

cis-Bis(butylamine- κ N)bis[sulfadiazine(1-)- κ^2 N,N']copper(II) pentahydrate

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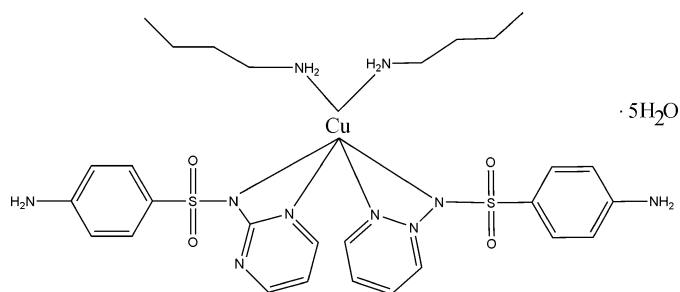
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; H-atom completeness 81%; disorder in main residue; R factor = 0.048; wR factor = 0.141; data-to-parameter ratio = 18.1.

In the title compound [systematic name: *cis*-bis[4-amino-*N*-(pyrimidin-2-yl)benzenesulfonamido- κ^2 N,N']bis(butylamine- κ N)copper(II) pentahydrate], $[\text{Cu}(\text{C}_{10}\text{H}_9\text{N}_4\text{O}_2\text{S})_2(\text{C}_4\text{H}_{11}\text{N})_2] \cdot 5\text{H}_2\text{O}$ or $[\text{Cu}(\text{sdz})_2(\text{ba})_2] \cdot 5\text{H}_2\text{O}$ [ba is butylamine and sdz = sulfadiazine(1-)], the copper(II) cation is six-coordinated by four N atoms of two sulfadiazine ligands and two N atoms of butylamine ligands. The copper(II) ion and one of the water molecules lie on twofold rotation axes. One of the butyl groups is disordered over two sites, with occupancies of 0.395 (8) and 0.605 (8). The geometry around the S atom is distorted tetrahedral. The crystal structure involves intermolecular $\text{N}-\text{H} \cdots \text{N}$ and $\text{N}-\text{H} \cdots \text{O}$ hydrogen bonds. $\text{N}-\text{H} \cdots \text{N}$ hydrogen bonds between sdz ligands lead to a sheet structure parallel to the *ab* plane.

Related literature

For related structures, see: Heren *et al.* (2006); Chung *et al.* (1975).



Experimental

Crystal data

$[\text{Cu}(\text{C}_{10}\text{H}_9\text{N}_4\text{O}_2\text{S})_2(\text{C}_4\text{H}_{11}\text{N})_2] \cdot 5\text{H}_2\text{O}$
 $M_r = 790.04$
 Orthorhombic, *Pbcn*
 $a = 22.623$ (6) Å
 $b = 10.342$ (5) Å
 $c = 16.250$ (6) Å
 $V = 3802$ (2) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.74$ mm⁻¹
 $T = 296$ K
 $0.34 \times 0.21 \times 0.19$ mm

Data collection

Stoe IPDS2 diffractometer
 Absorption correction: integration (*X-RED32*; Stoe & Cie, 2002)
 $T_{\min} = 0.821$, $T_{\max} = 0.899$
 57960 measured reflections
 4235 independent reflections
 2098 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.085$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.141$
 $S = 0.90$
 4235 reflections
 234 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.65$ e Å⁻³
 $\Delta\rho_{\min} = -0.45$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| <i>D</i> — <i>H</i> ··· <i>A</i> | <i>D</i> — <i>H</i> | <i>H</i> ··· <i>A</i> | <i>D</i> ··· <i>A</i> | <i>D</i> — <i>H</i> ··· <i>A</i> |
|----------------------------------|---------------------|-----------------------|-----------------------|----------------------------------|
| N5—H5A···N2 ⁱ | 0.86 | 2.53 | 3.359 (5) | 162 |
| N4—H4A···O5 ⁱⁱ | 0.90 | 2.45 | 3.337 (6) | 171 |
| N4—H4B···O4 ⁱⁱⁱ | 0.90 | 2.25 | 3.119 (6) | 161 |
| N5—H5B···O5 ^{iv} | 0.86 | 2.26 | 3.113 (5) | 170 |

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

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CF2216).

