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Di- μ -chlorido-bis{[2-(2-pyridylmethylamino)ethanesulfonato- κ^3 N,N',O]-copper(II)}

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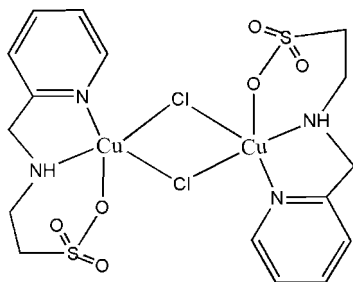
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 Key indicators: single-crystal X-ray study; $T = 291$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.024; wR factor = 0.068; data-to-parameter ratio = 16.0.

In the title compound, $[\text{Cu}_2(\text{C}_8\text{H}_{11}\text{N}_2\text{O}_3\text{S})_2\text{Cl}_2]$, the Cu atoms are five-coordinated in a distorted square-pyramidal geometry by three donor atoms of the deprotonated anionic 2-(2-pyridylmethylamino)ethanesulfonate (pmt) ligand and two Cl atoms. The Cl atoms bridge two Cu atoms, giving a binuclear structure; the centroid of the Cu_2Cl_2 ring lies on a crystallographic center of inversion. The complex is stabilized by hydrogen bonds and π - π stacking interactions [average interplanar distance = 3.4969 (1) Å and ring-centroid separation distance = 4.1068 (4) Å].

Related literature

 For related literature, see: Li *et al.* (2006, 2007a,b).


Experimental

Crystal data

$[\text{Cu}_2(\text{C}_8\text{H}_{11}\text{N}_2\text{O}_3\text{S})_2\text{Cl}_2]$	$\gamma = 113.043$ (2)°
$M_r = 628.48$	$V = 544.9$ (1) Å ³
Triclinic, $P\bar{1}$	$Z = 1$
$a = 8.294$ (1) Å	Mo $K\alpha$ radiation
$b = 8.362$ (1) Å	$\mu = 2.43$ mm ⁻¹
$c = 9.110$ (1) Å	$T = 291$ (2) K
$\alpha = 103.773$ (2)°	$0.33 \times 0.20 \times 0.09$ mm
$\beta = 98.118$ (2)°	

Data collection

Bruker APEXII CCD area-detector diffractometer	3270 measured reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	2385 independent reflections
$T_{\min} = 0.504$, $T_{\max} = 0.811$	2224 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.009$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.023$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.067$	$\Delta\rho_{\text{max}} = 0.48$ e Å ⁻³
$S = 1.10$	$\Delta\rho_{\text{min}} = -0.34$ e Å ⁻³
2385 reflections	
149 parameters	

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H1N}\cdots\text{O3}^{\text{i}}$	0.95 (3)	2.20 (3)	2.966 (2)	137 (2)

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

Data collection: APEX2 (Bruker, 2004); cell refinement: APEX2; data reduction: SAINT (Bruker, 2004); 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.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IM2051).

References

- Bruker (2004). APEX2 and SAINT. Bruker AXS Inc., Madison, Wisconsin, USA.
- Li, J.-X., Jiang, Y.-M. & Li, H.-Y. (2006). *Acta Cryst.* E62, m2984–m2986.
- Li, J.-X., Jiang, Y.-M. & Wang, J.-G. (2007a). *Acta Cryst.* E63, m213–m215.
- Li, J.-X., Jiang, Y.-M. & Wang, J.-G. (2007b). *Acta Cryst.* E63, m601–m603.
- Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst.* A64, 112–122.

supplementary materials

Acta Cryst. (2008). E64, m341 [doi:10.1107/S1600536808000779]

Di- μ -chlorido-bis{[2-(2-pyridylmethylamino)ethanesulfonato- κ^3N,N',O]copper(II)}

Z.-X. Du, J.-H. Qin and J.-G. Wang

Comment

Previously, one Co (Li *et al.*, 2006) and two Cu complexes (Li *et al.*, 2007a,b) containing the reduced schiff base ligand 2-pyridine-2-ylmethylamino-ethanesulfonic acid (Hpmt) have been reported. Herein we describe the structure of another dinuclear Cu compound, Figure 1.

