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Dichlorido[2-(phenyliminomethyl)-quinoline-*N,N'*]palladium(II)William M. Motswainyana,^a Martin O. Onani^{a*} and Abram M. Madiehe^b

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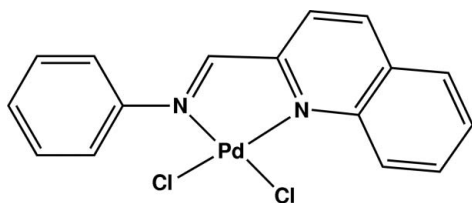
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Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.030; wR factor = 0.063; data-to-parameter ratio = 17.9.

In the title complex, $[\text{PdCl}_2(\text{C}_{16}\text{H}_{12}\text{N}_2)]$, the Pd^{II} ion is coordinated by two N atoms [$\text{Pd}-\text{N}$ 2.039 (2), 2.073 (2) Å] from a bidentate ligand and two chloride anions [$\text{Pd}-\text{Cl}$ 2.2655 (7), 2.2991 (7) Å] in a distorted square-planar geometry. In the crystal, $\pi-\pi$ interactions between the six-membered rings of the quinoline fragments [centroid-centroid distances = 3.815 (5), 3.824 (5) Å] link two molecules into centrosymmetric dimers.

Related literature

For the synthesis of quinolyl-imine ligands and their transition metal-based complexes, see: Ardizzoia *et al.* (2009); Tianpengfei *et al.* (2011); Wei *et al.* (2009). For related structures, see: Motswainyana *et al.* (2011); Onani & Motswainyana (2011); Massa & Dehghampour (2009); Keter *et al.* (2008); Singh *et al.* (2007); Doherty *et al.* (2002).



Experimental

Crystal data

 $[\text{PdCl}_2(\text{C}_{16}\text{H}_{12}\text{N}_2)]$ $M_r = 409.58$

Monoclinic, $P2_1/c$
 $a = 10.0980$ (4) Å
 $b = 15.8936$ (6) Å
 $c = 10.0010$ (3) Å
 $\beta = 112.005$ (2)°
 $V = 1488.17$ (9) Å³

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 1.60$ mm⁻¹
 $T = 173$ K
 $0.23 \times 0.16 \times 0.03$ mm

Data collection

Bruker SMART APEX CCD diffractometer
Absorption correction: multi-scan (SADABS; Bruker, 2009)
 $T_{\text{min}} = 0.710$, $T_{\text{max}} = 0.954$

49839 measured reflections
3400 independent reflections
2575 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.086$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.030$
 $wR(F^2) = 0.063$
 $S = 1.05$
3400 reflections

190 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 1.00$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.59$ e Å⁻³

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: X-SEED (Barbour, 2001); software used to prepare material for publication: SHELXL97.

Financial support from the NRF (Thuthuka) and University of the Western Cape Senate Research is greatly acknowledged. We also thank Professor Roger A. Lalancette for resolving the symmetry-related geometries.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CV5251).

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supporting information

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Dichlorido[2-(phenyliminomethyl)quinoline-*N,N'*]palladium(II)**William M. Motswainyana, Martin O. Onani and Abram M. Madiehe****S1. Comment**

The transition metal-diimine ligands are well known and have been used extensively as catalyst stabilisers in many reported catalytic processes (Motswainyana *et al.*, 2011; Massa *et al.*, 2009; Tiangpengfei *et al.*, 2011; Ardizzoia *et al.*, 2009). Several of these complexes, besides being heavily utilized in the catalytic field, equally double up as therapeutic agents (Keter *et al.*, 2008; Singh *et al.*, 2007).

