedron vector  $\text{O}_\text{X}$ . For ideal octahedron,  $\text{O}_\text{X}=0$

$\text{O}-\text{Cu}$  bonds has a unique structural position, since, besides mentioned polyhedra, it participates in six  $\text{O}\cdots\text{Cl}$  distances that form the  $\text{OCl}_6$  octahedral system. Tetrahedron  $\text{OCu}_4$  is bonding homogeneous (equivalent), its four bonds  $\text{O}-\text{Cu}$  are of the same kind. Similarly, the polyhedral systems  $\text{OCl}_6$  are homogeneous (equivalent) by distances  $\text{OCl}_6$ . By sum of four bond vectors  $\text{O}-\text{Cu}$ , the tetrahedral vector  $T_{\text{Cu}}$  can be calculated, by sum of six distance vectors  $\text{O}\cdots\text{Cl}$  the octahedral vector  $\text{O}_{\text{Cl}}$ .

The trigonal bipyramid  $\text{OCuCl}_3\text{N}$  is bonding non-homogeneous unit. However, it is composed of the bonding homogeneous equatorial  $\text{CuCl}_3$  subunit and the bonding non-homogeneous axial  $\text{O}-\text{Cu}-\text{N}$  subunit. Each of the bonds  $\text{Cu}-\text{Cl}$  and  $\text{Cu}-\text{N}$  must be transformed to bond vectors in the orthogonal coordinate system with  $x$ ,  $y$ , and  $z$  axes. The transformed bond vectors serve for calculation of the equatorial  $E_{\text{Cl}}$  and axial  $A_{\text{ON}}$  vectors. From these subunit vectors, the total polyhedral trigonal bipyramidal vector  $P_{\text{Cu}}$  can be calculated. Most complex vector is represented by the total molecular  $\text{Cu}_4\text{OCl}_6\text{N}_4$  vector. This vector can be obtained by summing of four trigonal bipyramidal  $\text{OCuCl}_3\text{N}$  vectors,  $P_{\text{Cu}}$ .

For transformation of the bonds and interatomic distances to bond vectors and distance vectors, respectively, the molecular structure is placed into orthogonal coordinate system with atom O in the origin of coordinate axes  $x$ ,  $y$ ,  $z$ . One of the  $\text{O}-\text{Cu}$  bonds, mostly  $\text{O}-\text{Cu1}$ , is placed into  $+z$  axis and other  $\text{O}-\text{Cu}$  bond, mostly  $\text{O}-\text{Cu2}$ , into  $x, -z$  plane, as shown in Fig. 2. After transformation, each of the vectors

has its origin in the axes origin and the end point is defined by calculated coordinates  $x$ ,  $y$ ,  $z$ . The distance between the origin and the end point measured in  $pm$  is a vector magnitude, in this case, vector length. The direction of the vector is represented by the vector coordinates  $x$ ,  $y$ ,  $z$ .

Polyhedron vectors of tetrahedron  $T_{Cu}$  and octahedron  $O_{Cl}$  are calculated from vectors of transformed bonds and distances according to the equations:

$$T_{Cu} = \Sigma(O1 - Cu(1 - 4)).$$

$$O_{Cl} = \Sigma(O1 \dots Cl(1 - 6)).$$

For trigonal bipyramid  $O_1Cu_1Cl_3N_1$ :

$$P_{Cu1} = E_{Cl} + A_{O1N1}.$$

$E_{Cl}$  = equatorial vector of bonds  $Cu_1Cl_3$ .

$A_{O1N1}$  = axial vector of bonds  $O1-Cu1$  and  $Cu1-N1(L)$ .

The principles of vector methods and calculations are described in literature (Ondrejovič and Moncol 2015, 2017).

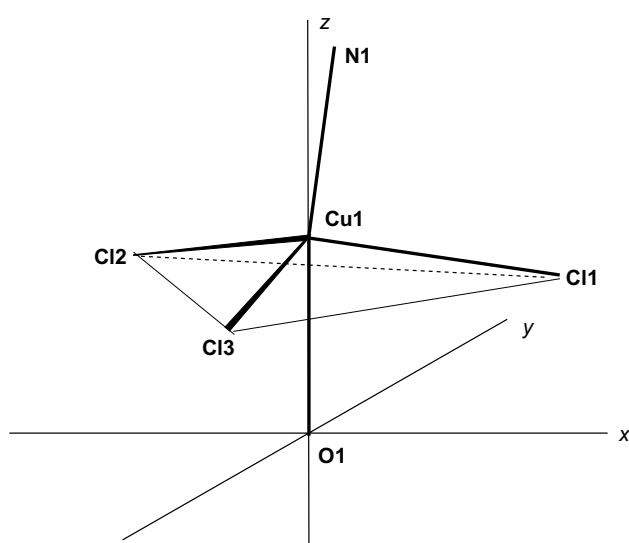
The use of the orthogonal coordinate system with the origin occupied by atom O of the  $OCu_4$  tetrahedron corresponds to the Schoenflies molecular structure parameter  $T_d$  and electron bonds representations of the  $OCu_4$  tetrahedral-bonding system. Both aspects, crystallographic and chemical, are connected by the same point group.

The  $Cu_4O_1X_6L_4$  structure consists of two basic polyhedra:

$O1 \rightarrow Cu$  giving total tetrahedron vector  $T_{Cu}$ . For ideal tetrahedron,  $T_{Cu} = 0$ .

Four trigonal bipyramidal coordination polyhedra (Fig. 4) of central copper(II) atoms are strongly bonding nonequivalent. Therefore,  $TB_{Cu} > 0$ .