The title compound is a binuclear Cu complex and each Cu center has square-pyramidal geometry formed by two N and one O atoms of an anionic pmt ligand and two chlorine atoms. The plane N1/N2/O1/Cl1A defines the base of the pyramid while Cl1 occupies the apical position. Cu1 is situated 0.168 (1) Å above the N1/N2/O1/Cl1A plane. The chlorine atoms bridge the Cu atoms to form this binuclear structure, with a Cu1...Cu1(-x, -y + 2, -z + 1) distance of 3.461 (2) Å.

The N—H donor and S=O acceptor (Table 1) groups of the pmt ligand participate in hydrogen bonding and they join the dinuclear complex units into a one-dimensional chain structure along *b* axis (Figure 2 and Table 1). These chains are further expanded into a two-dimensional network *via* π - π stacking between the pyridine rings of adjacent parallel chains. The interplanar average distance and ring-centroid separation distance are 3.4969 (1) Å and 4.1068 (4) Å, respectively.

Experimental

The ligand 2-pyridin-2-ylmethylamino-ethanesulfonic acid (Hpmt) was prepared according to the method of Li *et al.* (2006). 10 ml of an aqueous solution of CuCl₂ × 2 H₂O (0.171 g, 1 mmol) was dropped into 10 ml of a methanolic solution of Hpmt (0.216 g, 1 mmol). Then the mixture was stirred for 6 h at 343 K. The filtrate was left to stand under air for about one week to obtain blue block-shaped crystals. Analysis, found (%): C, 30.51; H, 3.50; N, 8.91; S, 10.18. [C₁₆H₂₂N₄O₆S₂Cl₂Cu₂] requires (%): C,30.55; H,5.55; N,12.96; S,14.81.

Refinement

H atoms bonded to C and N were positioned geometrically with C—H distances of 0.93 or 0.97 Å and a N—H distance of 0.95 Å, respectively, and treated as riding atoms with $U_{iso}(H) = 1.2 U_{eq}(C \text{ or } N)$.

Figures

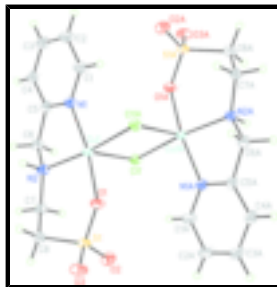


Fig. 1. The molecular structure of (I) with atom-numbering scheme. Atoms with the suffix A are at the symmetry position $(-x, -y + 2, -z + 1)$.

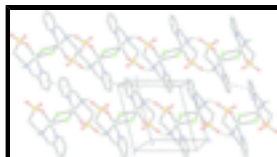


Fig. 2. Packing diagram, projected on the bc plane, showing the hydrogen bonding and π - π stacking interactions. H atoms have partially been omitted for the sake of clarity.

Di- μ -chlorido-bis[[2-(2-pyridylmethylamino)ethanesulfonato- κ^3N,N',O]copper(II)]

Crystal data

$[\text{Cu}_2(\text{C}_8\text{H}_{11}\text{N}_2\text{O}_3\text{S})_2\text{Cl}_2]$

$M_r = 628.28$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 8.294\ (1)\ \text{\AA}$

$b = 8.362\ (1)\ \text{\AA}$

$c = 9.110\ (1)\ \text{\AA}$

$\alpha = 103.773\ (2)^\circ$

$\beta = 98.118\ (2)^\circ$

$\gamma = 113.043\ (2)^\circ$

$V = 544.9\ (1)\ \text{\AA}^3$

$Z = 1$

$F_{000} = 318$

$D_x = 1.915\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation

$\lambda = 0.71073\ \text{\AA}$

Cell parameters from 2667 reflections

$\theta = 2.4\text{--}28.3^\circ$

$\mu = 2.43\ \text{mm}^{-1}$

$T = 291\ (2)\ \text{K}$

Block, blue

$0.33 \times 0.20 \times 0.09\ \text{mm}$

Data collection

Bruker APEXII CCD area-detector diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