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supplementary materials

Acta Cryst. (2008). E64, m1192 [doi:10.1107/S1600536808026457]

cis-Bis(butylamine- κN)bis[sulfadiazine(1-)- $\kappa^2 N, N'$]copper(II) pentahydrate

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Comment

In the title complex (I), the copper(II) ion is six-coordinated by four N atoms of sulfadiazine ligands and two N atoms of butylammonium ligands. The copper(II) ion and one of the water molecules lie on twofold rotation axes. It is found that the Cu–N_{sdz} and Cu–N_{ba} bond distances are nearly equal. The bond angles around the S atom correspond to a distorted tetrahedral geometry. The C4–N5 bond distance and the torsion angle C1–S1–N1–C7 are comparable to those observed in related structures (Heren *et al.*, 2006; Chung *et al.*, 1975). One of the butyl groups is disordered over two sites with occupancies of 0.395 (8):0.605 (8) (see Fig. 1).

The packing of (I) is stabilized by intermolecular N—H \cdots N and N—H \cdots O hydrogen bonds (Table 1). The N—H \cdots N hydrogen bond takes place between sdz ligands and it is seen that these hydrogen bonds generate a sheet structure parallel to the *ab* plane (Fig. 2). The H atoms of water molecules could not be located from a Fourier map. However, it is possible to see that water molecules are involved in hydrogen bonds with sdz and ba ligands on the basis of interatomic distances.

Experimental

A solution of butylamine (2 mmol) in ethanol (20 ml) was added dropwise with stirring to a solution of Cu(II) sulfadiazine (1 mmol) in methanol (40 ml). The solution was heated and stirred for 3 h at 343 K and then the mixture was cooled to room temperature. The blue crystals were filtered off, washed with cold distilled water and acetone, and dried *in vacuo*. Analysis calculated: C 42.62, H 5.84, N 17.76%; found: C 43.08, H 5.72, N 18.25%.

Refinement

One butyl group shows disorder and was modelled with two different orientations and site occupancies of 0.395 (8):0.605 (8). The H atoms of water molecules could not be located from a Fourier map. All other H atoms were placed in geometrically idealized positions with distances N—H = 0.86–0.90 Å, C—H = 0.93–0.97 Å, and were refined as riding atoms with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ and $U_{\text{iso}}(\text{H}_{\text{methyl}}) = 1.5U_{\text{eq}}(\text{C})$.

Figures

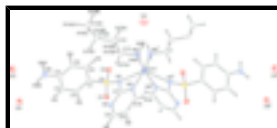


Fig. 1. The molecular structure of (I), with atom labels and 30% probability displacement ellipsoids. Only the major part of the disordered ba ligand is included. [Symmetry code: (i) $-x + 1, y, -z + 1/2$.]

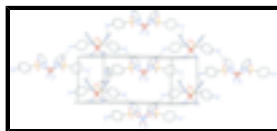


Fig. 2. A view of the complex showing the sheet structure parallel to the *ab* plane. The butyl groups, water molecules and some hydrogen bonds have been omitted for clarity. Other hydrogen bonds are shown as dashed lines. [Symmetry code: (i) $-x + 1/2, y - 1/2, z$.]

supplementary materials

cis-bis[4-amino-N-(pyrimidin-2-yl)benzenesulfonamido- κ^2 N,N']bis(butylamine- κ N)copper(II) pentahydrate}

