

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.029$
 $wR(F^2) = 0.058$
 $S = 1.129$
 4079 reflections
 262 parameters
 H-atom parameters
 constrained

$w = 1/\sigma^2(F_o^2)$
 $(\Delta/\sigma)_{\max} = -0.004$
 $\Delta\rho_{\max} = 0.326 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.776 \text{ e } \text{\AA}^{-3}$
 Extinction correction: none
 Scattering factors from
*International Tables for
 Crystallography* (Vol. C)

Table 1. Selected geometric parameters (\AA , $^\circ$)

Mo—C3	1.949 (3)	Mo—P	2.4994 (8)
Mo—C2	1.986 (2)	O1—C1	1.149 (4)
Mo—C1	2.012 (3)	O2—C2	1.151 (3)
Mo—C4	2.041 (3)	O3—C3	1.160 (4)
Mo—N	2.326 (2)	O4—C4	1.128 (4)
C3—Mo—C2	88.08 (10)	C3—Mo—P	99.42 (7)
C3—Mo—C1	89.89 (12)	C2—Mo—P	172.24 (6)
C2—Mo—C1	91.33 (11)	C1—Mo—P	90.66 (8)
C3—Mo—C4	91.74 (13)	C4—Mo—P	88.72 (8)
C2—Mo—C4	89.08 (11)	N—Mo—P	75.70 (5)
C1—Mo—C4	178.33 (11)	O1—C1—Mo	178.9 (2)
C3—Mo—N	175.00 (7)	O2—C2—Mo	178.3 (3)
C2—Mo—N	96.75 (8)	O3—C3—Mo	179.2 (2)
C1—Mo—N	91.28 (10)	O4—C4—Mo	178.6 (3)
C4—Mo—N	87.07 (11)		

All non-H atoms were located by direct methods and subsequent alternate cycles of difference Fourier synthesis and full-matrix least-squares refinement. The final structural model used anisotropic displacement parameters for the non-H atoms. The H atoms were included in geometrically idealized positions employing appropriate riding models with isotropic displacement parameters constrained to $1.2U_{\text{eq}}$ of their carrier atoms.

Data collection: local diffractometer software (Gomm, 1993). Cell refinement: local diffractometer software. Data reduction: local diffractometer software. Program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *XP* (Siemens, 1990). Software used to prepare material for publication: *SHELXL93*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1116). Services for accessing these data are described at the back of the journal.

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(2,2'-Bipyridine)(flavonolato)copper(II) Perchlorate, † [Cu(bpy)(fla)]ClO₄

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Abstract

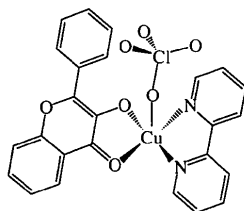
3-Hydroxyflavone coordinates to Cu^{II} together with the auxiliary ligand 2,2'-bipyridine to form a stable cationic flavonolato–copper(II) complex, [Cu(C₁₅H₉O₃)(C₁₀H₈N₂)]ClO₄. The complex has distorted square pyramidal geometry. Two O atoms of the flavonol and the N atoms of the chelating bpy ligand are in the basal position, while a perchlorate O atom occupies the apical position.

Comment

Quercetin 2,3-dioxygenase is a Cu^{II}-containing enzyme which catalyses the degradation of quercetin and related compounds into a depside (phenolic carboxylic acid ester) (Westlake, Talbot, Blakely & Simpson, 1959; Takamura & Ito, 1977). Quercetin coordinates to the copper(II) ion and flavonol forms stable Cu^I and Cu^{II} compounds (Speier, Fülöp & Párkányi, 1990; Balogh-Hergovich, Speier & Argay, 1991). Simple flavonolato–copper complexes have been used as model compounds and in model reactions (Utaka, Hojo, Fujii & Takeda, 1984; Utaka & Takeda, 1985). Following this, we report here the molecular structure of a cationic mixed-

† Alternative name: (2,2'-bipyridyl-*N,N'*)(4-oxo-2-phenyl-4*H*-chromen-3-olato-*O*³,*O*⁴)copper(II) perchlorate.

ligand flavonolato—Cu^{II} complex, (I), in order to gain information about the possible structure of the active site of the enzyme and its role in the catalytic process.