$T = 291\ (2)\ \text{K}$

ϕ and ω scans

Absorption correction: multi-scan (SADABS; Sheldrick, 1996)

$T_{\min} = 0.505$, $T_{\max} = 0.811$

3270 measured reflections

2385 independent reflections

2224 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.009$

$\theta_{\text{max}} = 27.5^\circ$

$\theta_{\text{min}} = 2.4^\circ$

$h = -10 \rightarrow 10$

$k = -10 \rightarrow 10$

$l = -9 \rightarrow 11$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.023$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.067$	$w = 1/[\sigma^2(F_o^2) + (0.037P)^2 + 0.2091P]$
$S = 1.10$	where $P = (F_o^2 + 2F_c^2)/3$
2385 reflections	$(\Delta/\sigma)_{\max} = 0.001$
149 parameters	$\Delta\rho_{\max} = 0.48 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	$\Delta\rho_{\min} = -0.34 \text{ e } \text{\AA}^{-3}$
	Extinction correction: none

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 $F^2 > \sigma(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
Cu1	-0.00815 (3)	0.79189 (3)	0.40050 (2)	0.02480 (9)
Cl1	0.21525 (6)	1.15056 (7)	0.48221 (5)	0.02695 (12)
S1	0.25729 (6)	0.72960 (7)	0.64825 (6)	0.02540 (12)
O1	0.09412 (19)	0.7556 (2)	0.59265 (16)	0.0311 (3)
O2	0.4160 (2)	0.9013 (2)	0.72554 (19)	0.0385 (4)
O3	0.2140 (2)	0.6050 (2)	0.74013 (19)	0.0381 (4)
N1	-0.1223 (2)	0.7825 (2)	0.18742 (18)	0.0257 (3)
N2	0.1380 (2)	0.6952 (2)	0.2771 (2)	0.0276 (3)
C1	-0.2789 (3)	0.7931 (4)	0.1445 (3)	0.0382 (5)
H1	-0.3426	0.8078	0.2188	0.046*
C2	-0.3480 (3)	0.7828 (4)	-0.0065 (3)	0.0443 (6)
H2	-0.4578	0.7878	-0.0340	0.053*