The vector analyses have also been applied to ten structures of the  $Cu_4OCl_6L_4$  complex molecules with  $L = N$ -donor ligands. Presented results demonstrate correlations between structural molecular parameters and supramolecular



**Fig. 4** The trigonal bipyramidal coordination polyhedra of central copper(II) atoms

inner- and intermolecular contacts by hydrogen bonds and van der Waals interactions.

Structures of  $Cu_4OX_6L_4$  complexes are of considerable interest, since about 116 structures are registered in the CCDC database (Groom et al. 2016).

## Experimental

### Synthesis of the complexes

Chemicals, syntheses and characterization of the  $Cu_4OCl_6(ron)_4$  (**1**) complexes instruments have been described in our previous report (Koman and Ondrejovič 2013; Ondrejovič et al. 2006). Single crystals of the  $Cu_4OCl_6(3-Mepy)_4$  (**2**) complex were prepared by diffusion of the 3-Mepy ligand into a methanol solution of the  $Cu_4OCl_6(methanol)_4$  complex as described elsewhere (Norman and Rose 1989; Löw et al. 2013) similar to the 1-methanolpyridine synthesis.

### X-ray crystallography

Intensity data for  $Cu_4OCl_6(ron)_4$  (**1**) were collected using a Siemens P4 diffractometer with graphite monochromated  $MoK\alpha$  radiation (Siemens 1990). The diffraction intensities were corrected for Lorentz, polarization effects and absorption correction with XSCANS (Siemens XSCANS and XEMP 1994). Intensity data for  $Cu_4OCl_6(3-Mepy)_4$  (**2**) were collected using diffractometer Stoe StadiVari using Pilatus3R 300 K HPAD detector and microfocused X-ray source Xenocs Genix3D Cu HF (Cu  $K\alpha$  radiation) at 100 K. The structures were solved using the programs SIR-2011 (Burla et al. 2012) or SHELXT (Sheldrick 2015a) and refined by the full-matrix least-squares method on all  $F^2$  data using the program SHELXL-2018/3 (Sheldrick 2015b). Geometrical analysis was performed using SHELXL-2018/3. The structures were drawn by OLEX2 (Dolomanov et al. 2009) software.

Crystal data and conditions of data collection and refinement for complexes **1** and **2** are reported in Table 1.

### Hirshfeld surface analysis

Hirshfeld surface analysis (Hirshfeld 1977; Spackman and Jayalitaka 2009) and associated fingerprint plots (Parkin et al. 2007; Spackman et al. 2002) have been made using program CrystalExplorer (version 17.5) (Turner et al. 2017). The Hirshfeld surface of **1** has been calculated including all orientations of the disordered molecule with their partial occupancies.

**Table 1** Crystal data and structure refinement for complexes **1** and **2**

	Cu <sub>4</sub> OCl <sub>6</sub> (ron) <sub>4</sub> — <b>1</b>	Cu <sub>4</sub> OCl <sub>6</sub> (3-Mepy) <sub>4</sub> — <b>2</b>
Chemical formula	C <sub>24</sub> H <sub>28</sub> Cl <sub>6</sub> Cu <sub>4</sub> N <sub>4</sub> O <sub>5</sub>	C <sub>24</sub> H <sub>28</sub> Cl <sub>6</sub> Cu <sub>4</sub> N <sub>4</sub> O
<i>M</i> <sub>r</sub>	919.41	855.36
Crystal system, space group	monoclinic, <i>P</i> 2 <sub>1</sub> / <i>n</i>	orthorhombic, <i>P</i> bca
Temperature (K)	293(1)	100(1)
<i>a</i>	11.346(1) Å	12.8364(4) Å
<i>b</i>	15.807(2) Å	30.3137(8) Å
<i>c</i>	19.027(2) Å	16.3451(4) Å
α (°)	90	90
β (°)	106.27(3)	90
γ (°)	90	90
<i>V</i> (Å <sup>3</sup> )	3275.8(6)	6360.2(3)
<i>Z</i>	4	8
Radiation type	0.71073 Å (Mo K <sub>α</sub> )	1.54186 (Cu K <sub>α</sub> )
2θ range for data collection	8.36°–52.73°	5.832°–143.198°
Index ranges	– 1 ≤ <i>h</i> ≤ 14, – 1 ≤ <i>k</i> ≤ 19 – 23 ≤ <i>l</i> ≤ 23	– 11 ≤ <i>h</i> ≤ 15, – 37 ≤ <i>k</i> ≤ 27, – 19 ≤ <i>l</i> ≤ 18
μ (mm <sup>–1</sup> )	3.092	7.860
Crystal size (mm)	0.45 × 0.38 × 0.35	0.14 × 0.07 × 0.03
Diffractometer	Siemens P4	Stoe StadiVari
Absorption correction	Psi-scan	Multi-scan
Reflections collected	8250	52,601
Data/redtrains/parameters	6667/24/396	6003/0/356
<i>R</i> <sub>int</sub>	0.0408	0.0783
<i>R</i> [ <i>F</i> <sup>2</sup> > 2 σ ( <i>F</i> <sup>2</sup> )], <i>wR</i> ( <i>F</i> <sup>2</sup> ), <i>S</i>	<i>R</i> <sub>1</sub> = 0.0657 <i>wR</i> <sub>2</sub> = 0.1953	<i>R</i> <sub>1</sub> = 0.0378 <i>wR</i> <sub>2</sub> = 0.0866
<i>S</i>	1.011	0.907
Largest diff. peak and hole (e. Å <sup>–3</sup> )	1.05 and – 1.52	0.75 and – 0.41

