

{N,N-Dimethyl-N'-[1-(2-pyridyl)ethylidene]propane-1,3-diamine}bis(thiocyanato- κN)copper(II)

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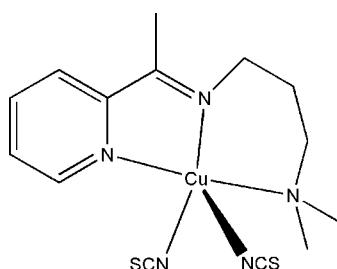
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Key indicators: single-crystal X-ray study; $T = 298\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$; disorder in main residue; R factor = 0.042; wR factor = 0.111; data-to-parameter ratio = 16.1.

In the title complex, $[\text{Cu}(\text{NCS})_2(\text{C}_{12}\text{H}_{19}\text{N}_3)]$, the Cu^{II} atom is five-coordinated in a square-pyramidal geometry defined by one pyridine N, one imine N, and one amine N atom of the tridentate Schiff base ligand and two N-bonded thiocyanate ions (one of the latter occupying the apical site). The three bridging C atoms and the two terminal C atoms of the Schiff base are disordered over two sets of sites, with occupancies of 0.465 (2) and 0.535 (2).

Related literature

For a related structure and background to Schiff bases, see: Xue *et al.* (2010).



Experimental

Crystal data

$[\text{Cu}(\text{NCS})_2(\text{C}_{12}\text{H}_{19}\text{N}_3)]$
 $M_r = 385.00$
Monoclinic, $P2_1/c$
 $a = 13.723$ (2) \AA
 $b = 7.2380$ (12) \AA
 $c = 18.237$ (3) \AA
 $\beta = 103.559$ (2) $^\circ$

$V = 1760.9$ (5) \AA^3
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 1.48\text{ mm}^{-1}$
 $T = 298\text{ K}$
 $0.23 \times 0.21 \times 0.21\text{ mm}$

Data collection

Bruker SMART CCD
diffractometer
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
 $T_{\min} = 0.727$, $T_{\max} = 0.746$

13886 measured reflections
3816 independent reflections
2698 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.111$
 $S = 1.04$
3816 reflections
237 parameters

16 restraints
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.57\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.47\text{ e \AA}^{-3}$

Table 1
Selected bond lengths (\AA).

| | | | |
|--------|-----------|--------|-----------|
| Cu1—N5 | 1.955 (3) | Cu1—N3 | 2.078 (3) |
| Cu1—N2 | 2.013 (3) | Cu1—N4 | 2.153 (3) |
| Cu1—N1 | 2.047 (3) | | |

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; 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*.

We thank the Top-Class Foundation and the Applied Chemistry Key Laboratory Foundation of Pingdingshan University.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5636).

References

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- Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Xue, L.-W., Zhao, G.-Q., Han, Y.-J. & Feng, Y.-X. (2010). *Acta Cryst. E* **66**, m1172–m1173.

supporting information

Acta Cryst. (2010). E66, m1274 [doi:10.1107/S1600536810036378]

{*N,N*-Dimethyl-*N'*-[1-(2-pyridyl)ethylidene]propane-1,3-diamine}bis(thiocyanato- κN)copper(II)

Ling-Wei Xue, Gan-Qing Zhao, Yong-Jun Han, Li-Hua Chen and Qin-Long Peng

S1. Comment

Recently, we have reported a copper(II) complex with a Schiff base ligand (Xue *et al.*, 2010). In this paper, a new thiocyanato-coordinated mononuclear copper(II) complex with the Schiff base *N,N*-dimethyl-*N'*-(1-pyridin-2-ylethylidene)propane-1,3-diamine, is reported.

