

Jin-Ling Wang,* Feng Ding and
Fang-Ming MiaoCollege of Chemical and Life Science, Tianjin
Normal University, Tianjin 300074, People's
Republic of ChinaCorrespondence e-mail:
wangjinling43@eyou.com

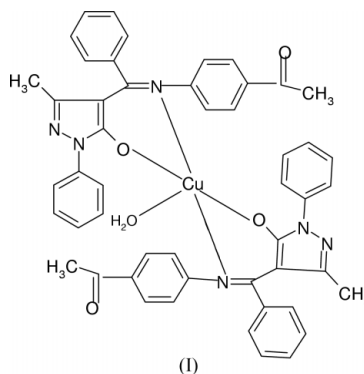
Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.008\text{ \AA}$
H-atom completeness 96%
 R factor = 0.055
 wR factor = 0.096
Data-to-parameter ratio = 10.4For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Bis{4-[α -(4-acetylphenylimino)benzyl]-
3-methyl-1-phenylpyrazol-5-onato}-
aquacopper(II)

The Cu atom in the title complex, $[\text{Cu}(\text{C}_{25}\text{H}_{20}\text{N}_3\text{O}_2)_2(\text{H}_2\text{O})]$, has a distorted trigonal-bipyramidal coordination environment, with two O atoms of the pyrazolone rings and a water O atom in the equatorial plane and two N atoms in axial positions [$\text{N}-\text{Cu}-\text{N} = 159.34(16)^\circ$]. The complex occupies a special position on a twofold axis running through the Cu atom and the water O atom. The $\text{Cu}-\text{O}(\text{H}_2\text{O})$ bond [$2.549(4)\text{ \AA}$] is substantially longer than the rest of the bonds in the coordination polyhedron [$\text{Cu}-\text{O} = 1.919(2)\text{ \AA}$ and $\text{Cu}-\text{N} = 1.990(3)\text{ \AA}$]. The pyrazolone ring is almost coplanar with the adjacent chelate ring, the dihedral angle formed by the mean plane of the pyrazolone ring and the planar part of the chelate ring being $2.91(3)^\circ$.

Comment

Schiff base compounds are relatively easy to obtain and they frequently show high biological activity. Therefore, studies of Schiff bases and their metal complexes are the focal point of many research groups working in the fields of coordination chemistry, biomedicine and pharmaceutical chemistry (Vigoto, 1987).



4-Acyl-5-pyrazolones are enolizable β -diketones which afford stable chelate complexes with several metals (Miyazaki *et al.*, 1989; Morales *et al.*, 1988). A number of compounds derived from 4-acyl-5-pyrazolones and aromatic amines have been synthesized (Wang, Yu, Xu & Miao, 2002; Wang, Yu, Yang & Miao, 2002; Yu *et al.*, 2002). The ligand prepared by condensation of 1-phenyl-3-methyl-4-benzoyl-5-pyrazolone (PMBP) with *p*-aminoacetophenone was used for the synthesis of the title copper complex, (I), the structure of which is reported here.

The molecule has crystallographic twofold rotation symmetry, the axis passing through Cu1 and the coordinated water molecule. Atom Cu1 has a distorted trigonal-bipyramidal environment, with atoms N3 and N3ⁱ [symmetry code:

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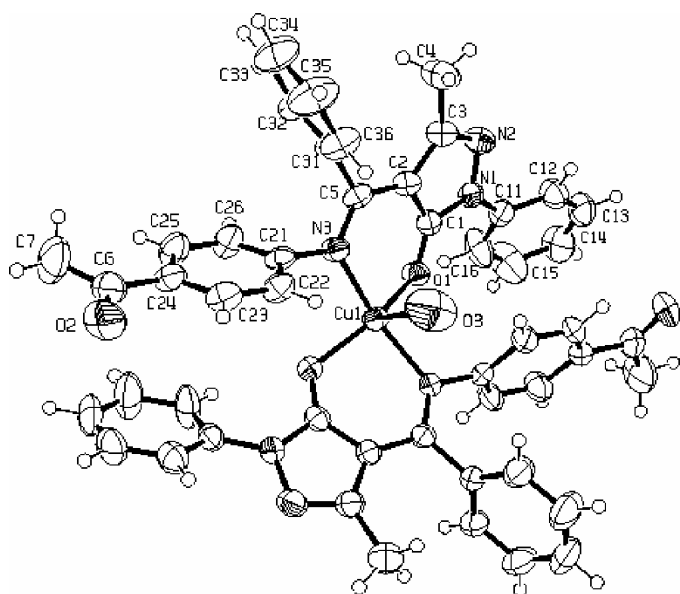


Figure 1
The molecular structure of (I), shown with 50% probability displacement ellipsoids.

(i) $-x, y, \frac{3}{2} - z$] in axial positions [N3—Cu1—N3ⁱ = 159.34 (16)°] and both pyrazolone and water O atoms (O1, O1ⁱ and O3) in the equatorial plane (Fig. 1). The Cu1—O3 bond [2.549 (4) Å] is substantially longer than other bonds in the coordination sphere of the Cu1 atom [Cu1—O1 = 1.919 (2) Å and Cu1—N3 = 1.990 (3) Å], which would be typical for square-planar Cu²⁺ complexes with additional coordination [4 + 1]. However, the description of the coordination as square-pyramidal, though in some sense logical, would be geometrically inaccurate, as the CuO₂N₂ fragment shows a pronounced tetrahedral distortion [O1—Cu1—O1ⁱ = 144.56 (14)°]. A pentacoordinated copper complex similar to (I) was recently reported for the thiosemicarbazide derivative, aqua(3-hydroxy-5-hydroxymethyl-2-methyl-pyridine-4-carboxaldehyde 3-methylisothiosemicarbazone)(nitrate)-copper(II) nitrate (Leovac *et al.*, 2002).

