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# Ricardo Baggio,<sup>a</sup>\* Maria Teresa Garland,<sup>b,c</sup> Ana María Atria,<sup>c,d</sup> Oscar Hidalgo<sup>d</sup> and Santiago Solezi<sup>d</sup>

<sup>a</sup>Departamento de Física, Comisión Nacional de Energía Atómica, Buenos Aires, Argentina, <sup>b</sup>Departamento de Física, Facultad de Ciencias Físicas y Matemáticas, Universidad de Chile, Casilla 487-3, Santiago 1, Chile, <sup>c</sup>Centro para la Investigación Interdisciplinaria Avanzada en Ciencias de los, Materiales, Casilla 233, Santiago 1, Chile, and <sup>d</sup>Facultad de Ciancias Químicas y Farmacéuticas, Universidad de Chile, Casilla 233, Santiago 1, Chile

Correspondence e-mail: baggio@cnea.gov.ar

#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.016 Å R factor = 0.048 wR factor = 0.108 Data-to-parameter ratio = 6.6

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

© 2001 International Union of Crystallography Printed in Great Britain – all rights reserved {2,2'-[1,2-Phenylenebis(nitrilomethylidyne)]diphenolato}copper(II) dichloromethane solvate

The title compound,  $[Cu(C_{20}H_{14}N_2O_2)]\cdot CH_2Cl_2$ , crystallizes as a monomer, with the copper ion in a square-planar environment provided by the tetradentate 2,2'-[1,2-phenylenebis-(nitrilomethylidyne)]diphenolate (L18) ligand. The planar molecules pack parallel to the (101) crystallographic planes. Received 23 November 2000 Accepted 4 December 2000 Online 14 December 2000

### Comment

2,2'-[1,2-Phenylenebis(nitrilomethylidyne)]diphenol (L18) is a well known Schiff base which can act as a tetradentate ligand, and its square-planar copper complex has been described as an isolated molecule in a number of crystalline forms, both on its own (Suresh *et al.*, 1996), (1), as well as stabilized by a variety of different solvates, *e.g.* water [(2); Yao *et al.*, 1997], thiourea [(3); Ferrari *et al.*, 1976] and tetracyanoquinodimethane [(4); Cassoux & Gleizes, 1980]. During the synthesis of a heteronuclear Cu–Gd(L18) complex, we obtained some crystals which happened to be a new unreported form of the Cu(L18) complex, (I), this time stabilized by a molecule of CH<sub>2</sub>Cl<sub>2</sub>, and which we report herein.



The molecular structure of (I) is quite similar to those described previously (Fig. 1), in spite of the very different crystalline forms, and a least-squares fit showed that the extreme differences were found when comparing it with the two independent moieties in (2), with a minimum mean-squares deviation of 0.079 Å and a maximum of 0.158 Å. The largest misfit appeared as deviations of the ligand from its planar character, rather than from the intrinsic bond lengths and angles, which remained fairly stable.

The planar molecules pack parallel to each other and to the (101) planes (Fig. 2), with no obvious interaction besides the usual van der Waals forces.

The CH<sub>2</sub>Cl<sub>2</sub> solvate molecule attaches to the complex through a medium strength  $C-H\cdots O$  interaction [H1*XB* $\cdots$ O1 2.313 (3) Å and C1*X*-H1*XB* $\cdots$ O1 171 (3)°].

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# metal-organic papers



#### Figure 1

The molecular structure of (I) showing the atom numbering and displacement ellipsoids at the 50% probability level.



#### Figure 2

Packing diagram showing the way in which the molecules organize parallel to (101).

## **Experimental**

To a solution containing 1 mmol of  $Cu(NO_3)_2 \cdot 2H_2O$  in  $CH_2Cl_2$ , 1 mmol of  $Gd(NO_3)_2$  and 1 mmol of L18 were added. The  $Cu(L18) \cdot CH_2Cl_2$  complex precipitated as a polycrystalline brown powder, which by filtering was separated from the solution containing the heteronuclear species.

#### Crystal data

$[Cu(C_{20}H_{14}N_2O_2)]\cdot CH_2Cl_2$	$D_x = 1.591 \text{ Mg m}^{-3}$		
$M_r = 462.80$	Mo $K\alpha$ radiation		
Monoclinic, Cc	Cell parameters from 25		
a = 15.787 (3)  Å	reflections		
b = 14.359(3) Å	$\theta = 7.5 - 12.5^{\circ}$		
c = 9.1630 (15)  Å	$\mu = 1.43 \text{ mm}^{-1}$		
$\beta = 111.59 \ (2)^{\circ}$	T = 293 (2) K		
V = 1931.5 (6) Å <sup>3</sup>	Prism, red-brown		
Z = 4	$0.15\times0.12\times0.10~\text{mm}$		

#### Data collection

Siemens R3m diffractometer  $R_{\rm int} = 0.043$  $\omega/2\theta$  scans Absorption correction:  $\psi$  scan (SHELXTL/PC; Sheldrick, 1994)  $T_{\min} = 0.80, \ T_{\max} = 0.84$ 3419 measured reflections 1718 independent reflections 1308 reflections with  $I > 2\sigma(I)$ Refinement Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.048$  $wR(F^2) = 0.108$ S = 1.021718 reflections 262 parameters

H atoms treated by a mixture of independent and constrained refinement

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\theta_{\text{max}} = 25^{\circ}
h = -18 \rightarrow 18
k = -17 \rightarrow 17
l = 0 \rightarrow 10
2 standard reflections
every 98 reflections
intensity decay: 4.8%
w = 1/[\sigma^2(F_o^2) + (0.061P)^2]
where P = (F_o^2 + 2F_c^2)/3
(\Delta/\sigma)_{\text{max}} = 0.002
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 $\begin{array}{l} (\Delta/\sigma)_{\rm max} = 0.002 \\ \Delta\rho_{\rm max} = 0.49 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.34 \ {\rm e} \ {\rm \AA}^{-3} \\ {\rm Absolute \ structure: \ Flack \ (1983);} \\ 119 \ {\rm Friedel \ pairs} \\ {\rm Flack \ parameter} = -0.01 \ (3) \end{array}$ 

Table 1Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D-\mathrm{H}\cdots A$
$C1X - H1XB \cdots O1$	0.89 (4)	2.32 (4)	3.20 (2)	171 (3)

H atoms attached to C atoms were included in idealized positions and refined using a riding scheme (C–H = 0.93 Å), both for the coordinates and the displacement parameters. The H atoms of the CH<sub>2</sub>Cl<sub>2</sub> molecule were found in the difference Fourier map and refined with similarity restraints in the C–H distances (C–H = 0.90 Å).

Data collection: *P3/P4–PC* (Siemens, 1991); cell refinement: *P3/P4–PC*; data reduction: *XDISK* in *SHELXTL/*PC (Sheldrick, 1994); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL/PC* (Sheldrick, 1994); software used to prepare material for publication: *SHELXTL* and *PARST* (Nardelli, 1983).

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