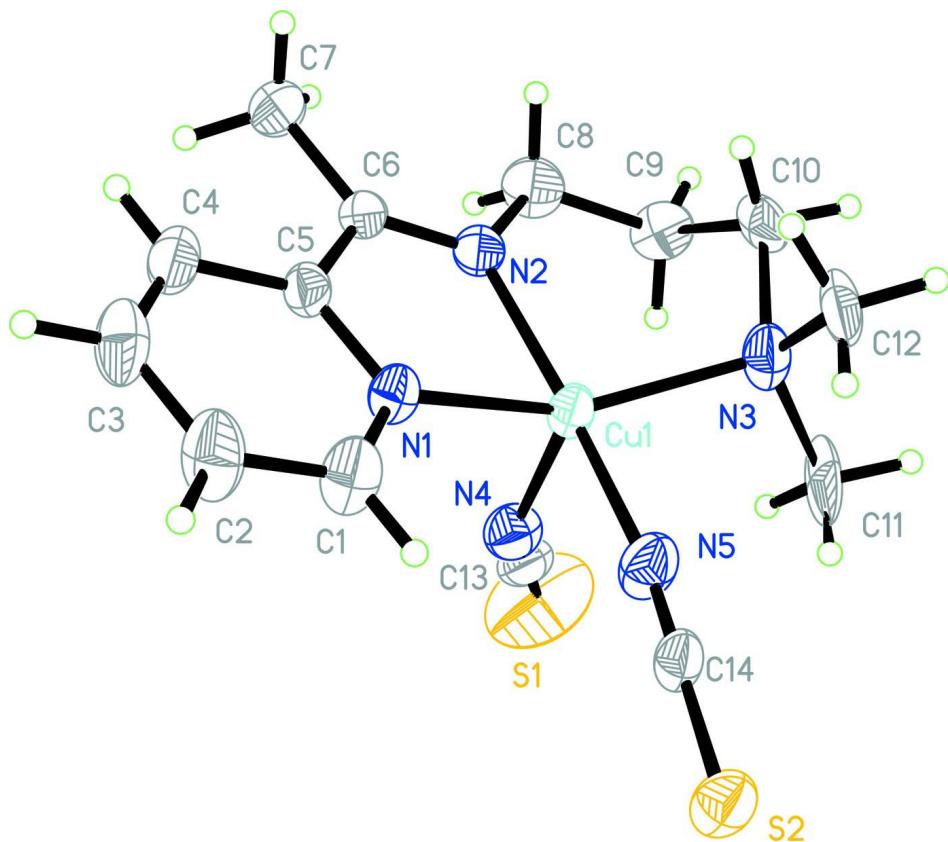
The Cu atom in the complex, Fig. 1, is five-coordinate in a square pyramidal geometry, with one pyridine N, one imine N, and one amine N atoms of a Schiff base ligand, and with one thiocyanate N atom, occupying the basal plane, and with another thiocyanate N atom occupying the apical position. The Cu atom displaced 0.306 (2) Å from the plane defined by the four basal donor atoms. The slight distortion of the square pyramidal coordination can be observed from the coordinate bond lengths and angles (Table 1).

S2. Experimental

2-Acetylpyridine (121 mg, 1.0 mmol), *N,N*dimethylpropane-1,3-diamine (102 mg, 1.0 mmol), ammonium thiocyanate (76 mg, 1.0 mmol), and copper acetate monohydrate (199.2 mg, 1.0 mmol) were dissolved in methanol (80 ml). The mixture was stirred for two hours at room temperature. The resulting solution was left in air for a few days, yielding blue blocks of (I).

S3. Refinement

H atoms were placed in idealized positions and constrained to ride on their parent atoms with C—H distances of 0.93–0.97 Å, and with $U_{\text{iso}}(\text{H})$ set at $1.2U_{\text{eq}}(\text{C})$ and $1.5U_{\text{eq}}(\text{C}_{\text{methyl}})$. The three bridging C atoms and the two terminal C atoms of the Schiff base ligand are disordered over two sites, with occupancies of 0.465 (2) and 0.535 (2).

**Figure 1**

The structure of the title complex with 30% probability displacement ellipsoids.

