

A new one-dimensional coordination polymer: $\{[\text{Cu}(\text{C}_{10}\text{H}_9\text{NO}_5\text{S})(\text{H}_2\text{O})]\cdot\text{H}_2\text{O}\}_n$

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Key indicators

Single-crystal X-ray study

$T = 291\text{ K}$

Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$

R factor = 0.029

wR factor = 0.073

Data-to-parameter ratio = 11.9

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, *catena*-poly[[copper(II)- μ -2- $\{[(E)$ -(5-formyl-2-oxidophenyl)methylidene]amino}-1-ethanesulfonato(2-)] monohydrate], $\{[\text{Cu}(\text{L})(\text{H}_2\text{O})]\cdot\text{H}_2\text{O}\}_n$, where $\text{H}_2\text{L} = 2\text{-}\{[(E)$ -(5-formyl-2-hydroxyphenyl)methylidene]amino}-1-ethanesulfonic acid ($\text{C}_{10}\text{H}_{11}\text{NO}_5\text{S}$), was synthesized in water-methanol solution and the crystal structure determined by X-ray diffraction analysis. The Cu atom is five-coordinate, forming a distorted square-pyramidal geometry. This geometry includes a weak bond between the Cu atom and the O atom of the formyl group of another ligand, giving a one-dimensional infinite chain structure. The structure is further stabilized by intermolecular hydrogen bonds.

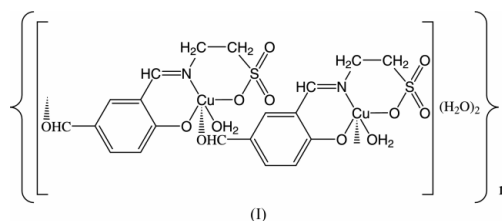
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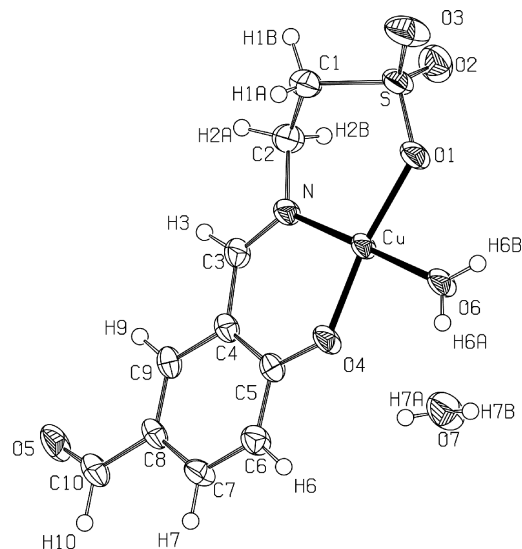
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Comment

5-Formylsalicylaldehyde is an important organic synthetic intermediate that can be used to synthesize macrocyclic complexes (Badri *et al.*, 1990) and photochromic compounds of spiropyrans (Sharmila & Bhaskar, 1996). Studies of Schiff base complexes containing sulfur and complexes of amino acid Schiff bases (Casella & Gullotti, 1981; Wang *et al.*, 1994; Casella & Gullotti, 1986) have aroused more and more interest because of their antiviral, anticancer and antibacterial activities. Recently, Zhang & Jiang (2002) have reported a Schiff base complex derived from taurine, an amino acid containing sulfur. We report here the synthesis and crystal structure of a new Schiff base copper(II) complex, (I), prepared by the reaction of $\text{CuSO}_4\cdot 5\text{H}_2\text{O}$ and the potassium salt of the Schiff base ligand 2- $\{[(E)$ -(2-hydroxy-5- $\{[(2\text{-sulfoethyl})\text{imino}]methyl\}$ phenyl)methylidene]amino}-1-ethanesulfonic acid, derived in turn from the reaction of taurine and 5-formylsalicylaldehyde.



As shown in Figs. 1 and 2, the one-dimensional infinite chain of the title compound, (I), is assembled from $[\text{Cu}(\text{C}_{10}\text{H}_9\text{NO}_5\text{S})(\text{H}_2\text{O})]\cdot\text{H}_2\text{O}$ units, in which the Cu atom is five-coordinate with two O atoms and one N atom of the tridentate ligand L, one O atom of the coordinated water and one O atom of the formyl group belonging to the ligand on an adjacent unit [symmetry code: (i) $1 + x, y, z$], giving a distorted square-pyramidal geometry. The plane O1/N/O4/O6 is the base of the pyramid, while O5ⁱ is the apex. The distance from


Figure 1

View of the asymmetric unit of the title compound, with displacement ellipsoids drawn at the 50% probability level.