Crystal data

| | |
|--|---|
| [Cu(C ₁₀ H ₉ N ₄ O ₂ S) ₂ (C ₄ H ₁₁ N) ₂] \cdot 5H ₂ O | $F_{000} = 1594$ |
| $M_r = 790.04$ | $D_x = 1.347 \text{ Mg m}^{-3}$ |
| Orthorhombic, <i>Pbcn</i> | Mo $K\alpha$ radiation |
| Hall symbol: -P 2n 2ab | $\lambda = 0.71069 \text{ \AA}$ |
| $a = 22.623 (6) \text{ \AA}$ | Cell parameters from 30097 reflections |
| $b = 10.342 (5) \text{ \AA}$ | $\theta = 1.8\text{--}27.1^\circ$ |
| $c = 16.250 (6) \text{ \AA}$ | $\mu = 0.74 \text{ mm}^{-1}$ |
| $V = 3802 (2) \text{ \AA}^3$ | $T = 296 \text{ K}$ |
| $Z = 4$ | Prism, blue |
| | $0.34 \times 0.21 \times 0.19 \text{ mm}$ |

Data collection

| | |
|--|--|
| Stoe IPDS2 diffractometer | 4235 independent reflections |
| Radiation source: fine-focus sealed tube | 2098 reflections with $I > 2\sigma(I)$ |
| Monochromator: graphite | $R_{\text{int}} = 0.085$ |
| $T = 296 \text{ K}$ | $\theta_{\text{max}} = 27.2^\circ$ |
| rotation method scans | $\theta_{\text{min}} = 1.8^\circ$ |
| Absorption correction: integration (X-RED32; Stoe & Cie, 2002) | $h = -28 \rightarrow 28$ |
| $T_{\text{min}} = 0.821$, $T_{\text{max}} = 0.900$ | $k = -13 \rightarrow 13$ |
| 57960 measured reflections | $l = -20 \rightarrow 20$ |

Refinement

| | |
|--|--|
| Refinement on F^2 | Secondary atom site location: difference Fourier map |
| Least-squares matrix: full | Hydrogen site location: inferred from neighbouring sites |
| $R[F^2 > 2\sigma(F^2)] = 0.048$ | H-atom parameters constrained |
| $wR(F^2) = 0.141$ | $w = 1/[\sigma^2(F_o^2) + (0.0787P)^2]$ |
| $S = 0.90$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| 4235 reflections | $(\Delta/\sigma)_{\text{max}} < 0.001$ |
| 234 parameters | $\Delta\rho_{\text{max}} = 0.65 \text{ e \AA}^{-3}$ |
| Primary atom site location: structure-invariant direct methods | $\Delta\rho_{\text{min}} = -0.45 \text{ e \AA}^{-3}$ |
| | 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)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ | Occ. (<1) |
|------|--------------|-------------|--------------|----------------------------------|-----------|
| C1 | 0.33524 (14) | 0.7316 (3) | 0.45323 (19) | 0.0565 (8) | |