The asymmetric unit of the title compound is shown in Fig. 1. In the title compound, the Pd^{II} ion is bidentately coordinated to two N atoms of the 2-quinolybenzylimine ligand and two chloride anions. The complex forms a distorted square planar geometry around the central metal. It is notable that the bond length of the Pd—Cl bond *trans* to the quinoly-N atom are similar indicating lack of *trans* influence. There are two π - π interactions between the quinoline ring systems as indicated by the interplanar centroid-centroid distances of 3.815 (5) and 3.824 (5) linking two molecules into a centrosymmetric dimers.

S2. Experimental

To a suspension of PdCl₂(cod) (0.0925 g, 0.324 mmol) in CH₂Cl₂ (5 ml) was added a solution of (2-quinolybenzyl)imine (0.0748 g, 0.322 mmol) in CH₂Cl₂ (10 ml). The yellow solution was stirred under reflux for 4 h, resulting in the formation of yellow precipitate. The precipitate was filtered to obtain a pure yellow solid. Recrystallization from a mixture of a minimum of CH₂Cl₂ and an excess of C₆H₁₄ solution gave single crystals suitable for X-ray diffraction studies. The product yield was 82%.

S3. Refinement

All hydrogen atoms were placed in idealized positions, and refined as riding, with $U_{\text{iso}} = 1.2 U_{\text{eq}}$ of the parent atoms.

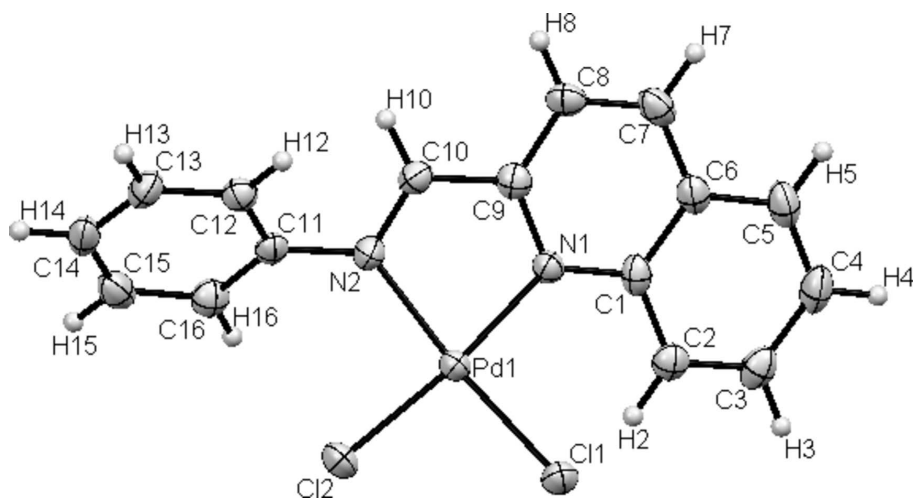


Figure 1

Molecular structure of the title compound showing the atomic numbering and 35% probability displacement ellipsoids.

Dichlorido[2-(phenyliminomethyl)quinoline-*N,N'*]palladium(II)

Crystal data

[PdCl₂(C₁₆H₁₂N₂)]

M_r = 409.58

Monoclinic, *P*2₁/*c*

Hall symbol: -*P* 2ybc

a = 10.0980 (4) Å

b = 15.8936 (6) Å

c = 10.0010 (3) Å

β = 112.005 (2)°

V = 1488.17 (9) Å³

Z = 4

F(000) = 808

D_x = 1.828 Mg m⁻³

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 49839 reflections

θ = 3.4–27.5°

μ = 1.60 mm⁻¹

T = 173 K

Plate, orange

0.23 × 0.16 × 0.03 mm

Data collection

Bruker SMART APEX CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

1.0° φ scans and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2009)

T_{min} = 0.710, *T_{max}* = 0.954

49839 measured reflections

3400 independent reflections

2575 reflections with *I* > 2σ(*I*)

R_{int} = 0.086

θ_{max} = 27.5°, θ_{min} = 3.4°

h = -13→13

k = -20→20

l = -12→12

Refinement

Refinement on *F*²

Least-squares matrix: full

R [*F*² > 2σ(*F*²)] = 0.030

wR (*F*²) = 0.063

S = 1.05

3400 reflections

190 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/[σ²(*F_o*²) + (0.0267*P*)² + 0.1302*P*]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = 0.001