{N,N-Dimethyl-N'-[1-(2-pyridyl)ethylidene]propane-1,3- diamine}bis(thiocyanato- κ N)copper(II)

Crystal data

[Cu(NCS)₂(C₁₂H₁₉N₃)]
 $M_r = 385.00$
Monoclinic, $P2_1/c$
Hall symbol: -P 2ybc
 $a = 13.723 (2)$ Å
 $b = 7.2380 (12)$ Å
 $c = 18.237 (3)$ Å
 $\beta = 103.559 (2)$ °
 $V = 1760.9 (5)$ Å³
 $Z = 4$

$F(000) = 796$
 $D_x = 1.452 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 2747 reflections
 $\theta = 2.3\text{--}24.5$ °
 $\mu = 1.48 \text{ mm}^{-1}$
 $T = 298$ K
Block, blue
 $0.23 \times 0.21 \times 0.21$ mm

Data collection

Bruker SMART CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
 $T_{\min} = 0.727$, $T_{\max} = 0.746$

13886 measured reflections
3816 independent reflections
2698 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$
 $\theta_{\max} = 27.0$ °, $\theta_{\min} = 2.3$ °
 $h = -15 \rightarrow 17$
 $k = -9 \rightarrow 9$
 $l = -23 \rightarrow 23$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.111$
 $S = 1.04$
 3816 reflections
 237 parameters
 16 restraints
 Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map
 Hydrogen site location: inferred from neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0492P)^2 + 0.6842P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.57 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.47 \text{ e } \text{\AA}^{-3}$

Special details

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

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ 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) |
|-----|-------------|--------------|--------------|----------------------------------|------------|
| Cu1 | 0.24814 (3) | 0.82783 (5) | 0.39009 (2) | 0.04378 (15) | |
| N1 | 0.3853 (2) | 0.8651 (4) | 0.36614 (15) | 0.0459 (7) | |
| N2 | 0.2846 (2) | 1.0789 (4) | 0.43585 (14) | 0.0457 (6) | |
| N4 | 0.2763 (2) | 0.7053 (4) | 0.50065 (18) | 0.0594 (8) | |
| N5 | 0.2289 (2) | 0.5969 (5) | 0.33292 (18) | 0.0624 (8) | |
| S1 | 0.2253 (2) | 0.6298 (3) | 0.63328 (10) | 0.1725 (10) | |
| S2 | 0.23557 (7) | 0.28351 (14) | 0.24673 (6) | 0.0611 (3) | |
| C1 | 0.4337 (3) | 0.7542 (6) | 0.3285 (2) | 0.0647 (10) | |
| H1 | 0.4028 | 0.6446 | 0.3090 | 0.078* | |
| C2 | 0.5270 (3) | 0.7936 (7) | 0.3170 (2) | 0.0752 (13) | |
| H2 | 0.5584 | 0.7124 | 0.2904 | 0.090* | |
| C3 | 0.5724 (3) | 0.9539 (7) | 0.3454 (2) | 0.0716 (12) | |
| H3 | 0.6358 | 0.9835 | 0.3388 | 0.086* | |
| C4 | 0.5236 (3) | 1.0714 (6) | 0.3839 (2) | 0.0601 (10) | |
| H4 | 0.5533 | 1.1821 | 0.4032 | 0.072* | |
| C5 | 0.4300 (2) | 1.0238 (5) | 0.39356 (17) | 0.0443 (7) | |
| C6 | 0.3702 (2) | 1.1400 (4) | 0.43399 (18) | 0.0452 (8) | |
| C7 | 0.4155 (3) | 1.3171 (5) | 0.4690 (2) | 0.0701 (12) | |
| H7A | 0.4090 | 1.4104 | 0.4307 | 0.105* | |
| H7B | 0.4851 | 1.2979 | 0.4921 | 0.105* | |
| H7C | 0.3812 | 1.3563 | 0.5065 | 0.105* | |
| N3 | 0.0939 (2) | 0.8665 (4) | 0.36804 (17) | 0.0606 (8) | 0.465 (11) |
| C8 | 0.2202 (3) | 1.1843 (5) | 0.4750 (2) | 0.0684 (11) | 0.465 (11) |
| H8A | 0.2116 | 1.3077 | 0.4538 | 0.082* | 0.465 (11) |
| H8B | 0.2549 | 1.1961 | 0.5276 | 0.082* | 0.465 (11) |

