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trans-Dichloridobis(3,4-dimethylpyridine)platinum(II)Alexander N. Chernyshev,^a Nadezhda A. Bokach,^a
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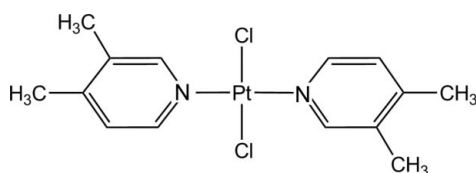
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Key indicators: single-crystal X-ray study; $T = 120$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.015; wR factor = 0.030; data-to-parameter ratio = 24.2.

In the title compound, $\text{trans}[\text{PtCl}_2(\text{C}_7\text{H}_9\text{N})_2]$, the Pt^{II} atom is located on an inversion center and is coordinated by two 3,4-dimethylpyridine ligands and two chloride ligands, resulting in a typical slightly distorted square-planar geometry. The crystallographic inversion centre forces the value of the C—N—N—C torsion angle to be linear and the 3,4-dimethylpyridine ligands to be coplanar.

Related literature

For related complexes see: Tessier & Rochon (1999); Eremenko *et al.* (1997); Shaver *et al.* (2000); Zordan *et al.* (2005); Rochon *et al.* (1996); Colamarino & Orioli (1975). For the geometry of the pyridine ligand, see: Bond & Davies (2002). For related literature, see: Orpen *et al.* (1989).



Experimental

Crystal data

$[\text{PtCl}_2(\text{C}_7\text{H}_9\text{N})_2]$
 $M_r = 480.29$
 Monoclinic, $P2_1/n$
 $a = 7.9763$ (5) Å
 $b = 7.1102$ (3) Å
 $c = 13.3586$ (7) Å
 $\beta = 98.247$ (5)°

$V = 749.77$ (7) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 9.70$ mm⁻¹
 $T = 120$ (2) K
 $0.21 \times 0.20 \times 0.10$ mm

Data collection

Nonius KappaCCD diffractometer
 Absorption correction: multi-scan
 (*SADABS*; Sheldrick, 2003)
 $T_{\min} = 0.201$, $T_{\max} = 0.381$
 17165 measured reflections
 2177 independent reflections
 1705 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.033$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.015$
 $wR(F^2) = 0.030$
 $S = 1.08$
 2177 reflections
 90 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.67$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.78$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Pt1—N1	2.0148 (18)	Pt1—Cl1	2.2901 (6)
N1—Pt1—Cl1	89.85 (6)		

Data collection: *COLLECT* (Bruker–Nonius, 2004); cell refinement: *EVALCCD* (Duisenberg *et al.*, 2003); data reduction: *EVALCCD* ; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *SHELXL97* .

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: KJ2108).

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supporting information

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***trans*-Dichloridobis(3,4-dimethylpyridine)platinum(II)**

Alexander N. Chernyshev, Nadezhda A. Bokach, Youlia A. Izotova and Matti Haukka

S1. Comment

The complex *trans*-[PtCl₂(C₇H₉N)₂] has an inversion symmetry and the Pt^{II} atom is situated at an inversion center and it is coordinated by two 3,4-dimethylpyridine ligands and two chloro ligands and exhibit *trans* configuration. Such arrangement of ligands leads to the square planar geometry. In the coordination polyhedron, all angles are very close to the ideal value of 90°. The crystallographic inversion centre forces the value of the torsion angle C1—N1—N1ⁱ—Cⁱ (symmetry operation *i*: -x, -y, -z) to be 180° and the the 3,4-dimethyl-pyridine ligands to be coplanar.

The geometry of 3,4-dimethylpyridine ligands resembles the geometry of the uncoordinated 3,4-dimethylpyridine, *i.e.* the C—C and C—N bond distances and angles in the coordinated 3,4-dimethylpyridine agree well with the expected value (Bond, Davies, 2002). The bond distance Pt—N (2.0148 (18) Å) is similar to the Pt—N bond lengths in other related compounds (Orpen *et al.*, 1989). The Pt—Cl bond lengths agree well with the reported values (See Table 2).

