Received 17 October 2006

Accepted 31 October 2006

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Yong Fang, Cui-Ying Huang, Zai-Ming Zhu, Xiao-Li Yu and Wan-Sheng You*

Institute of Chemistry for Functionalized Materials, College of Chemistry and Chemical Engineering, Liaoning Normal University, Dalian 116029, People's Republic of China

Correspondence e-mail: wsyou@Innu.edu.cn

Key indicators

Single-crystal X-ray study T = 150 K Mean σ (C–C) = 0.003 Å R factor = 0.039 wR factor = 0.097 Data-to-parameter ratio = 13.8

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

Dichlorobis{2-[(triphenylmethyl)amino]pyridyl}-cobalt(II)

In the title compound, $[CoCl_2(C_{24}H_{20}N_2)_2]$, Co^{II} is coordinated in an approximately tetrahedral geometry by two Cl atoms and two N atoms of the pyridine rings from two 2-[(triphenylmethyl)imino]pyridyl ligands.

Comment

Transition metal complexes have attracted interest because of their useful properties, such as microporosity, molecular magnetism or linear optical behavior. The title compound, (I) (Fig. 1), is a Co^{II} complex of the N-donor ligand 2-[*N*-(triphenylmethyl)imino]pyridyl. The complex exhibits approximate C_2 point symmetry, and the Co^{II} atom is coordinated in a slightly distorted tetrahedral geometry by atoms Cl1, Cl2, N1 and N2 (Table 1). The dihedral angle between the two pyridyl rings directly coordinated to Co^{II} is 109.8 (1)°. The formation of a four-coordinate complex, rather than a possible six-coordinate one, is attributed to the large volume of the 2-[*N*-(triphenylmethyl)imino]pyridyl ligand.



Experimental

In a 100 ml round-bottomed flask, 2-[N-(triphenyl)imino]pyridine (0.672 g, 2 mmol) was dissolved in THF (20 ml), and then CoCl₂



© 2006 International Union of Crystallography All rights reserved

Figure 1 Molecular structure of (I), showing displacement ellipsoids at the 30% probability level. H atoms have been omitted.

metal-organic papers

(0.13 g, 1 mmol) was added to the solution. The solution was refluxed for 6 h and then stirred overnight. The THF was removed under vacuum and the residue was dissolved in CH_2Cl_2 (20 ml). The filtrate was layered with hexane (20 ml), and blue crystals were obtained after one week. All manipulations were carried out under an atmosphere of Ar gas.

Crystal data

 $\begin{bmatrix} \text{CoCl}_2(\text{C}_{24}\text{H}_{20}\text{N}_2)_2 \end{bmatrix} \\ M_r = 802.67 \\ \text{Monoclinic, } P_{2_1}/n \\ a = 9.920 \text{ (2) Å} \\ b = 22.620 \text{ (5) Å} \\ c = 17.460 \text{ (4) Å} \\ \beta = 97.36 \text{ (3)}^{\circ} \\ V = 3885.6 \text{ (15) Å}^3 \\ \end{bmatrix}$

Data collection

Bruker APEXII CCD diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 2003) $T_{\min} = 0.812, T_{\max} = 0.836$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.039$ $wR(F^2) = 0.098$ S = 1.026823 reflections 496 parameters H-atom parameters constrained Z = 4 $D_x = 1.372 \text{ Mg m}^{-3}$ Mo K α radiation $\mu = 0.62 \text{ mm}^{-1}$ T = 150 (2) K Block, blue $0.35 \times 0.34 \times 0.30 \text{ mm}$

25600 measured reflections 6823 independent reflections 5433 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.081$ $\theta_{\text{max}} = 25.0^{\circ}$

$w = 1/[\sigma^2(F_o^2) + (0.039P)^2]$
+ 2.5598P]
where $P = (F_0^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} = 0.001$
$\Delta \rho_{\rm max} = 0.69 \ {\rm e} \ {\rm \AA}^{-3}$
$\Delta \rho_{\rm min} = -0.52 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (Å, $^\circ).$

Co1-Cl1	2.2415 (8)	Co1-N1	2.054 (2)
Co1-Cl2	2.2394 (8)	Co1-N3	2.0510 (18)
	111.52 (2)		110 (2 (()
CII - CoI - CI2	114.53 (3)	CI2-Co1-N1	110.63 (6)
Cl1-Co1-N1	110.26 (6)	Cl2-Co1-N3	109.08 (6)
Cl1-Co1-N3	109.41 (6)	N1-Co1-N3	102.21 (8)

All H atoms were placed in calculated positions and allowed to ride during subsequent refinement, with C–H = 0.95 Å, N–H = 0.88 Å, and $U_{iso}(H) = 1.2U_{eq}(C,N)$.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

This work was supported by the Education Office Foundation of Liaoning Province (project No. 605 L207).

References

Bruker (1997). SHELXTL. Version 5.10. Bruker AXS Inc., Madison, Wisconsin, USA.

Bruker (2005). APEX2 (Version 1.27) and SAINT. Bruker AXS Inc., Madison, Wisconsin, USA.

Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.

Sheldrick, G. M. (2003). SADABS. Version 2.10. University of Göttingen, Germany.