## Results and discussion

### Crystal structures

Complex **1** crystallizes in monoclinic space group *P*2<sub>1</sub>/*n*, and other hand compound **2** crystallizes in orthorhombic space group *P*bca. All copper atoms are joined by three μ<sub>2</sub>-chlorido bridging ligands and one μ<sub>4</sub>-oxido ligand. The coordination polyhedron around all copper atoms in both tetranuclear complexes is trigonal-bipyramide. The trigonal plane is built up by three chlorido ligands, and axial positions are occupied by one oxygen atom and one pyridine nitrogen atom of ronicol (3-hydroxymethylpyridine) (**1**) or 3-methylpyridine (**2**). The structures of title complexes Cu<sub>4</sub>OCl<sub>6</sub>(ron)<sub>4</sub>—**1** and Cu<sub>4</sub>OCl<sub>6</sub>(3-Mepy)<sub>4</sub>—**2** can be described as a system of three penetrating polyhedral. These polyhedral, the OCu<sub>4</sub> tetrahedron, the OCl<sub>6</sub> octahedron, and four CuOCl<sub>3</sub>N trigonal bipyramids can be distorted due to both intramolecular and intermolecular interactions (Ondrejovič et al. 2000). Selected interatomic distances are listed in Table 2. The molecular structures of Cu<sub>4</sub>OCl<sub>6</sub>(ron)<sub>4</sub>—**1** and Cu<sub>4</sub>OCl<sub>6</sub>(3-Mepy)<sub>4</sub>—**2** are shown in Fig. 5.

The crystal structures of **1** is drawn in Fig. 6. The molecules of Cu<sub>4</sub>OCl<sub>6</sub>(ron)<sub>4</sub> are connected through O–H•••O hydrogen bonds between hydroxyl oxygen atoms of 3-pyridylmethanol ligands [O2–H2•••O1, O2–H2•••O1A, O3–H3•••O4, O3–H3•••O4A, O4–H4•••O4 and O4A–H4A•••O4A, the O•••O distances are in the range 2.74–3.06 Å (See ESI Table S1)] and O1–H1•••Cl15 hydrogen bond between hydroxyl oxygen atom of 3-pyridylmethanol ligand (O1) and chlorine atom (Cl15) [the distance of O1•••Cl15 [3.218(19)] Å, (See ESI Table S1)] and forming 3D supramolecular network (Fig. 6). The O–H•••O and O–H•••Cl hydrogen-bond system of **1** is enriched by weaker C23–H23•••O3 hydrogen-bonding interactions between carbon atom of pyridine ring (C23) and hydroxyl oxygen atom of 3-pyridylmethanol ligand (O3) [the distance of C23•••O3 (3.403(12)) Å, (See ESI Table S1)]; and C–H•••Cl hydrogen-bonding interactions between carbon atoms of pyridine ring (C33) or methylene group (C16, C46) of 3-pyridylmethanol ligands and chlorine atoms (Cl11, Cl14Cl13) [C16–H16B•••Cl11, C33–H33•••Cl14 and C46–H46B•••Cl13, the C•••Cl distances are in the range 3.40–3.63 Å (See ESI Table S1)].

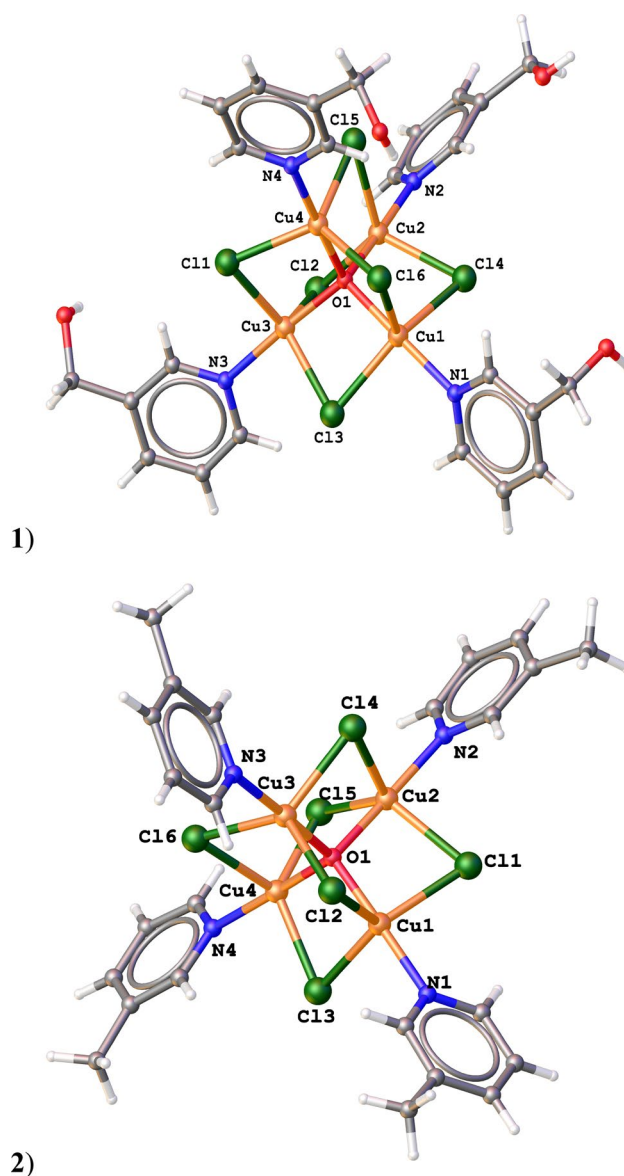
**Table 2** Selected bond distances [Å] for complexes **1** and **2**