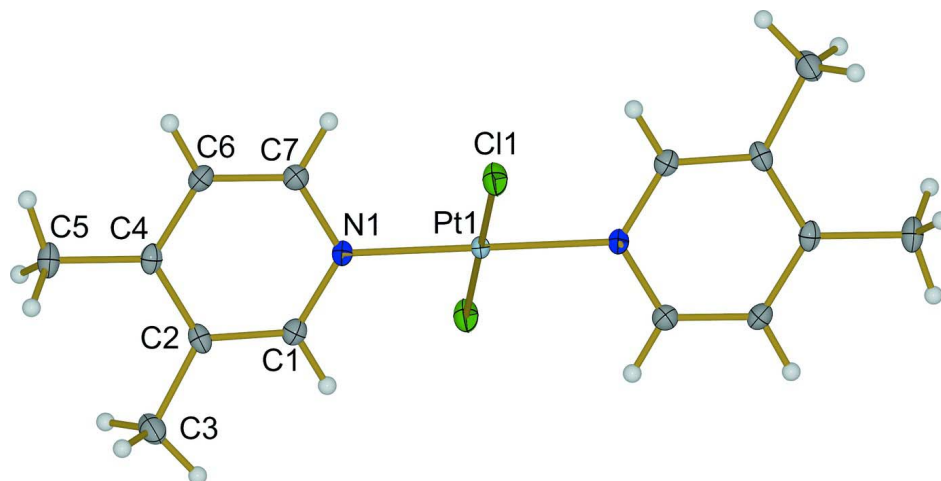
All *trans*-[PtCl₂L₂] complexes given in Table 2 have the same coordination environment as in the title compound. Indeed, they are square-planar and their pyridine rings lie in the same plane. The N—Pt—N and Cl—Pt—Cl angles in all observed compounds are equal to 180°, the angles N—Pt—Cl are very close to 90°.

S2. Experimental

3,4-dimethylpyridine (1 ml) was added to the powder of K₂[PtCl₄] (0.2 g) and the resulting mixture was heated to 150°C until the complete evaporation of the 3,4-dimethylpyridine. The resulting complex was recrystallized from CHCl₃ (yield 92%). Crystals were obtained by slow evaporation of CHCl₃ solution. Anal. calc. for C₁₄H₁₈N₂Cl₂Pt: C, 35.01; H, 3.78; N, 5.83%. Found: C, 35.30; H, 3.96; N, 5.54%.

S3. Refinement

Hydrogen atoms were positioned geometrically and constrained to ride on their parent atoms, with C—H = 0.95–0.98 Å, and $U_{iso} = 1.2–1.5 U_{eq}(\text{parent atom})$. The highest peak is located 0.87 Å from atom Pt1 and the deepest hole is located 0.83 Å from atom Pt1.

**Figure 1**

The molecular structure of the title molecule, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

trans-Dichloridobis(3,4-dimethylpyridine)platinum(II)

Crystal data

[PtCl₂(C₇H₉N)₂]

$M_r = 480.29$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 7.9763 (5) \text{ \AA}$

$b = 7.1102 (3) \text{ \AA}$

$c = 13.3586 (7) \text{ \AA}$

$\beta = 98.247 (5)^\circ$

$V = 749.77 (7) \text{ \AA}^3$

$Z = 2$

$F(000) = 456$

$D_x = 2.127 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 2339 reflections

$\theta = 1.0\text{--}20.0^\circ$

$\mu = 9.70 \text{ mm}^{-1}$

$T = 120 \text{ K}$

Block, pale yellow

$0.21 \times 0.20 \times 0.10 \text{ mm}$

Data collection

Nonius KappaCCD
diffractometer

Radiation source: fine-focus sealed tube

Horizontally mounted graphite crystal
monochromator

Detector resolution: 9 pixels mm^{-1} pixels mm^{-1}

φ scans and ω scans with κ offset

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 2003)

$T_{\min} = 0.201$, $T_{\max} = 0.381$

17165 measured reflections

2177 independent reflections

1705 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.033$

$\theta_{\max} = 30.0^\circ$, $\theta_{\min} = 2.8^\circ$

$h = -11 \rightarrow 10$

$k = -10 \rightarrow 9$

$l = -18 \rightarrow 18$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.015$

$wR(F^2) = 0.030$

$S = 1.08$

2177 reflections

90 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0064P)^2 + 0.7739P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$