supplementary materials

C3	-0.2519 (3)	0.7652 (3)	-0.1151 (2)	0.0391 (5)
H3	-0.2961	0.7581	-0.2174	0.047*
C4	-0.0881 (3)	0.7580 (3)	-0.0712 (2)	0.0332 (5)
H4	-0.0203	0.7481	-0.1430	0.040*
C5	-0.0275 (3)	0.7659 (3)	0.0813 (2)	0.0264 (4)
C6	0.1491 (3)	0.7631 (3)	0.1417 (2)	0.0314 (4)
H6A	0.1726	0.6835	0.0600	0.038*
H6B	0.2478	0.8857	0.1731	0.038*
C7	0.3187 (3)	0.7178 (3)	0.3579 (2)	0.0287 (4)
H7A	0.3988	0.8472	0.4099	0.034*
H7B	0.3725	0.6699	0.2812	0.034*
C8	0.2998 (3)	0.6170 (3)	0.4776 (2)	0.0296 (4)
H8A	0.4104	0.6049	0.5074	0.035*
H8B	0.2011	0.4944	0.4297	0.035*
H1N	0.051 (4)	0.570 (4)	0.233 (3)	0.045 (7)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cu1	0.02148 (14)	0.03819 (16)	0.02055 (13)	0.01626 (11)	0.00702 (9)	0.01323 (10)
Cl1	0.0222 (2)	0.0363 (3)	0.0269 (2)	0.01426 (19)	0.01029 (18)	0.01360 (18)
S1	0.0178 (2)	0.0297 (3)	0.0285 (2)	0.00845 (19)	0.00306 (18)	0.01438 (19)
O1	0.0270 (7)	0.0504 (9)	0.0255 (7)	0.0230 (7)	0.0082 (6)	0.0174 (6)
O2	0.0268 (8)	0.0346 (8)	0.0401 (8)	0.0044 (6)	0.0010 (6)	0.0087 (6)
O3	0.0324 (8)	0.0455 (9)	0.0432 (9)	0.0154 (7)	0.0091 (7)	0.0296 (7)
N1	0.0243 (8)	0.0332 (9)	0.0211 (7)	0.0132 (7)	0.0053 (6)	0.0109 (6)
N2	0.0229 (8)	0.0328 (9)	0.0281 (8)	0.0125 (7)	0.0071 (7)	0.0112 (7)
C1	0.0324 (11)	0.0617 (15)	0.0289 (10)	0.0259 (11)	0.0085 (9)	0.0196 (10)
C2	0.0384 (13)	0.0680 (17)	0.0339 (11)	0.0295 (12)	0.0035 (10)	0.0217 (11)
C3	0.0485 (14)	0.0455 (13)	0.0227 (10)	0.0210 (11)	0.0021 (9)	0.0139 (9)
C4	0.0422 (12)	0.0353 (11)	0.0224 (9)	0.0166 (10)	0.0103 (9)	0.0096 (8)
C5	0.0294 (10)	0.0245 (9)	0.0236 (9)	0.0101 (8)	0.0073 (8)	0.0080 (7)
C6	0.0297 (10)	0.0429 (12)	0.0262 (9)	0.0174 (9)	0.0123 (8)	0.0139 (8)
C7	0.0204 (9)	0.0343 (11)	0.0319 (10)	0.0118 (8)	0.0072 (8)	0.0115 (8)
C8	0.0246 (10)	0.0280 (10)	0.0388 (11)	0.0132 (8)	0.0078 (8)	0.0130 (8)

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

Cu1—O1	1.9775 (14)	C1—H1	0.9300
Cu1—N1	2.0051 (16)	C2—C3	1.371 (4)
Cu1—N2	2.0268 (17)	C2—H2	0.9300
Cu1—Cl1 ⁱ	2.2901 (5)	C3—C4	1.390 (3)
Cu1—Cl1	2.6796 (7)	C3—H3	0.9300
Cl1—Cu1 ⁱ	2.2901 (5)	C4—C5	1.386 (3)
S1—O2	1.4380 (16)	C4—H4	0.9300
S1—O3	1.4568 (15)	C5—C6	1.501 (3)
S1—O1	1.4916 (14)	C6—H6A	0.9700
S1—C8	1.774 (2)	C6—H6B	0.9700

