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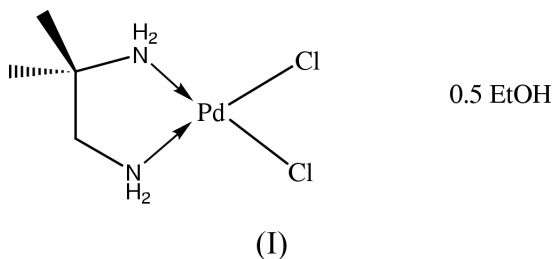
**Key indicators**

Single-crystal X-ray study  
 T = 150 K  
 Mean  $\sigma(\text{C}-\text{C}) = 0.004 \text{ \AA}$   
 Disorder in solvent or counterion  
 R factor = 0.026  
 wR factor = 0.058  
 Data-to-parameter ratio = 24.3

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

**cis-(2-Amino-1,1-dimethylethylamine)dichloro-palladium(II) ethanol hemisolvate**

The title complex,  $[\text{PdCl}_2(\text{C}_4\text{H}_{12}\text{N}_2)] \cdot 0.5\text{C}_2\text{H}_6\text{O}$ , (I), lies in a general position with a disordered molecule of ethanol solvate on a twofold symmetry axis. It adopts the expected square-planar geometry. The electron-donating properties of the  $\text{CMe}_2$  group attached to one N atom are apparent in the slightly shorter corresponding Pd–N distance and slightly longer Pd–Cl bond *trans* to this vector. There is a hydrogen-bonding network involving weak  $\text{N}-\text{H} \cdots \text{Cl}$  and stronger  $\text{O}-\text{H} \cdots \text{N}$  and  $\text{N}-\text{H} \cdots \text{O}$  interactions, in a chain parallel to the *c* axis.



**Experimental**

The title compound was crystallized from ethanol. It is a serendipitous hydrolysis product of a Schiff base palladium(II) complex.

*Crystal data*

$[\text{PdCl}_2(\text{C}_4\text{H}_{12}\text{N}_2)] \cdot 0.5\text{C}_2\text{H}_6\text{O}$   
 $M_r = 288.49$   
 Monoclinic,  $I2/a$   
 $a = 16.435 (5) \text{ \AA}$   
 $b = 7.562 (5) \text{ \AA}$   
 $c = 16.527 (4) \text{ \AA}$   
 $\beta = 90.28 (2)^\circ$   
 $V = 2053.9 (16) \text{ \AA}^3$   
 $Z = 8$

$D_x = 1.866 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation  
 Cell parameters from 3005 reflections  
 $\theta = 2-30.0^\circ$   
 $\mu = 2.27 \text{ mm}^{-1}$   
 $T = 150 (2) \text{ K}$   
 Rectangular block, yellow-green  
 $0.20 \times 0.08 \times 0.08 \text{ mm}$

*Data collection*

KappaCCD diffractometer  
 CCD rotation images, thick-slice scans  
 Absorption correction: multi-scan (Blessing, 1995)  
 $T_{\min} = 0.727$ ,  $T_{\max} = 0.834$   
 15 940 measured reflections

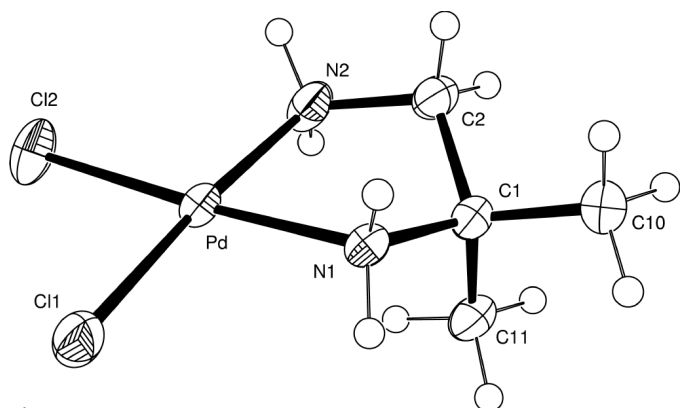
2993 independent reflections  
 2577 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.045$   
 $\theta_{\text{max}} = 30.0^\circ$   
 $h = -23 \rightarrow 23$   
 $k = -9 \rightarrow 10$   
 $l = -23 \rightarrow 23$