(I)

Compound (I) has a distorted square-pyramidal geometry around the copper(II) ion. Flavonol coordinates through its 3-hydroxy and 4-carbonyl groups with the O atoms occupying the basal positions together with the N atoms of the chelating bpy ligand. One of the perchlorate O atoms occupies the apical position. The Cu—O bond distance of the 4-carbonyl O atom is longer [Cu1—O1 1.970 (3) Å] than that found in [Cu(flav)₂] [1.944 (3) Å], and that of the 3-hydroxy O atom is shorter [Cu1—O2 1.897 (3) Å] than in [Cu(flav)₂] [1.901 (2) Å]. The apical Cu—O distance [Cu1—O6 2.425 (4) Å] is considerably longer than those in the basal position and as observed in a Cu^{II} perchlorate compound with *P,P,N,N*-coordination (Pilloni, Bandoli, Tisato & Corain, 1996). The O—Cu—O and N—Cu—N angles differ only slightly from the ideal 90°.

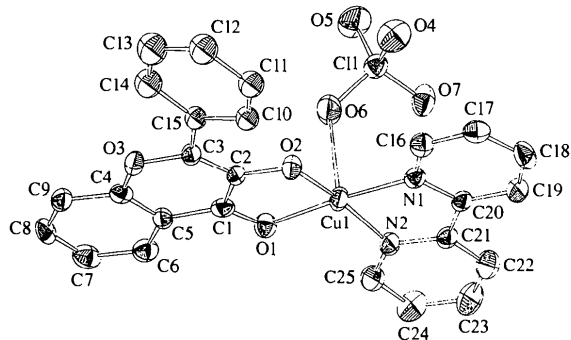


Fig. 1. The molecular structure showing 50% probability displacement ellipsoids. H atoms have been omitted for clarity.

Experimental

The title complex was prepared by stirring [Cu(CH₃CN)₄]ClO₄ with equimolar amounts of flavonol and 2,2'-bipyridine in acetonitrile at room temperature under a dioxigen atmosphere. The resulting green precipitate was crystallized from acetonitrile.

Crystal data

[Cu(C₁₅H₉O₃)(C₁₀H₈N₂)]-ClO₄

M_r = 556.40

Monoclinic

*P*2₁/*c*

Mo *K*α radiation

λ = 0.71073 Å

Cell parameters from 25 reflections

θ = 5.0–14.0°

a = 10.305 (3) Å
b = 22.511 (7) Å
c = 9.641 (3) Å
β = 83.25 (2)°
V = 2221.0 (12) Å³
Z = 4
D_x = 1.664 Mg m⁻³
D_m not measured

μ = 1.157 mm⁻¹

T = 293 (2) K

Prism

0.30 × 0.30 × 0.20 mm

Green

Data collection

Siemens (Nicolet Syntax) diffractometer

ω scan

Absorption correction:

ψ scan (*XDISK*; Siemens, 1990)

T_{min} = 0.573, *T_{max}* = 0.793

3481 measured reflections

3282 independent reflections

2960 reflections with

I > 2σ(*I*)

R_{int} = 0.017

θ_{max} = 23.5°

h = -5 → 11

k = -7 → 25

l = -10 → 10

3 standard reflections

every 50 reflections

intensity decay: 2%

Refinement

Refinement on *F*²

R(*F*) = 0.053

wR(*F*²) = 0.162

S = 1.911

3282 reflections

328 parameters

H atoms riding

w = 1/[σ²(*F_o*²) + (0.0738*P*)²]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = -0.058

Δρ_{max} = 2.764 e Å⁻³

Δρ_{min} = -0.857 e Å⁻³

Extinction correction: none

Scattering factors from

International Tables for

Crystallography (Vol. C)

Table 1. Selected geometric parameters (Å, °)

Cu1—O2	1.897 (3)	O2—C2	1.323 (5)
Cu1—N2	1.961 (4)	O3—C4	1.362 (5)
Cu1—O1	1.970 (3)	O3—C3	1.378 (5)
Cu1—N1	1.978 (4)	C1—C5	1.444 (6)
Cu1—O6	2.425 (4)	C1—C2	1.448 (6)
O1—C1	1.271 (5)	C2—C3	1.362 (6)
O2—Cu1—N2	170.29 (13)	C2—O2—Cu1	110.7 (3)
O2—Cu1—O1	85.64 (12)	C11—O6—Cu1	125.5 (2)
N2—Cu1—O1	95.64 (13)	C16—N1—Cu1	125.5 (3)
O2—Cu1—N1	96.12 (13)	C20—N1—Cu1	114.5 (3)
N2—Cu1—N1	82.07 (14)	C21—N2—Cu1	114.4 (3)
O1—Cu1—N1	176.29 (13)	C25—N2—Cu1	127.2 (3)
O2—Cu1—O6	93.68 (14)	O1—C1—C5	123.1 (4)
N2—Cu1—O6	96.01 (14)	O1—C1—C2	118.1 (4)
O1—Cu1—O6	85.82 (13)	O2—C2—C3	124.8 (4)
N1—Cu1—O6	97.30 (13)	O2—C2—C1	116.1 (4)
C1—O1—Cu1	109.4 (3)	C2—C3—O3	121.5 (4)