N1—C5	1.345 (3)	C7—C8	1.519 (3)
N1—C1	1.346 (3)	C7—H7A	0.9700
N2—C6	1.475 (3)	C7—H7B	0.9700
N2—C7	1.492 (2)	C8—H8A	0.9700
N2—H1N	0.95 (3)	C8—H8B	0.9700
C1—C2	1.382 (3)		
O1—Cu1—N1	169.95 (7)	C3—C2—C1	118.9 (2)
O1—Cu1—N2	92.73 (6)	C3—C2—H2	120.5
N1—Cu1—N2	81.18 (7)	C1—C2—H2	120.5
O1—Cu1—Cl1 ⁱ	89.03 (4)	C2—C3—C4	119.45 (19)
N1—Cu1—Cl1 ⁱ	95.57 (5)	C2—C3—H3	120.3
N2—Cu1—Cl1 ⁱ	169.68 (5)	C4—C3—H3	120.3
O1—Cu1—Cl1	96.00 (5)	C5—C4—C3	118.9 (2)
N1—Cu1—Cl1	92.76 (5)	C5—C4—H4	120.5
N2—Cu1—Cl1	97.82 (5)	C3—C4—H4	120.5
Cl1 ⁱ —Cu1—Cl1	92.091 (19)	N1—C5—C4	121.49 (19)
Cu1 ⁱ —Cl1—Cu1	87.910 (19)	N1—C5—C6	115.27 (16)
O2—S1—O3	114.61 (10)	C4—C5—C6	123.21 (18)
O2—S1—O1	112.23 (10)	N2—C6—C5	108.83 (16)
O3—S1—O1	109.38 (9)	N2—C6—H6A	109.9
O2—S1—C8	107.16 (10)	C5—C6—H6A	109.9
O3—S1—C8	106.96 (10)	N2—C6—H6B	109.9
O1—S1—C8	105.98 (9)	C5—C6—H6B	109.9
S1—O1—Cu1	134.45 (9)	H6A—C6—H6B	108.3
C5—N1—C1	119.10 (17)	N2—C7—C8	110.64 (16)
C5—N1—Cu1	114.53 (13)	N2—C7—H7A	109.5
C1—N1—Cu1	126.36 (14)	C8—C7—H7A	109.5
C6—N2—C7	111.18 (16)	N2—C7—H7B	109.5
C6—N2—Cu1	108.32 (13)	C8—C7—H7B	109.5
C7—N2—Cu1	120.32 (13)	H7A—C7—H7B	108.1
C6—N2—H1N	104.2 (16)	C7—C8—S1	113.05 (14)
C7—N2—H1N	112.8 (16)	C7—C8—H8A	109.0
Cu1—N2—H1N	98.3 (16)	S1—C8—H8A	109.0
N1—C1—C2	122.1 (2)	C7—C8—H8B	109.0
N1—C1—H1	119.0	S1—C8—H8B	109.0
C2—C1—H1	119.0	H8A—C8—H8B	107.8
O1—Cu1—Cl1—Cu1 ⁱ	-89.25 (4)	N1—Cu1—N2—C7	158.15 (16)
N1—Cu1—Cl1—Cu1 ⁱ	95.68 (5)	Cl1 ⁱ —Cu1—N2—C7	-129.6 (2)
N2—Cu1—Cl1—Cu1 ⁱ	177.15 (5)	Cl1—Cu1—N2—C7	66.55 (14)
Cl1 ⁱ —Cu1—Cl1—Cu1 ⁱ	0.0	C5—N1—C1—C2	1.6 (4)
O2—S1—O1—Cu1	86.40 (15)	Cu1—N1—C1—C2	-179.12 (19)
O3—S1—O1—Cu1	-145.24 (13)	N1—C1—C2—C3	-1.4 (4)
C8—S1—O1—Cu1	-30.26 (16)	C1—C2—C3—C4	0.0 (4)
N1—Cu1—O1—S1	70.6 (4)	C2—C3—C4—C5	1.1 (4)
N2—Cu1—O1—S1	18.24 (15)	C1—N1—C5—C4	-0.5 (3)
Cl1 ⁱ —Cu1—O1—S1	-171.93 (14)	Cu1—N1—C5—C4	-179.82 (16)

supplementary materials

C11—Cu1—O1—S1	-79.93 (14)	C1—N1—C5—C6	177.54 (19)
O1—Cu1—N1—C5	-68.7 (4)	Cu1—N1—C5—C6	-1.8 (2)
N2—Cu1—N1—C5	-15.48 (14)	C3—C4—C5—N1	-0.9 (3)
C11 ⁱ —Cu1—N1—C5	174.39 (13)	C3—C4—C5—C6	-178.7 (2)
C11—Cu1—N1—C5	82.02 (14)	C7—N2—C6—C5	-170.83 (17)
O1—Cu1—N1—C1	112.1 (4)	Cu1—N2—C6—C5	-36.51 (19)
N2—Cu1—N1—C1	165.2 (2)	N1—C5—C6—N2	25.8 (2)
C11 ⁱ —Cu1—N1—C1	-4.89 (19)	C4—C5—C6—N2	-156.21 (19)
C11—Cu1—N1—C1	-97.27 (19)	C6—N2—C7—C8	-169.70 (17)
O1—Cu1—N2—C6	-159.28 (14)	Cu1—N2—C7—C8	62.2 (2)
N1—Cu1—N2—C6	28.76 (14)	N2—C7—C8—S1	-74.07 (19)
C11 ⁱ —Cu1—N2—C6	101.0 (3)	O2—S1—C8—C7	-65.44 (17)
C11—Cu1—N2—C6	-62.84 (13)	O3—S1—C8—C7	171.21 (14)
O1—Cu1—N2—C7	-29.89 (15)	O1—S1—C8—C7	54.58 (17)

