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Key indicators

Single-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C}) = 0.005 \text{ \AA}$
R factor = 0.025
wR factor = 0.046
Data-to-parameter ratio = 6.8

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

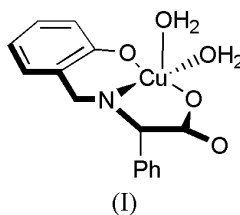
Diaqua(*N*-salicylidene-*L*-2-phenylglycinate- $\kappa^3\text{N},\text{O},\text{O}'$)copper(II)

The title compound, $[\text{Cu}(\text{C}_{15}\text{H}_{11}\text{NO}_3)(\text{H}_2\text{O})_2]$, was formed by the reaction of cupric perchlorate, *L*- α -phenylglycine and salicylaldehyde. Crystallographic analysis shows that the compound is a new Cu–Schiff base of an amino acid complex. The ligand chelates with Cu^{II} in a tridentate fashion. One water molecule is in this ligand plane and another is in an axial position. Extensive hydrogen-bonding interactions are present.

Comment

Transition metal complexes of chiral ligands attract widespread attention from the viewpoint of their structures, molecular recognition, and molecular processes (Whitesides *et al.*, 1995; Philp & Stoddart, 1996). As model systems for pyridoxal-potentiated enzymes, *N*-salicylidene-*L*-amino acids are being applied to synthesize a series of transition metal–chiral ligand complexes. Several *N*-salicylidene-*L*-amino acid–transition metal complexes have been synthesized and studied, using amino acids with different side-chain groups, such as valine (Rajak *et al.*, 1999), glutamic acid (Korhonen *et al.*, 1984), phenylalanine (Marinovich *et al.*, 1999), alanine (Warda, 1999) and methionine (Palacios *et al.*, 1989).

We report here a new Cu^{II} complex of an *N*-salicylidene-amino acid, (I), formed from copper(II), *L*-2-phenyl-glycine and salicylaldehyde.



The coordination of the molecule can be described as 4 + 1; each Cu^{II} ion is coordinated by the carboxylate and phenolate O atoms and amine N atom from the ligand, with bond distances of 1.950 (2), 1.932 (2) and 1.900 (2) \AA , respectively, together with one water molecule at 1.960 (2) \AA , forming a square-planar geometry. The axial position is occupied by another coordinated water molecule, with a Cu–O distance of 2.400 (3) \AA . Except for the phenyl group, all the atoms of the organic ligand and Cu^{II} are coplanar, with an average deviation of 0.084 \AA . In the crystal structure of the title compound, hydrogen-bonding interactions between the coordinated water molecules and free carboxylate and phenolate O atoms result in a two-dimensional layer structure, as shown in Fig. 2.

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Experimental

A 15.0 ml methanol solution of salicylaldehyde (100 ml, 1.0 mmol) and cupric perchlorate (576 mg, 1.0 mmol) was added to a 15.0 ml suspended aqueous suspension of L-2-phenyl-glycine (150 mg, 1.0 mmol). The resulting mixture was stirred until the L-2-phenyl-glycine completely dissolved. Green prismatic single crystals of the title complex (yield 72%), suitable for X-ray diffraction analysis, were obtained by evaporation of the filtered mixture at room temperature after 6 d.

Crystal data

[Cu(C₁₅H₁₁NO₃)(H₂O)₂]

M_r = 352.82

Monoclinic, *P*2₁

a = 5.7870 (12) Å

b = 9.4205 (19) Å

c = 13.242 (3) Å

β = 92.81 (3)°

V = 721.0 (3) Å³

Z = 2

D_x = 1.625 Mg m⁻³

Mo *K*α radiation

Cell parameters from 64

reflections

θ = 2.7–27.5°

μ = 1.54 mm⁻¹

T = 293 (2) K

Plate, green

0.40 × 0.36 × 0.16 mm

Data collection

Siemens SMART CCD
diffractometer

ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

T_{min} = 0.565, *T_{max}* = 0.782

2514 measured reflections

1757 independent reflections

1509 reflections with *I* > 2σ(*I*)