| | | | | | |
|------|------------|-------------|--------------|-------------|------------|
| C9 | 0.1197 (6) | 1.1045 (16) | 0.4713 (5) | 0.062 (3) | 0.465 (11) |
| H9A | 0.0797 | 1.1912 | 0.4924 | 0.074* | 0.465 (11) |
| H9B | 0.1263 | 0.9918 | 0.5009 | 0.074* | 0.465 (11) |
| C10 | 0.0677 (7) | 1.0625 (13) | 0.3896 (5) | 0.056 (3) | 0.465 (11) |
| H10A | 0.0886 | 1.1517 | 0.3566 | 0.067* | 0.465 (11) |
| H10B | -0.0043 | 1.0731 | 0.3830 | 0.067* | 0.465 (11) |
| C11 | 0.053 (3) | 0.696 (4) | 0.396 (2) | 0.092 (10) | 0.465 (11) |
| H11A | -0.0185 | 0.7058 | 0.3873 | 0.138* | 0.465 (11) |
| H11B | 0.0812 | 0.6838 | 0.4496 | 0.138* | 0.465 (11) |
| H11C | 0.0705 | 0.5903 | 0.3705 | 0.138* | 0.465 (11) |
| C12 | 0.066 (3) | 0.888 (6) | 0.2842 (6) | 0.071 (8) | 0.465 (11) |
| H12A | -0.0053 | 0.9054 | 0.2678 | 0.107* | 0.465 (11) |
| H12B | 0.0849 | 0.7789 | 0.2611 | 0.107* | 0.465 (11) |
| H12C | 0.0998 | 0.9931 | 0.2700 | 0.107* | 0.465 (11) |
| N3' | 0.0939 (2) | 0.8665 (4) | 0.36804 (17) | 0.0606 (8) | 0.535 (11) |
| C8' | 0.2202 (3) | 1.1843 (5) | 0.4750 (2) | 0.0684 (11) | 0.535 (11) |
| H8'A | 0.2415 | 1.3124 | 0.4803 | 0.082* | 0.535 (11) |
| H8'B | 0.2252 | 1.1336 | 0.5250 | 0.082* | 0.535 (11) |
| C9' | 0.1099 (5) | 1.1713 (11) | 0.4277 (7) | 0.077 (3) | 0.535 (11) |
| H9'A | 0.0703 | 1.2595 | 0.4484 | 0.093* | 0.535 (11) |
| H9'B | 0.1091 | 1.2112 | 0.3768 | 0.093* | 0.535 (11) |
| C10' | 0.0573 (6) | 0.9862 (12) | 0.4224 (6) | 0.069 (3) | 0.535 (11) |
| H10C | 0.0711 | 0.9278 | 0.4717 | 0.082* | 0.535 (11) |
| H10D | -0.0145 | 1.0035 | 0.4055 | 0.082* | 0.535 (11) |
| C11' | 0.036 (2) | 0.703 (3) | 0.3837 (17) | 0.071 (5) | 0.535 (11) |
| H11D | -0.0346 | 0.7331 | 0.3720 | 0.106* | 0.535 (11) |
| H11E | 0.0569 | 0.6702 | 0.4360 | 0.106* | 0.535 (11) |
| H11F | 0.0467 | 0.6009 | 0.3531 | 0.106* | 0.535 (11) |
| C12' | 0.052 (3) | 0.896 (5) | 0.2860 (6) | 0.071 (7) | 0.535 (11) |
| H12D | -0.0197 | 0.9133 | 0.2770 | 0.106* | 0.535 (11) |
| H12E | 0.0653 | 0.7900 | 0.2584 | 0.106* | 0.535 (11) |
| H12F | 0.0815 | 1.0035 | 0.2697 | 0.106* | 0.535 (11) |
| C13 | 0.2546 (3) | 0.6730 (5) | 0.5551 (2) | 0.0572 (9) | |
| C14 | 0.2316 (2) | 0.4674 (5) | 0.29685 (19) | 0.0481 (8) | |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|---------------|--------------|---------------|
| Cu1 | 0.0369 (2) | 0.0457 (2) | 0.0496 (2) | -0.00511 (18) | 0.01183 (16) | -0.00953 (18) |