Δρ_{max} = 1.00 e Å⁻³

Δρ_{min} = -0.59 e Å⁻³

Special details

Experimental. 'crystal mounted on a cryoloop'

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
Pd1	0.69195 (2)	0.038621 (12)	0.43401 (2)	0.02445 (9)
Cl1	0.65380 (7)	0.18083 (4)	0.39731 (8)	0.03194 (18)
Cl2	0.89245 (7)	0.06851 (4)	0.62746 (8)	0.03355 (18)
N1	0.5127 (2)	-0.00321 (13)	0.2642 (2)	0.0243 (5)
N2	0.6938 (2)	-0.08478 (13)	0.4906 (2)	0.0264 (5)
C1	0.4308 (3)	0.03303 (15)	0.1353 (3)	0.0268 (6)
C2	0.4815 (3)	0.10349 (15)	0.0822 (3)	0.0289 (6)
H2	0.5729	0.1264	0.1359	0.035*
C3	0.3981 (3)	0.13838 (16)	-0.0468 (3)	0.0333 (7)
H3	0.4329	0.1855	-0.0823	0.040*
C4	0.2618 (3)	0.10609 (17)	-0.1284 (3)	0.0387 (7)
H4	0.2045	0.1326	-0.2162	0.046*
C5	0.2123 (3)	0.03718 (18)	-0.0818 (3)	0.0374 (8)
H5	0.1211	0.0150	-0.1384	0.045*
C6	0.2947 (3)	-0.00179 (17)	0.0495 (3)	0.0295 (7)
C7	0.2488 (3)	-0.07536 (17)	0.0973 (3)	0.0338 (7)
H7	0.1566	-0.0977	0.0446	0.041*
C8	0.3373 (3)	-0.11471 (16)	0.2201 (3)	0.0307 (7)
H8	0.3102	-0.1663	0.2506	0.037*
C9	0.4687 (3)	-0.07709 (16)	0.2995 (3)	0.0270 (6)
C10	0.5726 (3)	-0.11936 (16)	0.4231 (3)	0.0269 (6)
H10	0.5507	-0.1724	0.4536	0.032*
C11	0.7958 (3)	-0.12731 (15)	0.6125 (3)	0.0251 (6)
C12	0.7546 (3)	-0.16047 (15)	0.7185 (3)	0.0277 (6)
H12	0.6589	-0.1544	0.7122	0.033*
C13	0.8534 (3)	-0.20272 (16)	0.8341 (3)	0.0327 (7)
H13	0.8264	-0.2240	0.9090	0.039*
C14	0.9911 (3)	-0.21379 (16)	0.8403 (3)	0.0361 (7)
H14	1.0581	-0.2444	0.9178	0.043*
C15	1.0315 (3)	-0.18044 (17)	0.7341 (3)	0.0368 (7)
H15	1.1266	-0.1882	0.7391	0.044*
C16	0.9356 (3)	-0.13606 (16)	0.6209 (3)	0.0313 (7)
H16	0.9647	-0.1117	0.5494	0.038*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Pd1	0.02349 (14)	0.02256 (14)	0.02361 (14)	-0.00179 (8)	0.00458 (10)	0.00024 (9)
Cl1	0.0326 (4)	0.0229 (3)	0.0349 (4)	-0.0023 (3)	0.0063 (3)	-0.0004 (3)
Cl2	0.0274 (4)	0.0340 (4)	0.0309 (4)	-0.0049 (3)	0.0014 (3)	-0.0011 (3)
N1	0.0253 (12)	0.0223 (12)	0.0220 (13)	-0.0001 (10)	0.0051 (10)	-0.0013 (10)
N2	0.0252 (13)	0.0244 (12)	0.0275 (13)	0.0008 (10)	0.0075 (11)	0.0050 (11)
C1	0.0287 (16)	0.0281 (15)	0.0206 (15)	0.0077 (12)	0.0057 (12)	-0.0020 (12)
C2	0.0332 (16)	0.0247 (15)	0.0274 (16)	0.0014 (12)	0.0098 (13)	-0.0038 (13)
C3	0.0464 (18)	0.0257 (15)	0.0267 (17)	0.0063 (13)	0.0125 (15)	-0.0004 (13)
C4	0.0468 (19)	0.0346 (17)	0.0262 (17)	0.0126 (14)	0.0039 (15)	0.0037 (14)
C5	0.0313 (17)	0.0417 (19)	0.0287 (18)	0.0085 (13)	-0.0008 (14)	-0.0035 (14)
C6	0.0290 (16)	0.0280 (16)	0.0267 (17)	0.0046 (12)	0.0049 (13)	-0.0055 (13)
C7	0.0272 (16)	0.0365 (17)	0.0314 (18)	-0.0032 (13)	0.0037 (14)	-0.0089 (14)
C8	0.0325 (16)	0.0267 (15)	0.0311 (17)	-0.0062 (12)	0.0098 (14)	-0.0066 (13)
C9	0.0298 (16)	0.0273 (15)	0.0232 (16)	0.0008 (12)	0.0092 (13)	-0.0032 (12)
C10	0.0320 (16)	0.0222 (14)	0.0260 (16)	-0.0007 (12)	0.0105 (13)	0.0006 (12)
C11	0.0276 (15)	0.0183 (13)	0.0250 (16)	0.0004 (11)	0.0048 (13)	-0.0029 (12)
C12	0.0273 (15)	0.0244 (14)	0.0303 (17)	-0.0005 (12)	0.0093 (13)	-0.0030 (13)
C13	0.0423 (18)	0.0248 (15)	0.0266 (17)	0.0021 (13)	0.0078 (14)	0.0043 (13)
C14	0.0371 (18)	0.0234 (15)	0.0330 (19)	0.0044 (13)	-0.0039 (14)	-0.0020 (13)
C15	0.0273 (16)	0.0316 (16)	0.045 (2)	0.0008 (13)	0.0059 (15)	-0.0090 (15)
C16	0.0343 (17)	0.0280 (15)	0.0309 (18)	0.0016 (13)	0.0114 (14)	-0.0025 (13)

