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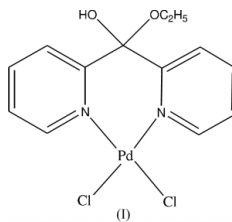
Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.006$ Å
 R factor = 0.032
 wR factor = 0.078
Data-to-parameter ratio = 14.4For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.Dichloro[2,2'-hydroxy(ethoxy)methylene-dipyridine- $\kappa^2\text{N},\text{N}'$]palladium(II)

In the title complex, $[\text{PdCl}_2(\text{C}_{13}\text{H}_{14}\text{N}_2\text{O}_2)]$, the central Pd^{II} atom is bonded to two pyridine N atoms and two terminal Cl atoms. The coordination geometry of the Pd atom is square planar with a slight tetrahedral distortion. The two Pd–N distances are 2.029 (3) and 2.057 (3) Å, and the N–Pd–N angle is 86.56 (13)°. The Pd–Cl distances are 2.2929 (11) and 2.2959 (11) Å, and the Cl–Pd–Cl angle is 90.84 (5)°.

Comment

The condensation products formed when metal complexes of di-2-pyridyl ketone (dpk) are reacted with some nucleophiles, including water, have been reported previously (Annibale *et al.*, 1981; Sommerer *et al.*, 1997). Herein we report the structure of the title compound, (I), which is formed *via* the reaction of dpk with PdCl₂ in CH₃CN/CH₃CH₂OH. In (I), the N1–Pd–N2 angle is narrower than the ideal value (for square-planar coordination) of 90° [86.56 (13)°]. This angle can be compared with corresponding angles in [Pd{dpk-(OH)₂}]Cl₂ and [Pd(2,2'-dipyridyl)]Cl₂, where the N–Pd–N angles are 87.1 (2) (Annibale *et al.*, 1981) and 80.5 (4)° (Beer *et al.*, 1997), respectively. The N2–Pd–N1 [86.56 (13)°], N2–Pd–Cl2 [90.71 (9)°], N1–Pd–Cl1 [91.91 (10)°] and Cl1–Pd–Cl2 [90.84 (5)°] angles approach the ideal value of 90°; this indicates that the coordination geometry of the Pd atom is square planar with a slight tetrahedral distortion.



Experimental

For the preparation of compound (I), *p*-phenylenediamine was combined with dpk in a 1:2 stoichiometric ratio in CH₃CN/CH₃CH₂OH, followed by addition of PdCl₂. The resulting solution was filtered and slow evaporation of the clear filtrate gave clear yellow crystals, suitable for X-ray diffraction studies (yield: 0.07 g, 75%). Analysis calculated for C₁₃H₁₄Cl₂N₂O₂Pd: C 38.43, H 3.48, N 6.90%; found: C 38.23, H 3.09, N 6.68%.

Crystal data

$[\text{PdCl}_2(\text{C}_{13}\text{H}_{14}\text{N}_2\text{O}_2)]$
 $M_r = 407.56$
Monoclinic, $P2_1/n$
 $a = 9.8277$ (1) Å
 $b = 12.0717$ (3) Å
 $c = 12.6885$ (1) Å
 $\beta = 93.162$ (2)°
 $V = 1503.03$ (2) Å³
 $Z = 4$

$D_x = 1.801$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 3244 reflections
 $\theta = 2.3$ – 25.0 °
 $\mu = 1.59$ mm⁻¹
 $T = 293$ (2) K
Column, yellow
 $0.36 \times 0.22 \times 0.20$ mm

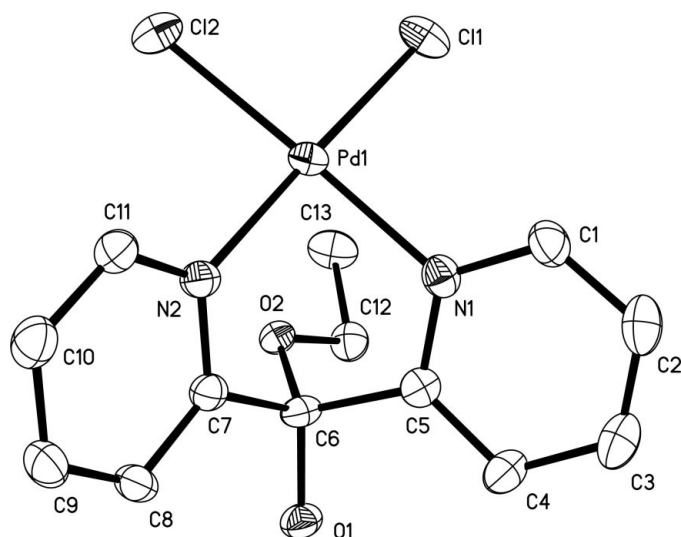


Figure 1
A view of the structure of the title compound, showing 30% probability displacement ellipsoids. H atoms have been omitted for clarity.

Data collection

Siemens SMART CCD
diffractometer
 φ and ω scans
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.558$, $T_{\max} = 0.727$
4510 measured reflections

2600 independent reflections
2270 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.019$
 $\theta_{\text{max}} = 25.0^\circ$
 $h = -9 \rightarrow 11$
 $k = -9 \rightarrow 14$
 $l = -15 \rightarrow 11$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.078$
 $S = 1.15$
2600 reflections
181 parameters
H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0225P)^2 + 2.9262P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.32 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.63 \text{ e } \text{\AA}^{-3}$

Table 1

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

Pd1—N2	2.029 (3)	Pd1—Cl1	2.2929 (11)
Pd1—N1	2.057 (3)	Pd1—Cl2	2.2959 (11)
N2—Pd1—N1	86.56 (13)	N1—Pd1—Cl2	176.93 (10)
N2—Pd1—Cl1	178.32 (10)	Cl1—Pd1—Cl2	90.84 (5)
N1—Pd1—Cl1	91.91 (10)	Cl1—N2—Pd1	122.0 (3)
N2—Pd1—Cl2	90.71 (9)	C7—N2—Pd1	119.1 (3)

H atoms were included in calculated positions and allowed to ride on their parent atoms, with C—H distances in the range 0.93–0.97 \AA .

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1994); data reduction: *SHELXTL* (Bruker, 1999); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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