| C2 | 0.29637 (16) | 0.8185 (4) | 0.4879 (2) | 0.0684 (9) | |
| H2 | 0.3109 | 0.8929 | 0.5129 | 0.082* | |
| C3 | 0.23634 (16) | 0.7970 (4) | 0.4860 (2) | 0.0752 (10) | |
| H3 | 0.2108 | 0.8559 | 0.5107 | 0.090* | |
| C4 | 0.21358 (15) | 0.6879 (4) | 0.4473 (2) | 0.0689 (9) | |
| C5 | 0.25293 (17) | 0.6011 (4) | 0.4128 (2) | 0.0728 (10) | |
| H5 | 0.2387 | 0.5271 | 0.3870 | 0.087* | |
| C6 | 0.31255 (15) | 0.6227 (3) | 0.4160 (2) | 0.0664 (9) | |
| H6 | 0.3382 | 0.5628 | 0.3927 | 0.080* | |
| C7 | 0.41743 (13) | 0.8242 (3) | 0.3032 (2) | 0.0565 (8) | |
| C8 | 0.36943 (17) | 0.9966 (4) | 0.2474 (3) | 0.0800 (11) | |
| H8 | 0.3452 | 1.0686 | 0.2536 | 0.096* | |
| C9 | 0.39032 (17) | 0.9691 (4) | 0.1702 (3) | 0.0847 (12) | |
| H9 | 0.3802 | 1.0190 | 0.1247 | 0.102* | |
| C10 | 0.42700 (16) | 0.8638 (4) | 0.1637 (2) | 0.0745 (10) | |
| H10 | 0.4427 | 0.8426 | 0.1125 | 0.089* | |
| C11 | 0.3906 (2) | 0.4981 (5) | 0.2216 (4) | 0.124 (2) | |
| H11A | 0.3706 | 0.5801 | 0.2136 | 0.149* | |
| H11B | 0.3925 | 0.4841 | 0.2806 | 0.149* | |
| C12 | 0.3518 (2) | 0.4000 (5) | 0.1892 (4) | 0.1116 (17) | |
| H12A | 0.3740 | 0.3199 | 0.1854 | 0.134* | |
| H12B | 0.3409 | 0.4246 | 0.1337 | 0.134* | |
| C13A | 0.2962 (4) | 0.3733 (8) | 0.2362 (5) | 0.088 (2) | 0.605 (8) |
| H13A | 0.2764 | 0.4549 | 0.2461 | 0.105* | 0.605 (8) |
| H13B | 0.3070 | 0.3376 | 0.2893 | 0.105* | 0.605 (8) |
| C13B | 0.2848 (5) | 0.4111 (13) | 0.1823 (9) | 0.088 (2) | 0.395 (8) |
| H13C | 0.2709 | 0.4851 | 0.2134 | 0.105* | 0.395 (8) |
| H13D | 0.2733 | 0.4223 | 0.1253 | 0.105* | 0.395 (8) |
| C14A | 0.2505 (12) | 0.277 (3) | 0.1934 (18) | 0.108 (4) | 0.605 (8) |
| H14A | 0.2169 | 0.2658 | 0.2286 | 0.162* | 0.605 (8) |
| H14B | 0.2692 | 0.1954 | 0.1841 | 0.162* | 0.605 (8) |
| H14C | 0.2380 | 0.3132 | 0.1418 | 0.162* | 0.605 (8) |
| C14B | 0.261 (2) | 0.299 (4) | 0.213 (3) | 0.108 (4) | 0.395 (8) |
| H14D | 0.2186 | 0.3043 | 0.2115 | 0.162* | 0.395 (8) |
| H14E | 0.2736 | 0.2879 | 0.2693 | 0.162* | 0.395 (8) |
| H14F | 0.2741 | 0.2270 | 0.1810 | 0.162* | 0.395 (8) |
| N1 | 0.43499 (11) | 0.7422 (3) | 0.36300 (16) | 0.0586 (7) | |