Geometric parameters (\AA , $^\circ$)

Pd1—N2	2.039 (2)	C6—C7	1.406 (4)
Pd1—N1	2.073 (2)	C7—C8	1.370 (4)
Pd1—Cl2	2.2655 (7)	C7—H7	0.9500
Pd1—Cl1	2.2991 (7)	C8—C9	1.400 (4)
N1—C9	1.348 (3)	C8—H8	0.9500
N1—C1	1.371 (3)	C9—C10	1.452 (4)
N2—C10	1.279 (3)	C10—H10	0.9500
N2—C11	1.436 (3)	C11—C12	1.380 (4)
C1—C2	1.415 (3)	C11—C16	1.390 (4)
C1—C6	1.431 (4)	C12—C13	1.385 (4)
C2—C3	1.366 (4)	C12—H12	0.9500
C2—H2	0.9500	C13—C14	1.380 (4)
C3—C4	1.407 (4)	C13—H13	0.9500
C3—H3	0.9500	C14—C15	1.379 (4)
C4—C5	1.357 (4)	C14—H14	0.9500
C4—H4	0.9500	C15—C16	1.377 (4)
C5—C6	1.407 (4)	C15—H15	0.9500
C5—H5	0.9500	C16—H16	0.9500
N2—Pd1—N1	80.41 (8)	C8—C7—C6	119.8 (3)
N2—Pd1—Cl2	93.00 (6)	C8—C7—H7	120.1