R_{int} = 0.034

θ_{max} = 27.5°

h = -7 → 7

k = -12 → 12

l = -17 → 17

Refinement

Refinement on *F*²

R[*F*² > 2σ(*F*²)] = 0.025

wR(*F*²) = 0.046

S = 0.85

1757 reflections

260 parameters

All H-atom parameters refined

w = 1/[σ²(*F_o*²) + (0.0224*P*)²]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = 0.116

Δρ_{max} = 0.23 e Å⁻³

Δρ_{min} = -0.29 e Å⁻³

Extinction correction: *SHELXL97*

Extinction coefficient: 0.0031 (12)

Absolute structure: Flack (1983),

xxxx Friedel pairs [PLEASE
SUPPLY]

Flack parameter = 0.061 (18)

Table 1

Selected geometric parameters (Å, °).

N—Cu	1.900 (2)	Cu—O5	2.400 (3)
Cu—O1	1.932 (2)	O4—H4B	0.81 (4)
Cu—O2	1.950 (2)	O5—H5A	0.75 (4)
Cu—O4	1.960 (3)	O5—H5B	1.13 (7)
N—Cu—O1	93.93 (10)	O2—Cu—O4	93.88 (11)
N—Cu—O2	84.43 (10)	N—Cu—O5	90.74 (10)
O1—Cu—O2	173.10 (11)	O1—Cu—O5	98.30 (11)
N—Cu—O4	174.51 (13)	O2—Cu—O5	88.44 (10)
O1—Cu—O4	87.12 (11)	O4—Cu—O5	94.45 (11)

One reflection, 001, was omitted due to extinction.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SMART*; data reduction: *SMART* (Siemens, 1994) and *XPREP* in *SHELXTL* (Siemens, 1994); program(s) used to solve structure: *SHELXTL*; program(s) used to refine structure: *SHELXTL*; mole-

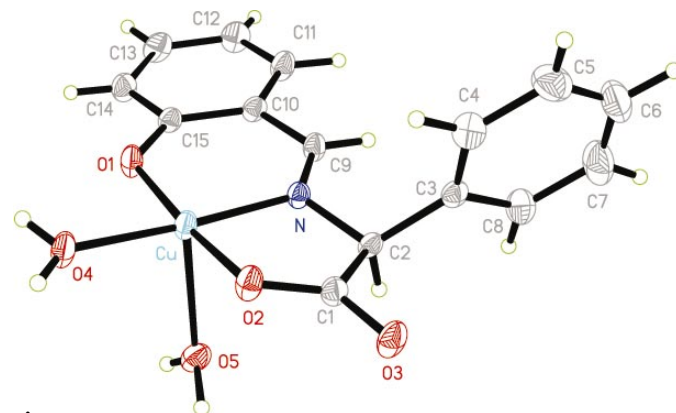


Figure 1

A displacement ellipsoid plot at the 30% probability level.

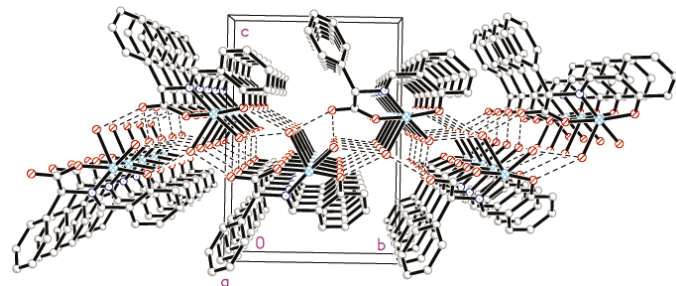


Figure 2

A view of the packing, as viewed down the *a* axis.

cular graphics: *SHELXL97* (Siemens, 1996); software used to prepare material for publication: *SHELXL97* (Siemens, 1996).

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