supplementary materials

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|-----|--------------|-------------|--------------|-------------|
| N2 | 0.38163 (12) | 0.9265 (3) | 0.31428 (17) | 0.0635 (7) |
| N3 | 0.44067 (12) | 0.7910 (3) | 0.22883 (16) | 0.0596 (7) |
| N4 | 0.44912 (14) | 0.5158 (3) | 0.1947 (2) | 0.0896 (10) |
| H4A | 0.4478 | 0.5356 | 0.1408 | 0.108* |
| H4B | 0.4677 | 0.4391 | 0.1990 | 0.108* |
| N5 | 0.15418 (15) | 0.6661 (4) | 0.4447 (2) | 0.1001 (11) |
| H5A | 0.1407 | 0.5976 | 0.4213 | 0.120* |
| H5B | 0.1302 | 0.7209 | 0.4664 | 0.120* |
| O1 | 0.42208 (10) | 0.8844 (2) | 0.48919 (14) | 0.0690 (6) |
| O2 | 0.43907 (10) | 0.6530 (2) | 0.49927 (15) | 0.0731 (7) |
| O3 | 0.02149 (14) | 0.6126 (3) | 0.4017 (2) | 0.1193 (11) |
| O4 | 0.0000 | 0.7493 (5) | 0.2500 | 0.151 (2) |
| O5 | 0.43200 (14) | 0.3833 (3) | 0.5011 (3) | 0.1410 (15) |
| S2 | 0.41156 (4) | 0.75774 (8) | 0.45398 (5) | 0.0588 (2) |
| Cu1 | 0.5000 | 0.64823 (6) | 0.2500 | 0.0647 (2) |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C1 | 0.0482 (17) | 0.062 (2) | 0.0590 (17) | -0.0025 (16) | 0.0034 (15) | 0.0033 (16) |
| C2 | 0.056 (2) | 0.068 (2) | 0.081 (2) | -0.0050 (18) | 0.0009 (18) | -0.0092 (19) |
| C3 | 0.053 (2) | 0.080 (2) | 0.093 (3) | 0.0044 (19) | 0.0133 (19) | -0.005 (2) |
| C4 | 0.051 (2) | 0.078 (2) | 0.078 (2) | -0.0120 (19) | -0.0005 (17) | 0.007 (2) |
| C5 | 0.059 (2) | 0.077 (2) | 0.083 (2) | -0.013 (2) | -0.0008 (19) | -0.005 (2) |
| C6 | 0.054 (2) | 0.069 (2) | 0.076 (2) | -0.0018 (17) | 0.0045 (17) | -0.0079 (18) |
| C7 | 0.0426 (17) | 0.064 (2) | 0.063 (2) | -0.0072 (16) | 0.0046 (15) | 0.0020 (16) |
| C8 | 0.067 (2) | 0.084 (3) | 0.088 (3) | 0.012 (2) | -0.003 (2) | 0.020 (2) |
| C9 | 0.068 (2) | 0.105 (3) | 0.081 (3) | 0.004 (2) | -0.003 (2) | 0.028 (2) |
| C10 | 0.057 (2) | 0.100 (3) | 0.066 (2) | -0.001 (2) | 0.0069 (17) | 0.007 (2) |
| C11 | 0.089 (3) | 0.096 (3) | 0.187 (6) | -0.029 (3) | 0.050 (3) | -0.047 (4) |
| C12 | 0.080 (3) | 0.108 (4) | 0.146 (4) | -0.006 (3) | 0.002 (3) | -0.049 (3) |
| C13A | 0.079 (4) | 0.092 (5) | 0.092 (5) | -0.012 (4) | 0.002 (4) | -0.018 (4) |
| C13B | 0.079 (4) | 0.092 (5) | 0.092 (5) | -0.012 (4) | 0.002 (4) | -0.018 (4) |
| C14A | 0.081 (11) | 0.103 (9) | 0.140 (15) | -0.017 (6) | -0.014 (8) | -0.034 (8) |
| C14B | 0.081 (11) | 0.103 (9) | 0.140 (15) | -0.017 (6) | -0.014 (8) | -0.034 (8) |
| N1 | 0.0480 (15) | 0.0615 (16) | 0.0664 (16) | 0.0056 (13) | 0.0059 (12) | 0.0030 (14) |
| N2 | 0.0588 (17) | 0.0636 (17) | 0.0679 (17) | 0.0086 (14) | 0.0022 (14) | 0.0019 (15) |
| N3 | 0.0506 (16) | 0.0697 (17) | 0.0585 (16) | -0.0061 (14) | 0.0041 (12) | -0.0026 (14) |
| N4 | 0.070 (2) | 0.087 (2) | 0.112 (3) | 0.0022 (18) | 0.0034 (19) | -0.021 (2) |
| N5 | 0.0505 (18) | 0.110 (3) | 0.140 (3) | -0.0131 (19) | 0.004 (2) | -0.009 (2) |
| O1 | 0.0574 (14) | 0.0741 (15) | 0.0755 (14) | -0.0064 (12) | -0.0044 (12) | -0.0138 (12) |
| O2 | 0.0606 (15) | 0.0799 (16) | 0.0787 (15) | 0.0053 (13) | -0.0093 (12) | 0.0243 (13) |
| O3 | 0.079 (2) | 0.125 (3) | 0.155 (3) | 0.0017 (19) | 0.018 (2) | -0.008 (2) |
| O4 | 0.174 (6) | 0.091 (3) | 0.188 (6) | 0.000 | 0.053 (4) | 0.000 |
| O5 | 0.073 (2) | 0.096 (2) | 0.253 (5) | 0.0007 (18) | -0.015 (3) | -0.001 (3) |
| S2 | 0.0469 (4) | 0.0667 (5) | 0.0629 (5) | -0.0008 (4) | -0.0020 (4) | 0.0035 (4) |
| Cu1 | 0.0516 (3) | 0.0697 (4) | 0.0726 (4) | 0.000 | 0.0051 (3) | 0.000 |