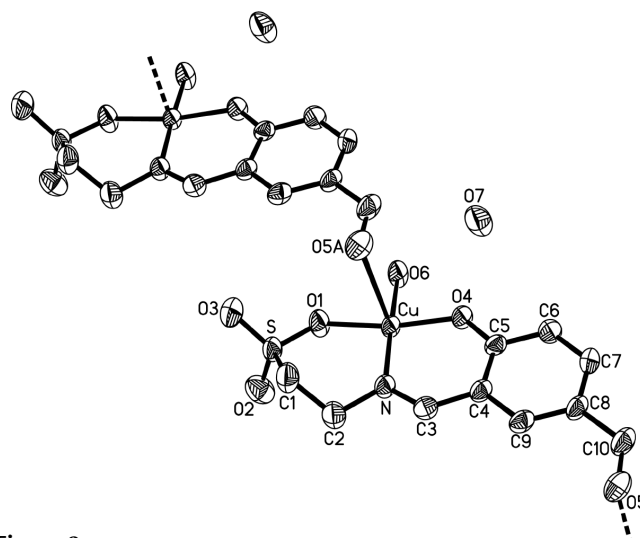
Cu to the least-squares plane O1/N/O4/O6 is 0.0133 (1) Å. The length of the bond Cu—O5ⁱ, which links the structural units to form a one-dimensional infinite chain structure, is 2.877 (3) Å. This bond is much longer than a normal coordinate bond length, so can be called a partly coordinated bond (Procter *et al.*, 1968). The Cu atom forms two six-membered chelate rings with the ligand. The bond length of Cu—O4 [1.8786 (18) Å] is the shortest coordinate bond in (I), while Cu—O1 is 1.9526 (18) Å, indicating that the sulfonate group can coordinate to the metal ion in competition with water molecules.

There are a number of hydrogen bonds in (I). The hydrogen bond involving the coordinated and uncoordinated water molecules is O6—H6A···O7. The polymer chains are linked by four intermolecular hydrogen bonds (Table 2) O6—H6B···O5ⁱ, O7—H7A···O2ⁱⁱ, O7—H7B···O3ⁱⁱⁱ and C10—H10···O1^{iv} (symmetry codes as in Table 2) to form a three-dimensional structure. This is why, despite the weakness of the Cu—O5ⁱ bond, the polymer is stable.

Fig. 1 also shows that there is only one imine group in the complex. This is because, in the case of the imine *ortho* to the hydroxyl, the imino-N, hydroxy-O and Cu can form a six-membered chelate ring that can stabilize the imine, while the other imine has no such effect; as a result, the C=N double bond is broken down by attack of water molecules.

Experimental

A solution of 2.0 mmol of taurine and 2.0 mmol of KOH in 10 ml anhydrous methanol was dropped into a solution of 1.0 mmol of 5-formylsalicylaldehyde in 10 ml anhydrous methanol. The mixture was stirred and refluxed at 323 K for 2 h, and then was cooled to room temperature to give a yellow precipitate, which was collected by filtration, recrystallized from anhydrous methanol, washed successively with anhydrous methanol and ether, and dried *in vacuo* to yield 0.36 g of a yellow product. Analysis, found (%): C 32.85, H 3.12,


Figure 2

View of the interactions between adjacent molecules. H atoms have been omitted for clarity.