*Refinement*

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.026$   
 $wR(F^2) = 0.058$   
 $S = 1.06$   
 2993 reflections  
 123 parameters  
 Only H-atom  $U$ 's refined

$w = 1/[\sigma^2(F_o^2) + (0.0222P)^2 + 4.1385P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.002$   
 $\Delta\rho_{\text{max}} = 0.67 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.64 \text{ e \AA}^{-3}$

Received 20 September 2001  
 Accepted 26 September 2001  
 Online 6 October 2001



**Figure 1**  
View of (I) (70% probability displacement ellipsoids). The disordered solvent molecule is not shown.

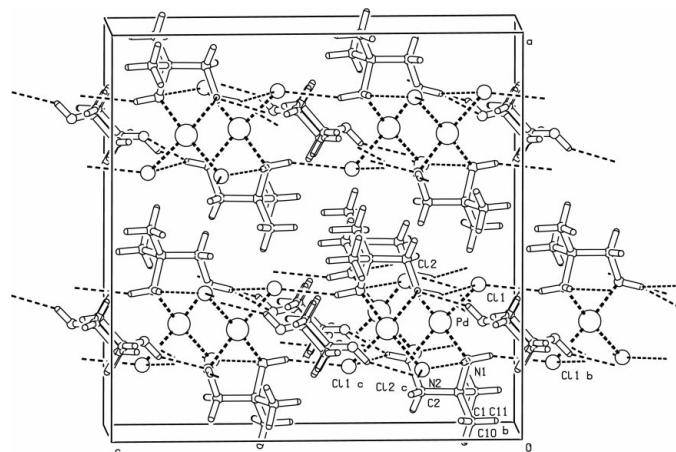
**Table 1**  
Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

Pd—N1	2.0197 (19)	N2—C2	1.488 (3)
Pd—N2	2.036 (2)	C1—C11	1.523 (4)
Pd—Cl1	2.3068 (8)	C1—C10	1.526 (3)
Pd—Cl2	2.3213 (9)	C1—C2	1.527 (3)
N1—C1	1.501 (3)		
N1—Pd—N2	83.28 (8)	N1—Pd—Cl2	175.58 (5)
N1—Pd—Cl1	90.69 (6)	N2—Pd—Cl2	92.38 (6)
N2—Pd—Cl1	173.96 (6)	Cl1—Pd—Cl2	93.66 (3)

**Table 2**  
Hydrogen-bonding geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H12 $\cdots$ Cl2 <sup>i</sup>	0.92	2.59	3.405 (3)	148
N1—H12 $\cdots$ O100 <sup>ii</sup>	0.92	2.64	3.280 (9)	127
N1—H11 $\cdots$ Cl1 <sup>iii</sup>	0.92	2.55	3.443 (2)	163
N2—H21 $\cdots$ Cl2 <sup>iv</sup>	0.92	2.44	3.352 (3)	170
N2—H22 $\cdots$ O100 <sup>v</sup>	0.92	2.07	2.873 (6)	145
N2—H22 $\cdots$ Cl1 <sup>i</sup>	0.92	2.81	3.538 (3)	137
O100—H100 $\cdots$ N2 <sup>vi</sup>	0.84	2.15	2.873 (6)	144

Symmetry codes: (i)  $\frac{3}{2}-x, \frac{1}{2}-y, \frac{1}{2}-z$ ; (ii)  $\frac{3}{2}-x, \frac{3}{2}-y, \frac{1}{2}-z$ ; (iii)  $\frac{3}{2}-x, y, -z$ ; (iv)  $\frac{3}{2}-x, -\frac{1}{2}-y, \frac{1}{2}-z$ ; (v)  $x, y-1, z$ ; (vi)  $x, 1+y, z$ .



**Figure 2**  
Unit-cell packing diagram of (I) viewed along the  $b$  axis.

Data collection: *COLLECT* (Nonius, 1997–2000); cell refinement: *HKL SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *HKL DENZO* (Otwinowski & Minor, 1997) and *WinGX* (Farrugia, 1999); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* for Windows (Farrugia, 1997); software used to prepare material for publication: *WinGX* publication routines.

The EPSRC is thanked for a grant towards the purchase of a KappaCCD diffractometer and an Oxford Cryosystems Cryostream cooler.

## References

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