Geometric parameters (Å, °)

| | | | |
|----------|-----------|---------------------|------------|
| C1—C6 | 1.377 (5) | C12—C13B | 1.524 (13) |
| C1—C2 | 1.378 (5) | C12—H12A | 0.970 |
| C1—S2 | 1.748 (3) | C12—H12B | 0.970 |
| C2—C3 | 1.377 (5) | C13A—C14A | 1.592 (16) |
| C2—H2 | 0.930 | C13A—H13A | 0.970 |
| C3—C4 | 1.391 (5) | C13A—H13B | 0.970 |
| C3—H3 | 0.930 | C13B—C14B | 1.37 (5) |
| C4—N5 | 1.363 (5) | C13B—H13C | 0.970 |
| C4—C5 | 1.382 (5) | C13B—H13D | 0.970 |
| C5—C6 | 1.368 (5) | C14A—H14A | 0.960 |
| C5—H5 | 0.930 | C14A—H14B | 0.960 |
| C6—H6 | 0.930 | C14A—H14C | 0.960 |
| C7—N2 | 1.344 (4) | C14B—H14D | 0.960 |
| C7—N1 | 1.351 (4) | C14B—H14E | 0.960 |
| C7—N3 | 1.361 (4) | C14B—H14F | 0.960 |
| C8—N2 | 1.334 (5) | N1—S2 | 1.579 (3) |
| C8—C9 | 1.371 (6) | N3—Cu1 | 2.025 (3) |
| C8—H8 | 0.930 | N4—Cu1 | 2.002 (3) |
| C9—C10 | 1.373 (5) | N4—H4A | 0.900 |
| C9—H9 | 0.930 | N4—H4B | 0.900 |
| C10—N3 | 1.336 (4) | N5—H5A | 0.860 |
| C10—H10 | 0.930 | N5—H5B | 0.860 |
| C11—N4 | 1.407 (5) | O1—S2 | 1.449 (2) |
| C11—C12 | 1.441 (6) | O2—S2 | 1.450 (2) |
| C11—H11A | 0.970 | Cu1—N4 ⁱ | 2.002 (3) |
| C11—H11B | 0.970 | Cu1—N3 ⁱ | 2.025 (3) |
| C12—C13A | 1.496 (9) | | |
| C6—C1—C2 | 118.4 (3) | C12—C13A—H13B | 108.3 |
| C6—C1—S2 | 119.9 (3) | C14A—C13A—H13B | 108.3 |
| C2—C1—S2 | 121.8 (3) | H13A—C13A—H13B | 107.4 |
| C3—C2—C1 | 121.0 (3) | C14B—C13B—C12 | 107 (2) |
| C3—C2—H2 | 119.5 | C14B—C13B—H13C | 110.2 |
| C1—C2—H2 | 119.5 | C12—C13B—H13C | 110.2 |
| C2—C3—C4 | 120.4 (4) | C14B—C13B—H13D | 110.2 |
| C2—C3—H3 | 119.8 | C12—C13B—H13D | 110.2 |
| C4—C3—H3 | 119.8 | H13C—C13B—H13D | 108.5 |
| N5—C4—C5 | 121.0 (4) | C13A—C14A—H14A | 109.5 |
| N5—C4—C3 | 120.8 (4) | C13A—C14A—H14B | 109.5 |
| C5—C4—C3 | 118.1 (3) | H14A—C14A—H14B | 109.5 |
| C6—C5—C4 | 120.9 (4) | C13A—C14A—H14C | 109.5 |
| C6—C5—H5 | 119.6 | H14A—C14A—H14C | 109.5 |
| C4—C5—H5 | 119.6 | H14B—C14A—H14C | 109.5 |
| C5—C6—C1 | 121.2 (3) | C13B—C14B—H14D | 109.5 |
| C5—C6—H6 | 119.4 | C13B—C14B—H14E | 109.5 |
| C1—C6—H6 | 119.4 | H14D—C14B—H14E | 109.5 |