$\text{Cu}_4\text{OCl}_6(\text{ron})_4\text{—1}$			
Cu1–O1	1.916(4)	Cu1–N1	1.979(5)
Cu2–O1	1.909(4)	Cu2–N2	1.969(5)
Cu3–O1	1.909(4)	Cu3–N3	1.996(6)
Cu4–O1	1.916(4)	Cu4–N4	1.989(5)
Cu1–Cl3	2.363(2)	Cu3–Cl1	2.401(2)
Cu1–Cl4	2.402(2)	Cu3–Cl2	2.395(2)
Cu1–Cl6	2.452(2)	Cu3–Cl3	2.402(2)
Cu2–Cl2	2.386(2)	Cu4–Cl1	2.444(2)
Cu2–Cl4	2.447(2)	Cu4–Cl5	2.428(2)
Cu2–Cl5	2.471(2)	Cu4–Cl6	2.395(2)
$\text{Cu}_4\text{OCl}_6(3\text{-Mepy})_4\text{—2}$			
Cu1–O1	1.898(2)	Cu1–N1	1.979(3)
Cu2–O1	1.901(3)	Cu2–N2	1.983(3)
Cu3–O1	1.921(3)	Cu3–N3	1.980(3)
Cu4–O1	1.904(3)	Cu4–N4	1.969(3)
Cu1–Cl1	2.381(1)	Cu3–Cl2	2.400(1)
Cu1–Cl2	2.436(1)	Cu3–Cl4	2.395(1)
Cu1–Cl3	2.378(1)	Cu3–Cl6	2.361(1)
Cu2–Cl1	2.416(1)	Cu4–Cl3	2.398(1)
Cu2–Cl4	2.464(1)	Cu4–Cl5	2.391(1)
Cu2–Cl5	2.376(1)	Cu4–Cl6	2.486(1)

On the other hand, complex **2** forms also 3D supramolecular network (Fig. 7), but complex molecules  $\text{Cu}_4\text{OCl}_6(3\text{-Mepy})_4$  are joined only via weak  $\text{C}\cdots\text{H}\cdots\text{Cl}$  hydrogen-bonding interactions. The  $\text{C}\cdots\text{H}\cdots\text{Cl}$  hydrogen-bonding interactions in crystal structure of **2** are observed between carbon atoms of pyridine ring (C3, C4) or methyl group (C12, C18, C24) of 3-methylpyridine ligands, and chlorine atoms (Cl3, Cl4C6) [ $\text{C3}\text{--H3}\cdots\text{Cl6}$ ,  $\text{C4}\text{--H4}\cdots\text{Cl3}$ ,  $\text{C12}\text{--H12C}\cdots\text{Cl4}$ ,  $\text{C18}\text{--H18C}\cdots\text{Cl4}$  and  $\text{C24}\text{--H24B}\cdots\text{Cl6}$ , the  $\text{C}\cdots\text{Cl}$  distances are in the range 3.52–3.82 Å (See ESI Table S1)]. The crystal structure of **2** exhibits also  $\pi\text{--}\pi$  stacking interactions (Janiak 2000) between pyridine rings [ $\text{N2}/\text{C7}\text{--}\text{C11}$ ] and [ $\text{N4}/\text{C19}\text{--}\text{C23}$ ] with centroid–centroid distance of 3.63 Å and shift distance of 0.63 Å.

### Hirshfeld surface analysis

Hirshfeld surface analysis was used to further study the intermolecular interactions of the crystal structures of both compounds. Figures 8, 9 show the 3D Hirshfeld surface of **1** and **2**, respectively. The 3D Hirshfeld surfaces have been mapped over  $d_{\text{norm}}$  shape index (Figs. 8, 9). The surfaces are shown as transparent to allow visualization of the molecular moiety around which they were calculated. As shown in Figs. 8, 9, the deep red spots on the  $d_{\text{norm}}$  Hirshfeld surfaces indicate the close-contact interactions, which are mainly

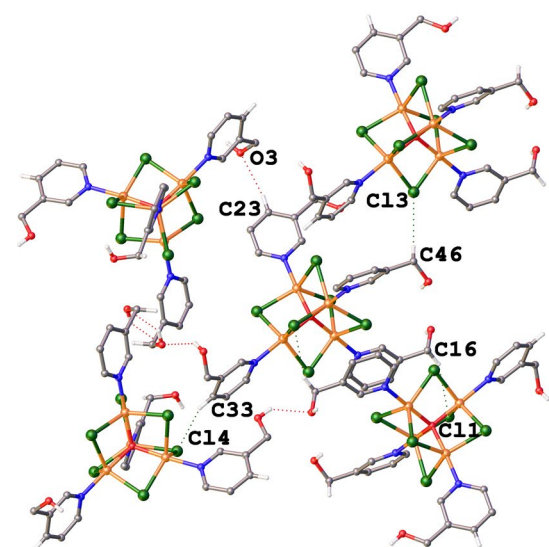
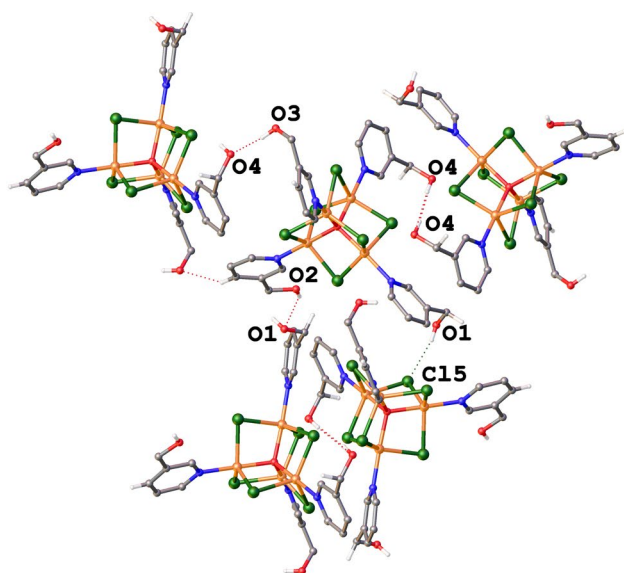