| N1 | 0.0412 (15) | 0.0515 (17) | 0.0463 (15) | -0.0058 (12) | 0.0131 (12) | -0.0059 (12) |
| N2 | 0.0448 (16) | 0.0426 (16) | 0.0496 (15) | 0.0014 (13) | 0.0109 (12) | -0.0037 (12) |
| N4 | 0.066 (2) | 0.057 (2) | 0.0533 (18) | -0.0049 (15) | 0.0097 (15) | 0.0003 (15) |
| N5 | 0.0559 (19) | 0.0598 (19) | 0.072 (2) | -0.0085 (16) | 0.0162 (16) | -0.0233 (17) |
| S1 | 0.264 (3) | 0.178 (2) | 0.1026 (12) | -0.0650 (18) | 0.0974 (15) | 0.0160 (12) |
| S2 | 0.0560 (6) | 0.0568 (6) | 0.0762 (6) | -0.0069 (4) | 0.0269 (5) | -0.0187 (5) |
| C1 | 0.055 (2) | 0.078 (3) | 0.064 (2) | -0.011 (2) | 0.0212 (19) | -0.025 (2) |
| C2 | 0.051 (2) | 0.107 (4) | 0.075 (3) | -0.005 (2) | 0.031 (2) | -0.023 (2) |
| C3 | 0.042 (2) | 0.106 (4) | 0.071 (3) | -0.012 (2) | 0.0205 (19) | 0.002 (2) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|-------------|--------------|
| C4 | 0.045 (2) | 0.069 (3) | 0.064 (2) | -0.0139 (19) | 0.0068 (17) | 0.003 (2) |
| C5 | 0.0366 (17) | 0.0494 (19) | 0.0435 (17) | -0.0030 (15) | 0.0024 (13) | 0.0060 (15) |
| C6 | 0.0428 (19) | 0.0389 (18) | 0.0486 (18) | -0.0020 (14) | 0.0004 (14) | 0.0025 (14) |
| C7 | 0.062 (2) | 0.046 (2) | 0.093 (3) | -0.0065 (19) | -0.001 (2) | -0.012 (2) |
| N3 | 0.0378 (16) | 0.073 (2) | 0.069 (2) | -0.0027 (15) | 0.0079 (14) | -0.0122 (17) |
| C8 | 0.069 (3) | 0.059 (2) | 0.079 (3) | 0.009 (2) | 0.023 (2) | -0.021 (2) |
| C9 | 0.062 (6) | 0.061 (7) | 0.078 (7) | 0.003 (5) | 0.047 (5) | -0.009 (5) |
| C10 | 0.034 (5) | 0.066 (7) | 0.072 (6) | 0.017 (4) | 0.019 (4) | 0.007 (5) |
| C11 | 0.019 (8) | 0.155 (18) | 0.096 (19) | -0.026 (9) | 0.003 (9) | 0.037 (10) |
| C12 | 0.032 (9) | 0.105 (17) | 0.065 (12) | -0.007 (8) | -0.012 (8) | 0.027 (10) |
| N3' | 0.0378 (16) | 0.073 (2) | 0.069 (2) | -0.0027 (15) | 0.0079 (14) | -0.0122 (17) |
| C8' | 0.069 (3) | 0.059 (2) | 0.079 (3) | 0.009 (2) | 0.023 (2) | -0.021 (2) |
| C9' | 0.067 (6) | 0.058 (6) | 0.126 (10) | 0.003 (4) | 0.061 (7) | -0.008 (6) |
| C10' | 0.058 (5) | 0.074 (6) | 0.083 (7) | 0.005 (4) | 0.034 (5) | 0.001 (5) |
| C11' | 0.038 (12) | 0.120 (12) | 0.053 (6) | -0.007 (6) | 0.007 (8) | 0.019 (6) |
| C12' | 0.050 (13) | 0.061 (10) | 0.100 (15) | -0.006 (7) | 0.013 (8) | 0.010 (9) |
| C13 | 0.066 (2) | 0.043 (2) | 0.061 (2) | -0.0099 (18) | 0.0108 (19) | 0.0024 (18) |
| C14 | 0.0348 (17) | 0.054 (2) | 0.0550 (19) | -0.0070 (15) | 0.0103 (14) | -0.0051 (17) |