supplementary materials

| | | | |
|--------------------|------------|--------------------------------------|-------------|
| N2—C7—N1 | 125.1 (3) | C13B—C14B—H14F | 109.5 |
| N2—C7—N3 | 123.4 (3) | H14D—C14B—H14F | 109.5 |
| N1—C7—N3 | 111.5 (3) | H14E—C14B—H14F | 109.5 |
| N2—C8—C9 | 124.2 (4) | C7—N1—S2 | 120.8 (2) |
| N2—C8—H8 | 117.9 | C8—N2—C7 | 116.2 (3) |
| C9—C8—H8 | 117.9 | C10—N3—C7 | 118.1 (3) |
| C8—C9—C10 | 116.3 (4) | C10—N3—Cu1 | 134.4 (2) |
| C8—C9—H9 | 121.8 | C7—N3—Cu1 | 106.8 (2) |
| C10—C9—H9 | 121.8 | C11—N4—Cu1 | 119.4 (3) |
| N3—C10—C9 | 121.7 (4) | C11—N4—H4A | 107.5 |
| N3—C10—H10 | 119.1 | Cu1—N4—H4A | 107.5 |
| C9—C10—H10 | 119.1 | C11—N4—H4B | 107.5 |
| N4—C11—C12 | 123.4 (4) | Cu1—N4—H4B | 107.5 |
| N4—C11—H11A | 106.5 | H4A—N4—H4B | 107.0 |
| C12—C11—H11A | 106.5 | C4—N5—H5A | 120.0 |
| N4—C11—H11B | 106.5 | C4—N5—H5B | 120.0 |
| C12—C11—H11B | 106.5 | H5A—N5—H5B | 120.0 |
| H11A—C11—H11B | 106.5 | O1—S2—O2 | 113.88 (15) |
| C11—C12—C13A | 117.0 (5) | O1—S2—N1 | 113.99 (15) |
| C11—C12—C13B | 125.3 (6) | O2—S2—N1 | 104.77 (15) |
| C11—C12—H12A | 108.0 | O1—S2—C1 | 107.76 (15) |
| C13A—C12—H12A | 108.0 | O2—S2—C1 | 108.16 (15) |
| C13B—C12—H12A | 125.1 | N1—S2—C1 | 108.02 (15) |
| C11—C12—H12B | 108.0 | N4—Cu1—N4 ⁱ | 93.7 (2) |
| C13A—C12—H12B | 108.0 | N4—Cu1—N3 ⁱ | 162.98 (13) |
| C13B—C12—H12B | 70.1 | N4 ⁱ —Cu1—N3 ⁱ | 92.38 (13) |
| H12A—C12—H12B | 107.3 | N4—Cu1—N3 | 92.38 (13) |
| C12—C13A—C14A | 116.0 (14) | N4 ⁱ —Cu1—N3 | 162.98 (13) |
| C12—C13A—H13A | 108.3 | N3 ⁱ —Cu1—N3 | 86.36 (16) |
| C14A—C13A—H13A | 108.3 | | |
| C6—C1—C2—C3 | 0.5 (5) | C9—C10—N3—Cu1 | -169.6 (3) |
| S2—C1—C2—C3 | 179.3 (3) | N2—C7—N3—C10 | -0.2 (5) |
| C1—C2—C3—C4 | -1.4 (6) | N1—C7—N3—C10 | 179.9 (3) |
| C2—C3—C4—N5 | -179.6 (4) | N2—C7—N3—Cu1 | 171.8 (2) |
| C2—C3—C4—C5 | 1.4 (6) | N1—C7—N3—Cu1 | -8.1 (3) |
| N5—C4—C5—C6 | -179.5 (4) | C12—C11—N4—Cu1 | -176.0 (5) |
| C3—C4—C5—C6 | -0.5 (6) | C7—N1—S2—O1 | -55.6 (3) |
| C4—C5—C6—C1 | -0.4 (6) | C7—N1—S2—O2 | 179.3 (2) |
| C2—C1—C6—C5 | 0.4 (5) | C7—N1—S2—C1 | 64.1 (3) |
| S2—C1—C6—C5 | -178.4 (3) | C6—C1—S2—O1 | 174.2 (3) |
| N2—C8—C9—C10 | -1.4 (6) | C2—C1—S2—O1 | -4.6 (3) |
| C8—C9—C10—N3 | 1.1 (6) | C6—C1—S2—O2 | -62.3 (3) |
| N4—C11—C12—C13A | 166.6 (7) | C2—C1—S2—O2 | 118.9 (3) |
| N4—C11—C12—C13B | -149.2 (9) | C6—C1—S2—N1 | 50.6 (3) |
| C11—C12—C13A—C14A | 172.7 (14) | C2—C1—S2—N1 | -128.2 (3) |
| C13B—C12—C13A—C14A | 59.0 (16) | C11—N4—Cu1—N4 ⁱ | 108.9 (5) |
| C11—C12—C13B—C14B | -132 (2) | C11—N4—Cu1—N3 ⁱ | -140.5 (5) |