The low θ_{max} value (23.5°) is related to the low reflectivity beyond 24°. The high value of *S* (1.911) is probably a result of the somewhat higher rest density near the Cu atom. Despite the experimental absorption correction (ψ scan), the high value of Δρ_{max} (2.764 e Å⁻³) could not be reduced (0.9 Å from Cu).

Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1079). Services for accessing these data are described at the back of the journal.

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[*N,N*-Bis(3,5-dimethylpyrazol-1-yl-methyl)-1-hydroxy-2-aminoethane](3,5-dimethylpyrazole)copper(II) Diperchlorate†

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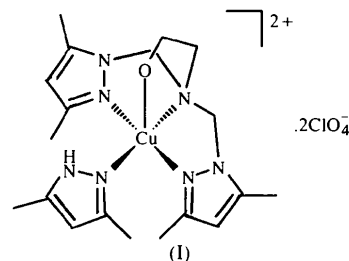
Abstract

The title compound, [Cu(C₅H₈N₂)(C₁₄H₂₃N₅O)](ClO₄)₂, consists of a five-coordinate square-pyramidal copper(II) complex as its perchlorate salt. The copper(II) ion is coordinated in the basal plane by azole and amine N atoms of the *N,N*-bis[(3,5-dimethyl-1-pyrazolyl)methyl]-1-hydroxy-2-aminoethane (bpmhe) ligand and 3,5-dimethylpyrazole, and in the apical position by an alcohol O atom.

† Alternative name: {2-[*N,N*-bis(3,5-dimethylpyrazol-1-yl-*N*²)methyl]amino-*N*}ethanol-*O*}(3,5-dimethylpyrazole-*N*²)copper(II) diperchlorate.

Comment

Copper(II) complexes of the tripodal ligand bpmhe (Driessen, 1982) of the type [Cu(bpmhe)*L*]*X* (*L* = Br, Cl, NO₃, N₃; *X* = Cl, Br, BF₄) have been reported as mononuclear five-coordinate complexes (Malachowski, Davidson & Hoffman, 1989). The ligand bpmhe may act as a tetradentate ligand with two pyrazole nitrogen, one tertiary amine and one alcohol oxygen donor. We have been interested in the formation of such complexes with *L* as a neutral donor ligand and have isolated the title compound [Cu(bpmhe)*L*](ClO₄)₂, where *L* is 3,5-dimethylpyrazole.



The coordination geometry around the copper(II) ion consists of two pyrazole N atoms, an alcohol O atom and an amine N atom of the bpmhe ligand and one N atom of the 3,5-dimethylpyrazole molecule in a square-pyramidal environment. The four N atoms form the base of the pyramid with the alcohol O atom at the apex. The Cu—N bond distances and angles are similar to those found in other *N,N*-bis(pyrazol-1-ylmethyl)amine complexes of copper(II) (Blonk, Driessen & Reedijk, 1985; Driessen, De Graaff & Wiesmeijer, 1987). The

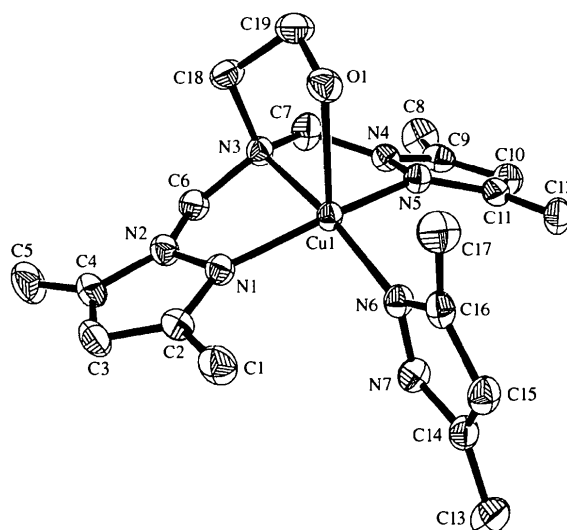


Fig. 1. The structure of the cation showing 50% probability displacement ellipsoids. H atoms have been omitted for clarity.