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

| | | | |
|--------|-----------|-----------|-----------|
| Cu1—N5 | 1.955 (3) | N3—C11 | 1.494 (9) |
| Cu1—N2 | 2.013 (3) | N3—C10 | 1.537 (7) |
| Cu1—N1 | 2.047 (3) | C8—C9 | 1.483 (7) |
| Cu1—N3 | 2.078 (3) | C8—H8A | 0.9700 |
| Cu1—N4 | 2.153 (3) | C8—H8B | 0.9700 |
| N1—C1 | 1.331 (4) | C9—C10 | 1.525 (9) |
| N1—C5 | 1.343 (4) | C9—H9A | 0.9700 |
| N2—C6 | 1.264 (4) | C9—H9B | 0.9700 |
| N2—C8 | 1.472 (4) | C10—H10A | 0.9700 |
| N4—C13 | 1.126 (4) | C10—H10B | 0.9700 |
| N5—C14 | 1.151 (4) | C11—H11A | 0.9600 |
| S1—C13 | 1.600 (4) | C11—H11B | 0.9600 |
| S2—C14 | 1.622 (4) | C11—H11C | 0.9600 |
| C1—C2 | 1.375 (5) | C12—H12A | 0.9600 |
| C1—H1 | 0.9300 | C12—H12B | 0.9600 |
| C2—C3 | 1.361 (6) | C12—H12C | 0.9600 |
| C2—H2 | 0.9300 | C9'—C10' | 1.514 (9) |
| C3—C4 | 1.373 (5) | C9'—H9'A | 0.9700 |
| C3—H3 | 0.9300 | C9'—H9'B | 0.9700 |
| C4—C5 | 1.380 (5) | C10'—H10C | 0.9700 |
| C4—H4 | 0.9300 | C10'—H10D | 0.9700 |
| C5—C6 | 1.486 (5) | C11'—H11D | 0.9600 |
| C6—C7 | 1.500 (4) | C11'—H11E | 0.9600 |
| C7—H7A | 0.9600 | C11'—H11F | 0.9600 |
| C7—H7B | 0.9600 | C12'—H12D | 0.9600 |
| C7—H7C | 0.9600 | C12'—H12E | 0.9600 |
| N3—C12 | 1.494 (9) | C12'—H12F | 0.9600 |

