

[*N'*-Benzoyl-*N,N*-bis(2-pyridylmethylmethyl- κ N)ethylenediamine- κ N]-chlorocopper(II) hexafluorophosphate

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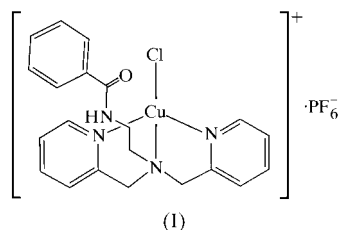
Data validation number: IUC0000107

The title mononuclear copper(II) complex, [CuCl(C₂₁H₂₂N₄O)]PF₆, shows a distorted square-planar coordination and the benzoylamino N atom does not coordinate to the Cu atom.

Comment

Crystal structures and catalytic activities of copper(II) complexes containing *N,N'*-bis(2-pyridylmethyl)- β -alanine-amide ligands were reported by Okuno *et al.* (1997). The copper(II) complexes with similar tripodal ligands containing a peptide group convert supercoiled plasmid DNA into a mixture of nicked and linear forms in the presence of hydrogen peroxide (Kobayashi *et al.*, 1998).

In the crystals of the title compound, [Cu(dpab)Cl]PF₆ [(I), dpab is *N'*-benzoyl-*N,N*-bis(2-pyridylmethyl)ethylenediamine], the benzoylamino N4 atom does not coordinate to the Cu atom; Cu...N4 = 2.827 (4) Å. The N4—H1 moiety is incorporated in a weak bifurcated hydrogen bond with the F4 and F5 atoms of PF₆; the N4...F4 and N4...F5 distances are 3.421 (9) and 3.383 (10) Å, respectively.



Experimental

The ligand and its chlorocopper(II) complex were prepared as described previously (Okuno *et al.*, 1997; Kobayashi *et al.*, 1998). Crystals of the title compound were grown from an aqueous methanol/acetonitrile solution.

Crystal data

[CuCl(C₂₁H₂₂N₄O)]PF₆
M_r = 590.4
Monoclinic, P2₁/n
a = 13.064 (3) Å
b = 13.347 (4) Å
c = 14.077 (3) Å
β = 101.52 (2)°
V = 2405.0 (10) Å³
Z = 4

D_x = 1.631 Mg m⁻³
Mo Kα radiation
Cell parameters from 25 reflections
θ = 10–15°
μ = 1.155 mm⁻¹
T = 296 K
Plate, blue
0.6 × 0.5 × 0.1 mm

Data collection

Rigaku AFC-5S diffractometer
θ–2θ scans
Absorption correction: by integration (Coppens *et al.*, 1965)
T_{min} = 0.868, T_{max} = 0.966
5998 measured reflections
5513 independent reflections
4029 reflections with I > 2σ(I)

R_{int} = 0.012
θ_{max} = 27.5°
h = 0 → 17
k = 0 → 17
l = -18 → 18
3 standard reflections every 100 reflections
intensity decay: none

Refinement

Refinement on F²
R(F) = 0.065
wR(F²) = 0.196
S = 1.38
5513 reflections
316 parameters

H-atom parameters not refined
w = 1/[σ²(F_o²) + {0.1(F_o² + 2F_c²)/3}]
(Δ/σ)_{max} = 0.004
Δρ_{max} = 0.67 e Å⁻³
Δρ_{min} = -0.84 e Å⁻³

Table 1

Selected geometric parameters (Å).

| | | | |
|---------|-----------|--------|-----------|
| Cu1—Cl1 | 2.249 (1) | Cu1—N2 | 2.048 (3) |
| Cu1—N1 | 1.986 (3) | Cu1—N3 | 1.991 (4) |

Positional parameters of all the H atoms were calculated geometrically and fixed with U(H) = 1.2U_{eq}(parent atom).

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1993); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN* (Molecular Structure Corporation, 1999); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *TEXSAN*; software used to prepare material for publication: *TEXSAN*.

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