Symmetry codes: (i) $-x, -y+2, -z+1$.

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2—H1N \cdots O3 ⁱⁱ	0.95 (3)	2.20 (3)	2.966 (2)	137 (2)

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

Fig. 1

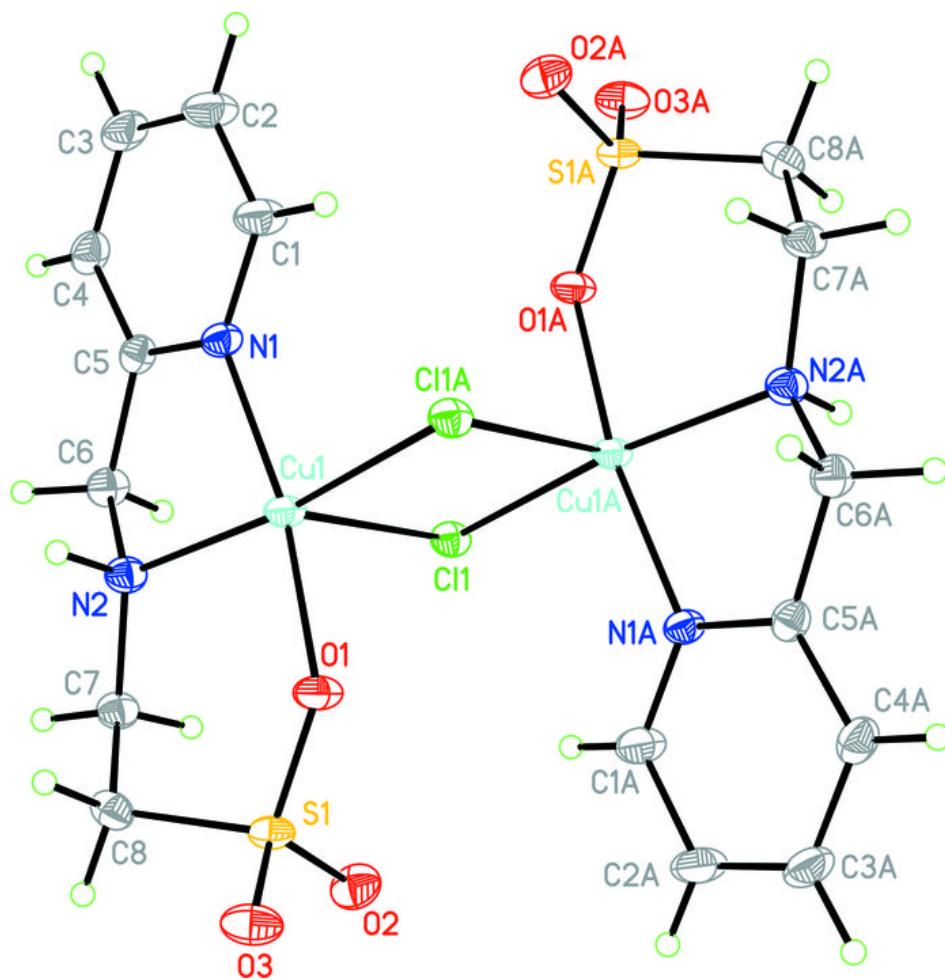


Fig. 2