| | | | |
|------------|-------------|----------------|------------|
| N5—Cu1—N2 | 169.17 (12) | C11—N3—Cu1 | 105.6 (16) |
| N5—Cu1—N1 | 90.84 (12) | C10—N3—Cu1 | 111.3 (4) |
| N2—Cu1—N1 | 79.51 (11) | N2—C8—C9 | 115.7 (4) |
| N5—Cu1—N3 | 90.46 (12) | N2—C8—H8A | 108.3 |
| N2—Cu1—N3 | 95.82 (11) | C9—C8—H8A | 108.3 |
| N1—Cu1—N3 | 152.40 (12) | N2—C8—H8B | 108.3 |
| N5—Cu1—N4 | 96.85 (13) | C9—C8—H8B | 108.3 |
| N2—Cu1—N4 | 90.64 (11) | H8A—C8—H8B | 107.4 |
| N1—Cu1—N4 | 106.39 (12) | C8—C9—C10 | 109.8 (7) |
| N3—Cu1—N4 | 100.82 (12) | C8—C9—H9A | 109.7 |
| C1—N1—C5 | 117.9 (3) | C10—C9—H9A | 109.7 |
| C1—N1—Cu1 | 128.7 (3) | C8—C9—H9B | 109.7 |
| C5—N1—Cu1 | 113.4 (2) | C10—C9—H9B | 109.7 |
| C6—N2—C8 | 119.9 (3) | H9A—C9—H9B | 108.2 |
| C6—N2—Cu1 | 116.6 (2) | C9—C10—N3 | 110.5 (7) |
| C8—N2—Cu1 | 123.3 (2) | C9—C10—H10A | 109.5 |
| C13—N4—Cu1 | 152.2 (3) | N3—C10—H10A | 109.5 |
| C14—N5—Cu1 | 169.4 (3) | C9—C10—H10B | 109.5 |
| N1—C1—C2 | 123.3 (4) | N3—C10—H10B | 109.5 |
| N1—C1—H1 | 118.4 | H10A—C10—H10B | 108.1 |
| C2—C1—H1 | 118.4 | N3—C11—H11A | 109.5 |
| C3—C2—C1 | 118.6 (4) | N3—C11—H11B | 109.5 |
| C3—C2—H2 | 120.7 | H11A—C11—H11B | 109.5 |
| C1—C2—H2 | 120.7 | N3—C11—H11C | 109.5 |
| C2—C3—C4 | 119.2 (4) | H11A—C11—H11C | 109.5 |
| C2—C3—H3 | 120.4 | H11B—C11—H11C | 109.5 |
| C4—C3—H3 | 120.4 | N3—C12—H12A | 109.5 |
| C3—C4—C5 | 119.4 (4) | N3—C12—H12B | 109.5 |
| C3—C4—H4 | 120.3 | H12A—C12—H12B | 109.5 |
| C5—C4—H4 | 120.3 | N3—C12—H12C | 109.5 |
| N1—C5—C4 | 121.6 (3) | H12A—C12—H12C | 109.5 |
| N1—C5—C6 | 114.3 (3) | H12B—C12—H12C | 109.5 |
| C4—C5—C6 | 124.1 (3) | C10'—C9'—H9'A | 107.7 |
| N2—C6—C5 | 116.1 (3) | C10'—C9'—H9'B | 107.7 |
| N2—C6—C7 | 125.6 (3) | H9'A—C9'—H9'B | 107.1 |
| C5—C6—C7 | 118.3 (3) | C9'—C10'—H10C | 109.9 |
| C6—C7—H7A | 109.5 | C9'—C10'—H10D | 109.9 |
| C6—C7—H7B | 109.5 | H10C—C10'—H10D | 108.3 |
| H7A—C7—H7B | 109.5 | H11D—C11'—H11E | 109.5 |
| C6—C7—H7C | 109.5 | H11D—C11'—H11F | 109.5 |
| H7A—C7—H7C | 109.5 | H11E—C11'—H11F | 109.5 |
| H7B—C7—H7C | 109.5 | H12D—C12'—H12E | 109.5 |
| C12—N3—C11 | 114 (2) | H12D—C12'—H12F | 109.5 |
| C12—N3—C10 | 98.4 (18) | H12E—C12'—H12F | 109.5 |
| C11—N3—C10 | 122.9 (19) | N4—C13—S1 | 178.9 (4) |
| C12—N3—Cu1 | 102.7 (18) | N5—C14—S2 | 179.4 (3) |