**Fig. 5** Molecular structures of the  $\text{Cu}_4\text{OCl}_6(\text{ron})_4\text{—1}$  and  $\text{Cu}_4\text{OCl}_6(3\text{-Mepy})_4\text{—2}$

responsible for the significant intermolecular hydrogen-bonding interactions.

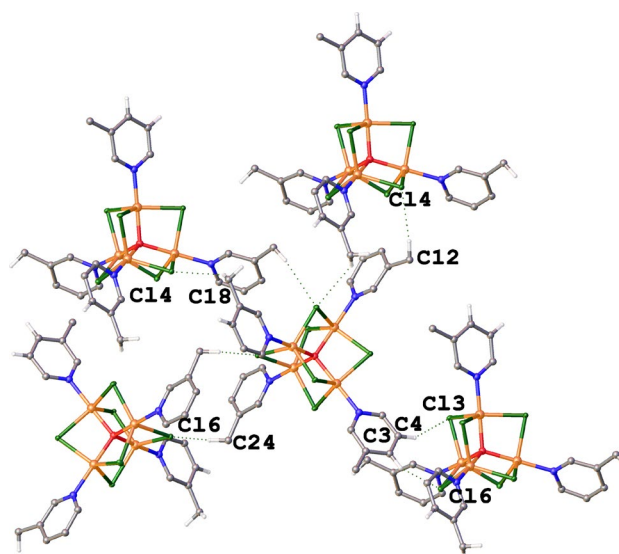
The 3D Hirshfeld surface illustration of **1** (Fig. 8) shows the deep red areas representing  $\text{O}\cdots\text{H}\cdots\text{O}$ ,  $\text{O}\cdots\text{H}\cdots\text{Cl}$ , and also weaker  $\text{C}\cdots\text{H}\cdots\text{Cl}$  hydrogen-bonding interactions. The 3D Hirshfeld surface illustration of **2** (Fig. 9) also shows weaker  $\text{C}\cdots\text{H}\cdots\text{Cl}$  hydrogen-bonding interactions. The Hirshfeld surface plotted over shape index of **2** visualizes the  $\pi\text{--}\pi$  stacking interactions (Janiak 2000) by the presence of adjacent red and blue triangles (Fig. 9).

The Hirshfeld 2D fingerprints of **1** and **2** compounds are illustrated in supplementary material (See ESI Figs. S3 and S4). The Hirshfeld 2D fingerprint plots allow a



**Fig. 6** The hydrogen-bonding network in crystal structure of  $\text{Cu}_4\text{OCl}_6(\text{ron})_4$  (**1**). The stronger  $\text{O}\cdots\text{H}\cdots\text{O}$  and  $\text{O}\cdots\text{H}\cdots\text{Cl}$  hydrogen bonds are drawn in top. The weaker  $\text{C}\cdots\text{H}\cdots\text{O}$  and  $\text{C}\cdots\text{H}\cdots\text{Cl}$  hydrogen-bonding interactions are pictured in bottom

quick and easy identification of the significant intermolecular interaction map on the molecular surface. As shown in Fig. S3 in supplementary materials, the strong  $\text{H}\cdots\text{O}/\text{O}\cdots\text{H}$ , and weak  $\text{H}\cdots\text{C}/\text{C}\cdots\text{H}$ , and  $\text{H}\cdots\text{Cl}/\text{Cl}\cdots\text{H}$  hydrogen-bonding interactions cover the 10.0, 10.1 and 22.8%, respectively, of the total Hirshfeld surface with two distinct spikes in the 2D fingerprint plots, indicating hydrogen-bonding interactions are the most significant interactions in the crystal. As shown in Fig. S3 in supplementary material, in the middle of scattered points in the 2D fingerprint plots,  $\text{H}\cdots\text{H}$  interactions cover 42.0% of the total Hirshfeld surface. As shown in Fig. S4 in supplementary



**Fig. 7** The weak  $\text{C}\cdots\text{H}\cdots\text{Cl}$  hydrogen-bonding interactions in crystal structure of  $\text{Cu}_4\text{OCl}_6(3\text{-Mepy})_4$  (**2**)