N 6.27; C₁₂H₁₄K₂N₂O₇S₂ requires (%): C 32.73, H 3.18, N 6.36. IR (KBr, ν cm⁻¹): 1048.8, 1152.4, 1197.2 (ν SO₃), 1636.2 (ν C=N), 3442.3 (ν O—H). 1.0 mmol of the ligand, the potassium salt of 2-[(*E*)-(2-hydroxy-5-[(2-sulfoethyl)imino]methyl)phenyl)methylidene]amino]-1-ethanesulfonic acid, was dissolved in 25 ml of aqueous methanol. To this solution, 1.0 mmol of CuSO₄·5H₂O was added, and the mixture was stirred and refluxed at 323 K for 6 h, then cooled to room temperature. After filtration, the filtrate was left to stand at room temperature. Dark-green crystals suitable for X-ray diffraction were obtained in a yield of 43%. Analysis, found (%): C 33.73, H 3.61, N 4.07; C₁₀H₁₃CuNO₇S requires (%): C 33.82, H 3.66, N 3.95%. IR (KBr, ν cm⁻¹): 1039.5, 1129.8, 1156.1, 1182.8 (ν SO₃), 1619.52 (ν C=N), 3433.5 (ν O—H).

Crystal data

[Cu(C₁₀H₁₁NO₅S)(H₂O)]·H₂O
M_r = 354.81
 Monoclinic, *P*2₁/*c*
a = 8.589 (2) Å
b = 17.585 (2) Å
c = 9.479 (2) Å
 β = 115.69 (1)°
V = 1290.1 (4) Å³
Z = 4

D_x = 1.827 Mg m⁻³
 Mo *K* α radiation
 Cell parameters from 33 reflections
 θ = 3.5–15.3°
 μ = 1.89 mm⁻¹
T = 291 (2) K
 Plate, green
 0.58 × 0.46 × 0.20 mm

Data collection

Siemens *P4* diffractometer
 ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.405, *T_{max}* = 0.686
 2761 measured reflections
 2409 independent reflections
 1963 reflections with *I* > 2 σ (*I*)

R_{int} = 0.014
 θ_{max} = 25.5°
h = 0 → 10
k = 0 → 21
l = -11 → 10
 3 standard reflections every 97 reflections
 intensity decay: 5.2%

Refinement

Refinement on *F*²
R[*F*² > 2 σ (*F*²)] = 0.029
wR(*F*²) = 0.073
S = 1.04
 2409 reflections
 202 parameters
 H atoms treated by a mixture of independent and constrained refinement

w = 1/[$\sigma^2(F_o^2) + (0.0403P)^2$]
 where *P* = (*F_o*² + 2*F_c*²)/3
 $(\Delta/\sigma)_{\text{max}}$ = 0.001
 $\Delta\rho_{\text{max}}$ = 0.34 e Å⁻³
 $\Delta\rho_{\text{min}}$ = -0.26 e Å⁻³
 Extinction correction: SHELXL97
 Extinction coefficient: 0.0077 (7)

Table 1
Selected geometric parameters (Å, °).

Cu—O4	1.8786 (18)	Cu—N	1.960 (2)
Cu—O6	1.9346 (19)	Cu—O5 ⁱ	2.877 (3)
Cu—O1	1.9526 (18)		
O4—Cu—O6	84.54 (8)	O1—Cu—N	96.72 (8)
O4—Cu—O1	168.50 (8)	O4—Cu—O5 ⁱ	94.80 (8)
O6—Cu—O1	84.18 (8)	O6—Cu—O5 ⁱ	87.34 (8)
O4—Cu—N	94.68 (8)	O1—Cu—O5 ⁱ	82.43 (8)
O6—Cu—N	176.48 (10)	N—Cu—O5 ⁱ	96.16 (8)

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

Table 2
Hydrogen-bonding 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>
O6—H6A···O7	0.813 (10)	1.842 (11)	2.654 (3)	177 (3)
O6—H6B···O5 ⁱⁱ	0.808 (10)	1.905 (11)	2.713 (3)	178 (3)
O7—H7A···O2 ⁱⁱⁱ	0.820 (10)	2.078 (17)	2.866 (4)	161 (4)
O7—H7B···O3 ^{iv}	0.813 (10)	1.99 (2)	2.768 (3)	159 (6)
C10—H10···O1 ^v	1.02 (3)	2.58 (3)	3.417 (3)	139 (2)

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

The H atoms on O6, O7 and C10 were located in a difference Fourier map and their positions and isotropic displacement parameters were refined, with the O—H distance restrained to 0.82 Å; the C10—H10 refined distance is 1.02 (3) Å. All other H atoms were

positioned geometrically and were treated as riding atoms, with C—H distances of 0.93–0.97 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: *XSCANS* (Siemens, 1994); cell refinement: *XSCANS*; data reduction: *SHELXTL* (Siemens, 1994); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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