The earlier studied PMBP complexes, however, had metal centers with higher coordination numbers [e.g. (PMBP)₂Sn(*t*-C₄H₉)₂ (Pettinari *et al.*, 1991) and (PMBP)₂SnMe₂ (Bovio *et al.*, 1993)].

The chelate ring Cu1—O1—C1—C2—C5—N3 has a sofa conformation with Cu1 displaced by 0.617 (4) Å from the mean O1/C1/C2/C5/N3 plane [maximum deviation of 0.042 (2) Å for atom C5]. The latter plane is almost coplanar with the mean pyrazolone ring plane; the dihedral angle formed by these planes [2.91 (3)°] is close to analogous dihedral angles in the Co(PMBP-*m*-chloroaniline) complex [2.47 (5) and 1.69 (6)°; Wang *et al.*, 2003]. Owing to steric hindrance, the chelate ring plane is nearly orthogonal to the rings planes C21—C26 and C31—C36 [respective dihedral angles are 84.39 (10) and 77.29 (9)°].

Though it was impossible to locate the water H atoms, it seems very likely that the intermolecular O2...O3ⁱⁱ

2.725 (4) Å contact [symmetry code: (ii) $\frac{1}{2} + x, \frac{1}{2} - y, \frac{1}{2} + z$], in fact, corresponds to a hydrogen bond. These bonds link the molecules into infinite chains along the [101] direction.

Experimental

A mixture of 10 ml of 0.1 M *p*-aminoacetophenone ethanol solution and 10 ml of 0.1 M PMBP ethanol solution was refluxed for 4 h over a steam bath. The excess solvent was removed by evaporation. The Schiff base ligand was collected as a cream-colored powder and dried in air. 10 ml of 0.1 M Cu(NO₃)₂·H₂O ethanol solution was added dropwise to 10 ml of a 0.2 M solution of the ligand in ethanol. After vigorous stirring for 6–8 h at 343–353 K, a dark-brown product precipitated; this was filtered off, washed with cold anhydrous ethanol solution several times and dried in a vacuum desiccator over CaCl₂. The Cu complex was recrystallized from ethanol solution. Elemental analyses for *L* and CuL₂ respectively are as follows (calculated values in parentheses): C 76.60 (76.12) and 69.15 (69.00)%, H 5.41 (5.11) and 5.12 (4.86), N 10.81 (10.65) and 10.13 (9.66)%. IR spectra in the region 4000–400 cm⁻¹ were determined with a KBr pellet on an AVATAR 360 (USA) IR spectrophotometer. The C=O absorption bands, observed at 1637 and 1505 cm⁻¹ for PMBP, shifted to 1672 and 1662 cm⁻¹ for the uncoordinated ligand, and to 1660 and 1630 cm⁻¹ for the complex; the C=N absorption bands, observed at 1572 and 1541 cm⁻¹ in the ligand, red shifted to 1570 and 1533 cm⁻¹ in the complex, indicating that the carbonyl O and imine N atoms of the ligand are coordinated to the Cu atom in the complex.

Crystal data

[Cu(C ₂₅ H ₂₀ N ₃ O ₂) ₂ (H ₂ O)]	$D_x = 1.359 \text{ Mg m}^{-3}$
$M_r = 870.35$	Mo K α radiation
Monoclinic, <i>C</i> 2/ <i>c</i>	Cell parameters from 790 reflections
$a = 22.652$ (6) Å	$\theta = 2.3$ –20.9°
$b = 16.228$ (5) Å	$\mu = 0.57 \text{ mm}^{-1}$
$c = 15.250$ (5) Å	$T = 293$ (2) K
$\beta = 130.636$ (14)°	Block, dark brown
$V = 4254$ (2) Å ³	0.28 × 0.25 × 0.22 mm
$Z = 4$	

Data collection

Bruker SMART 1000 CCD area-detector diffractometer	2050 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\text{int}} = 0.085$
Absorption correction: none	$\theta_{\text{max}} = 25.0^\circ$
8399 measured reflections	$h = -26 \rightarrow 25$
3765 independent reflections	$k = -17 \rightarrow 19$
	$l = -9 \rightarrow 18$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0266P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.055$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.096$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 0.97$	$\Delta\rho_{\text{max}} = 0.27 \text{ e \AA}^{-3}$
3765 reflections	$\Delta\rho_{\text{min}} = -0.27 \text{ e \AA}^{-3}$
362 parameters	Extinction correction: <i>SHELXL97</i>
All H-atom parameters refined	Extinction coefficient: 0.00032 (12)

All aryl and methyl H atoms were located in a difference map and refined with isotropic displacement parameters [C—H = 0.83 (4)–1.02 (4) Å]. The H atoms of the aqua ligand (O3) were not located.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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