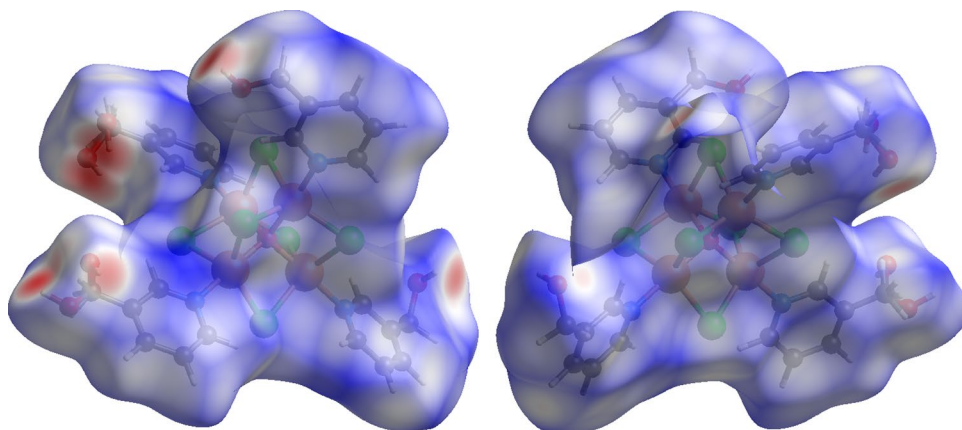
material, in scattered points in the 2D fingerplots,  $\text{H}\cdots\text{C}/\text{C}\cdots\text{H}$  and  $\text{H}\cdots\text{Cl}/\text{Cl}\cdots\text{H}$  interactions cover in the 17.5 and 29.1%, respectively, of the total Hirshfeld surface. In scattered points of the 2D fingerplot in **2** (See Fig. S4),  $\text{H}\cdots\text{H}$  and  $\text{C}\cdots\text{C}$  interactions illustrate covering of 43.3 and 2.5%, respectively.

### Vector analysis

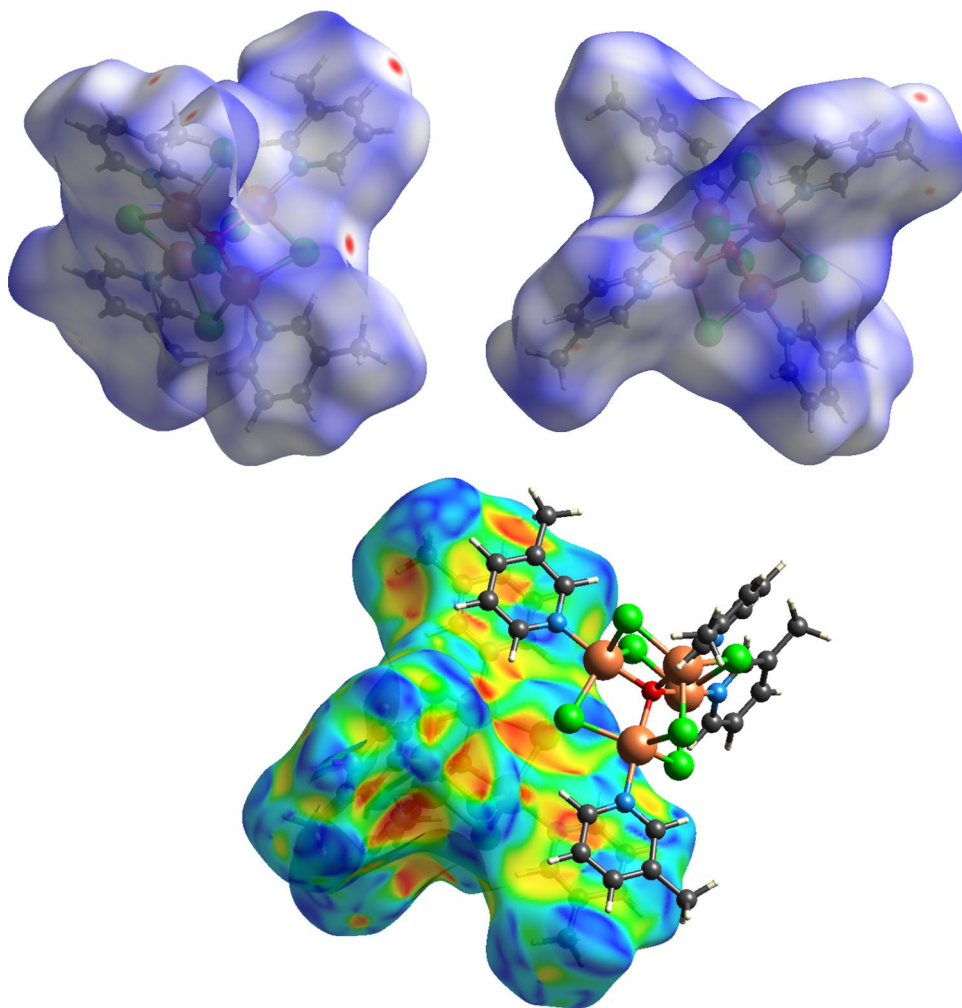
Analyzed ten structures of the  $\text{Cu}_4\text{OCl}_6\text{L}_4$  complexes are presented in Table 3. The structures are characterized by crystallographic data including codes of CCDC database. Calculated vector parameters of tetrahedron  $TCu$ , octahedron  $OCl$ , trigonal bipyramids  $\text{Cu}1$ ,  $\text{Cu}2$ ,  $\text{Cu}3$ ,  $\text{Cu}4$ , and corresponding molecule  $MOL$  are characterized by length and sector of the three-dimensional Cartesian coordinate system (three combined symbols of + and -). Intramolecular and intermolecular interactions are demonstrated by similar way. Red symbols demonstrate the identical directions.

The results presented in Tables 3, 4 are not surprising because of strong differences in bioactivity of ligands  $L$ . There is some classification of bioactive ligands  $L$  through the graphical comparison of vectors in Dependence 1 (See ESI Fig. S3) and Dependence 2 (See ESI Fig. S4). In Dependence 1, it is clearly seen how correspond tetrahedral  $TCu$ , octahedral  $OCl$ , and molecular  $MOL$  vectors. It is clearly seen that disorders have very weak influence on vector data. However, in Dependence 2, it is clearly seen that molecular interactions for complexes 4A, 4B are different, for complexes 5A, 5B very different, but for complexes 6A, 6B, there is practically no difference.