| | | | |
|--------------------|------------|----------------------------|-----------|
| C13A—C12—C13B—C14B | -42 (2) | C11—N4—Cu1—N3 | -55.2 (4) |
| N2—C7—N1—S2 | 2.7 (4) | C10—N3—Cu1—N4 | -79.3 (4) |
| N3—C7—N1—S2 | -177.4 (2) | C7—N3—Cu1—N4 | 110.6 (2) |
| C9—C8—N2—C7 | 0.9 (6) | C10—N3—Cu1—N4 ⁱ | 169.9 (4) |
| N1—C7—N2—C8 | 179.8 (3) | C7—N3—Cu1—N4 ⁱ | -0.1 (5) |
| N3—C7—N2—C8 | -0.1 (5) | C10—N3—Cu1—N3 ⁱ | 83.7 (3) |
| C9—C10—N3—C7 | -0.4 (5) | C7—N3—Cu1—N3 ⁱ | -86.4 (2) |

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

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> |
|----------------------------|-------------|---------------|-----------------------|-------------------------|
| N5—H5A...N2 ⁱⁱ | 0.86 | 2.53 | 3.359 (5) | 162 |
| N4—H4A...O5 ⁱⁱⁱ | 0.90 | 2.45 | 3.337 (6) | 171 |
| N4—H4B...O4 ^{iv} | 0.90 | 2.25 | 3.119 (6) | 161 |
| N5—H5B...O5 ^v | 0.86 | 2.26 | 3.113 (5) | 170 |

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

Fig. 1

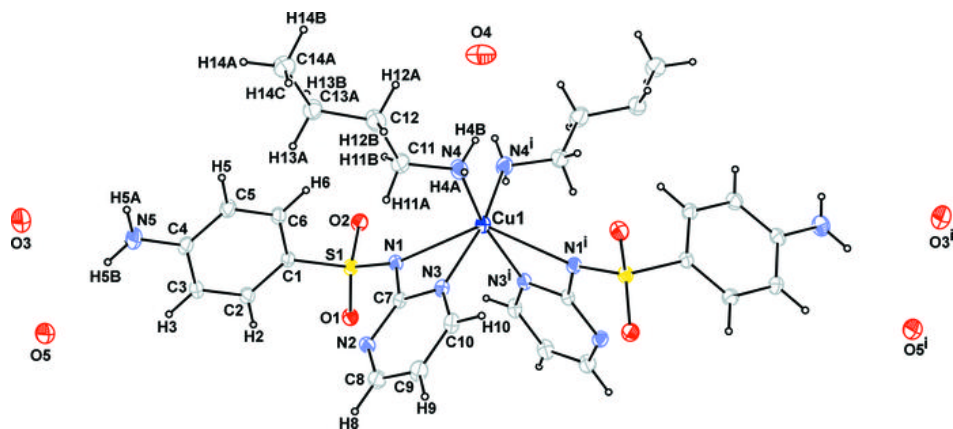


Fig. 2