$$\Delta\rho_{\max} = 0.67 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.78 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Pt1	0.0000	0.0000	0.0000	0.01078 (3)
Cl1	-0.12594 (8)	-0.27477 (8)	0.03883 (4)	0.01897 (11)
N1	0.0259 (2)	0.0765 (3)	0.14663 (13)	0.0127 (4)
C1	-0.0247 (3)	0.2464 (3)	0.17546 (16)	0.0143 (4)
H1	-0.0716	0.3324	0.1246	0.017*
C2	-0.0115 (3)	0.3015 (3)	0.27567 (16)	0.0141 (4)
C3	-0.0729 (3)	0.4918 (3)	0.30137 (17)	0.0227 (5)
H3A	-0.1179	0.5582	0.2390	0.034*
H3B	0.0215	0.5637	0.3379	0.034*
H3C	-0.1623	0.4781	0.3440	0.034*
C4	0.0634 (3)	0.1776 (3)	0.35029 (16)	0.0138 (4)
C5	0.0883 (3)	0.2318 (4)	0.45953 (16)	0.0188 (5)
H5A	0.1668	0.3383	0.4702	0.028*
H5B	0.1353	0.1249	0.5006	0.028*
H5C	-0.0209	0.2676	0.4795	0.028*
C6	0.1162 (3)	0.0044 (4)	0.31969 (15)	0.0161 (4)
H6	0.1677	-0.0824	0.3688	0.019*
C7	0.0948 (3)	-0.0435 (3)	0.21858 (17)	0.0163 (5)
H7	0.1297	-0.1645	0.1994	0.020*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Pt1	0.01278 (5)	0.01094 (5)	0.00822 (5)	0.00154 (6)	0.00017 (3)	-0.00158 (5)
Cl1	0.0265 (3)	0.0165 (2)	0.0140 (2)	-0.0048 (2)	0.0032 (2)	-0.0015 (2)
N1	0.0139 (9)	0.0141 (8)	0.0096 (8)	0.0001 (8)	0.0003 (7)	-0.0017 (7)
C1	0.0145 (10)	0.0156 (11)	0.0126 (9)	-0.0004 (9)	0.0014 (8)	0.0006 (8)
C2	0.0168 (11)	0.0133 (11)	0.0132 (10)	-0.0030 (9)	0.0050 (8)	-0.0033 (8)
C3	0.0370 (13)	0.0158 (10)	0.0165 (10)	0.0026 (13)	0.0077 (9)	-0.0002 (11)
C4	0.0122 (10)	0.0181 (11)	0.0111 (10)	-0.0041 (9)	0.0019 (8)	-0.0013 (8)
C5	0.0222 (12)	0.0232 (12)	0.0109 (10)	-0.0042 (10)	0.0018 (9)	-0.0026 (9)
C6	0.0168 (9)	0.0174 (10)	0.0131 (9)	0.0019 (11)	-0.0013 (7)	0.0011 (11)

C7	0.0166 (11)	0.0173 (12)	0.0147 (10)	0.0027 (8)	0.0008 (8)	-0.0010 (8)
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Geometric parameters (Å, °)

Pt1—N1 ⁱ	2.0148 (18)	C3—H3B	0.9800
Pt1—N1	2.0148 (18)	C3—H3C	0.9800
Pt1—Cl1 ⁱ	2.2901 (6)	C4—C6	1.382 (3)
Pt1—Cl1	2.2901 (6)	C4—C5	1.495 (3)
N1—C7	1.343 (3)	C5—H5A	0.9800
N1—C1	1.347 (3)	C5—H5B	0.9800
C1—C2	1.384 (3)	C5—H5C	0.9800
C1—H1	0.9500	C6—C7	1.380 (3)
C2—C4	1.398 (3)	C6—H6	0.9500
C2—C3	1.495 (3)	C7—H7	0.9500
C3—H3A	0.9800		
N1 ⁱ —Pt1—N1	180.0	C2—C3—H3C	109.5
N1 ⁱ —Pt1—Cl1 ⁱ	89.85 (6)	H3A—C3—H3C	109.5
N1—Pt1—Cl1 ⁱ	90.15 (6)	H3B—C3—H3C	109.5
N1 ⁱ —Pt1—Cl1	90.15 (6)	C6—C4—C2	117.88 (19)
N1—Pt1—Cl1	89.85 (6)	C6—C4—C5	121.0 (2)
Cl1 ⁱ —Pt1—Cl1	180.0	C2—C4—C5	121.1 (2)
C7—N1—C1	118.30 (19)	C4—C5—H5A	109.5
C7—N1—Pt1	119.91 (15)	C4—C5—H5B	109.5
C1—N1—Pt1	121.79 (15)	H5A—C5—H5B	109.5
N1—C1—C2	123.1 (2)	C4—C5—H5C	109.5
N1—C1—H1	118.5	H5A—C5—H5C	109.5
C2—C1—H1	118.5	H5B—C5—H5C	109.5
C1—C2—C4	118.5 (2)	C7—C6—C4	120.6 (2)
C1—C2—C3	119.8 (2)	C7—C6—H6	119.7
C4—C2—C3	121.78 (19)	C4—C6—H6	119.7
C2—C3—H3A	109.5	N1—C7—C6	121.6 (2)
C2—C3—H3B	109.5	N1—C7—H7	119.2
H3A—C3—H3B	109.5	C6—C7—H7	119.2

Symmetry code: (i) $-x, -y, -z$.