**Fig. 8** View of the three-dimensional Hirshfeld surface of **1** plotted over  $d_{\text{norm}}$  in the range – 1.1524 to 1.2567 a.u.



**Fig. 9** View of the three-dimensional Hirshfeld surface of **2** plotted over  $d_{\text{norm}}$  in the range – 0.0827 to 1.6290 a.u. (top) and shape index (bottom)



## Conclusions

1. The crystal structure of both complexes shows 3D supramolecular networks. The 3D supramolecular network of  $\text{Cu}_4\text{OCl}_6(\text{ron})_4$  (**1**) is formed through  $\text{O}-\text{H}\cdots\text{O}$ ,  $\text{O}-$

$\text{H}\cdots\text{Cl}$ , and  $\text{C}-\text{H}\cdots\text{Cl}$  hydrogen bonds. The  $\text{C}-\text{H}\cdots\text{Cl}$  hydrogen bonds form the 3D supramolecular network of  $\text{Cu}_4\text{OCl}_6(3\text{-Mepy})_4$  (**2**). The chlorine atoms of both complexes are acceptors of hydrogen bonds, which are also confirmed by Hirshfeld surfaces analysis.

**Table 3** Ligands and crystallographic data of selected  $\text{Cu}_4\text{OCl}_6\text{L}_4$  complexes, where L = N-donor ligands

LIGAND L NUMBER	LIGAND L SOLVATE, Disorder	CRYSTALLOGRAPHIC DATA, Refcode
1	<i>Chlor-promazine</i> 4.C <sub>6</sub> H <sub>6</sub>	Monoclinic <i>P2<sub>1</sub>/a</i> <i>R</i> = 6.5% PIVHAD (Yamada et al. 1994)
2A	<i>3-Pyridylmethanol, ronicol</i> A: O1-H1, O4-H4 Disorder	Monoclinic <i>P2<sub>1</sub>/n</i> <i>R</i> = 6.57% (This paper—1)
3	<i>2-Ethylpyrazine</i>	Monoclinic <i>P2<sub>1</sub>/n</i> <i>R</i> = 3.56% IDICUT (Nather et al. 2002)
4A	<i>3,5-Dimethylpyrazole</i> A: C14 Disorder	Monoclinic <i>P2/n</i> <i>R</i> = 3.9% BOGCEG (Stibrany et al. 2007)
4B	<i>3,5-Dimethylpyrazole</i> B: C14A Disorder	Monoclinic <i>P2/n</i> <i>R</i> = 3.9% BOGCEG (Stibrany et al. 2007)
5A	<i>3,5-Dimethylpyrazole</i> 4.C <sub>2</sub> H <sub>5</sub> OH A: C16, <i>Oc</i> : 0.6 Disorder	Triclinic <i>P-1</i> <i>R</i> = 8.3% RIRKOT (Jacimovic et al. 2007)
5B	<i>3,5-Dimethylpyrazole</i> 4.C <sub>2</sub> H <sub>5</sub> OH B: C17, <i>Oc</i> : 0.4 Disorder	Triclinic <i>P-1</i> <i>R</i> = 8.3% RIRKOT (Jacimovic et al. 2007)
6A	<i>3,5-Diisopropyl-1H-pyrazole</i> A: C14, <i>Oc</i> : 0.5 Disorder	Monoclinic <i>C2/c</i> <i>R</i> = 3.03% <i>T</i> = 100 K DIFTIX (Kashyap et al. 2013)
6B	<i>3,5-Diisopropyl-1H-pyrazole</i> B: C14A, <i>Oc</i> : 0.5 Disorder	Monoclinic <i>C2/c</i> <i>R</i> = 3.03% <i>T</i> = 100 K DIFTIX (Kashyap et al. 2013)
7	<i>3,4-Dimethyl-5-phenyl-pyrazole</i>	Monoclinic <i>I2/a</i> <i>R</i> = 5.2% JIWKAB (Keij et al. 1991)
8	<i>5-(2,4,6-Trimethyl-phenyl)pyrazole</i>	Monoclinic <i>C2/c</i> <i>R</i> = 3.6% WUXBOG (Liu et al. 2003)
9	<i>3-Methyl-5-phenyl-1H-pyrazole</i>	Tetragonal <i>P4/n</i> <i>R</i> = 6.65% IJIWII (He 2011)
10	<i>3-Methylpyridine</i>	Orthorhombic <i>Pbca</i> <i>R</i> = 3.78 (This paper—2)



**Table 4** Vector and interaction parameters [ $pm$ ] for structure of selected  $Cu_4OCl_6L_4$  complexes  
L = N-donor ligands

Ligand	Tetra	Octa	Trigonal polyhedron				Molecule
L	$T_{Cu}$	$O_{Cl}$	Cu1	Cu2	Cu3	Cu4	MOL
1	11.14	16.70	62.35	59.69	61.12	59.37	26.45
	(---)	(---)	(---)	(++)	(+-)	(+++)	(-++)
	Interactions		1112.1	2301.1	1835.5	1431.1	3006.4
			(++)	(+++)	(+-)	(-+-)	(-+-)
2A	4.81	47.23	329.08	322.10	322.23	329.30	98.65
	(+-)	(+++)	(++)	(+-)	(---)	(-+-)	(+++)
	Interactions		1319.85	563.69	637.72	870.25	1364.79
			(+++)	(---)	(-+-)	(-+-)	(-+-)
3	2.10	30.09	329.40	324.40	323.16	320.55	67.13
	(+-)	(-+-)	(++)	(+-)	(---)	(-+-)	(-+-)
	Interactions		990.64	825.74	883.63	829.35	772.31
			(++)	(+-)	(---)	(-+-)	(-+-)
			<i>Cu1</i>	<i>Cu2</i>	<i>Cu1A</i>	<i>Cu2A</i>	
4A	4.55	79.41	325.81	322.30	321.33	322.32	157.34
	(-+-)	(-+-)	(++)	(+-)	(---)	(-+-)	(-+-)
	Interactions		826.79	690.82	765.48	529.20	471.02
			(+++)	(+-)	(---)	(-+-)	(-+-)
4B	4.66	79.39	321.33	322.30	325.22	322.32	157.34
	(-+-)	(+-)	(+-)	(+-)	(---)	(-+-)	(+-)
	Interactions		957.93	695.40	659.23	528.16	904.66
			(+++)	(+-)	(---)	(-+-)	(+++)
5A	13.78	96.53	321.33	324.68	321.19	321.22	185.65
	(-+-)	(+-)	(+-)	(+-)	(-+-)	(---)	(+-)
	Interactions		662.62	265.67	648.86	734.25	445.33
			(+++)	(+-)	(-+-)	(---)	(+++)
5B	13.78	101.95	321.33	324.68	323.09	323.71	213.48
	(-+-)	(-+-)	(+-)	(+-)	(-+-)	(---)	(-+-)
	Interactions		850.61	461.40	907.42	656.08	1636.93
			(-+-)	(+-)	(-+-)	(-+-)	(-+-)
			<i>Cu1</i>	<i>Cu2</i>	<i>Cu1A</i>	<i>Cu2A</i>	
6A	9.56	69.18	326.14	325.60	314.18	325.59	142.94
	(-+-)	(---)	(-+-)	(+-)	(-+-)	(---)	(---)
	Interactions		231.87	788.21	298.31	898.07	519.66
			(-+-)	(+-)	(+++)	(---)	(-+-)
6B	9.56	69.18	314.42	325.60	325.92	325.59	142.88
	(-+-)	(---)	(+++)	(+-)	(-+-)	(---)	(-+-)
	Interactions		298.41	788.21	231.28	898.07	500.48
			(-+-)	(+-)	(-+-)	(---)	(---)
			<i>Cu1</i>	<i>Cu2</i>	<i>Cu1B</i>	<i>Cu2B</i>	
7	2.24	6.24	322.76	327.00	322.78	327.03	14.62
	(+-)	(+-)	(-+-)	(+-)	(-+-)	(---)	(+-)
	Interactions		749.94	1473.63	677.10	1473.16	494.07
			(+-)	(+-)	(+-)	(-+-)	(+-)
8	6.95	8.90	327.80	318.88	327.78	318.87	45.59
	(+-)	(-+-)	(-+-)	(+-)	(---)	(-+-)	(-+-)
	Interactions		1137.55	878.87	1179.03	878.76	974.48
			(-+-)	(-+-)	(+-)	(-+-)	(-+-)
			<i>Cu1</i>	<i>Cu1A</i>	<i>Cu1F</i>	<i>Cu1G</i>	
9	0.02	0.05	325.86	325.85	325.90	325.85	0.06
	(+0)	(-+-)	(+++)	(+-)	(---)	(-+-)	(---)

**Table 4** (continued)

Ligand	Tetra	Octa	Trigonal polyhedron				Molecule
	Interactions		698.18	698.29	698.28	698.30	0.28
			(--+)	(++-)	(---)	(+-)	(+-)
10	3.92	74.68	327.03	320.30	337.94	320.41	170.85
	(---)	(+-)	(++)	(+-)	(+-)	(---)	(+-)
	Interactions		565.48	741.59	502.04	741.33	1220.29
			(--+)	(+-)	(+-)	(+-)	(+-)

- The Hirshfeld surfaces analysis of  $\text{Cu}_4\text{OCl}_6(3\text{-Mepy})_4$  (**2**) confirms also  $\pi$ - $\pi$  stacking interactions between pyridine rings.
- This paper presents structural data of two tetrameric copper(II) complexes which contain a  $\mu_4$ -oxo group tetrahedrally coordinated to four copper(II) centers. Each pair of copper(II) centers is bridged by a single chlorine atoms. The coordination sphere about each copper(II) is trigonal-bipyramidal, which three chlorine atoms in the trigonal plane. One apical position about each copper(II) is occupied by oxygen atom, which is tetrahedrally coordinated to four copper(II) atoms. The second axial position is occupied by ligands with nitrogen donor atom.
- The vector analyses combine chemical and structural aspects of coordination compounds into one quantitative parameter—structural vector which is correlated with intra- and intermolecular interactions. As a method of crystallochemistry, it can be applied to arbitrary molecule.
- Structural vector parameters of biomolecules L coordinated to  $\text{Cu}_4\text{OX}_6$  receptors provide useful quantitative information about possible interaction activity of biomolecules in the real bioenvironment.
- Vectors of nonvalence supramolecular, hydrogen bond, and Van der Waals interactions correlate with the bond vectors of tetrahedrons  $\text{OCu}_4$ , distance vectors of octahedrons  $\text{OCl}_6$ , and total molecular vectors in molecular structures of  $\text{Cu}_4\text{OX}_6\text{L}_4$  complexes.

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