N1—Pd1—C12	173.36 (6)	C6—C7—H7	120.1
N2—Pd1—C11	166.73 (6)	C7—C8—C9	118.4 (3)
N1—Pd1—C11	98.16 (6)	C7—C8—H8	120.8
C12—Pd1—C11	88.45 (3)	C9—C8—H8	120.8
C9—N1—C1	118.1 (2)	N1—C9—C8	124.0 (2)
C9—N1—Pd1	109.69 (16)	N1—C9—C10	115.0 (2)
C1—N1—Pd1	132.02 (17)	C8—C9—C10	120.9 (2)
C10—N2—C11	119.0 (2)	N2—C10—C9	119.6 (2)
C10—N2—Pd1	111.05 (17)	N2—C10—H10	120.2
C11—N2—Pd1	128.13 (16)	C9—C10—H10	120.2
N1—C1—C2	120.7 (2)	C12—C11—C16	120.5 (2)
N1—C1—C6	120.5 (2)	C12—C11—N2	120.4 (2)
C2—C1—C6	118.8 (2)	C16—C11—N2	119.2 (2)
C3—C2—C1	119.6 (3)	C11—C12—C13	119.8 (2)
C3—C2—H2	120.2	C11—C12—H12	120.1
C1—C2—H2	120.2	C13—C12—H12	120.1
C2—C3—C4	121.4 (3)	C14—C13—C12	119.9 (3)
C2—C3—H3	119.3	C14—C13—H13	120.1
C4—C3—H3	119.3	C12—C13—H13	120.1
C5—C4—C3	120.1 (3)	C15—C14—C13	120.0 (3)
C5—C4—H4	120.0	C15—C14—H14	120.0
C3—C4—H4	120.0	C13—C14—H14	120.0
C4—C5—C6	120.7 (3)	C16—C15—C14	120.7 (3)
C4—C5—H5	119.7	C16—C15—H15	119.6
C6—C5—H5	119.7	C14—C15—H15	119.6
C7—C6—C5	121.9 (2)	C15—C16—C11	119.1 (3)
C7—C6—C1	118.9 (2)	C15—C16—H16	120.5
C5—C6—C1	119.3 (3)	C11—C16—H16	120.5
N2—Pd1—N1—C9	17.70 (17)	C2—C1—C6—C5	-3.0 (4)
C12—Pd1—N1—C9	24.5 (6)	C5—C6—C7—C8	174.9 (3)
C11—Pd1—N1—C9	-148.95 (16)	C1—C6—C7—C8	-3.5 (4)
N2—Pd1—N1—C1	-168.0 (2)	C6—C7—C8—C9	3.9 (4)
C12—Pd1—N1—C1	-161.1 (4)	C1—N1—C9—C8	-7.0 (4)
C11—Pd1—N1—C1	25.4 (2)	Pd1—N1—C9—C8	168.3 (2)
N1—Pd1—N2—C10	-17.17 (18)	C1—N1—C9—C10	169.1 (2)
C12—Pd1—N2—C10	163.62 (18)	Pd1—N1—C9—C10	-15.7 (3)
C11—Pd1—N2—C10	67.7 (4)	C7—C8—C9—N1	1.5 (4)
N1—Pd1—N2—C11	178.4 (2)	C7—C8—C9—C10	-174.4 (2)
C12—Pd1—N2—C11	-0.8 (2)	C11—N2—C10—C9	-179.8 (2)
C11—Pd1—N2—C11	-96.8 (3)	Pd1—N2—C10—C9	14.1 (3)
C9—N1—C1—C2	-170.4 (2)	N1—C9—C10—N2	1.4 (4)
Pd1—N1—C1—C2	15.6 (4)	C8—C9—C10—N2	177.6 (2)
C9—N1—C1—C6	7.1 (4)	C10—N2—C11—C12	-47.7 (3)
Pd1—N1—C1—C6	-166.84 (18)	Pd1—N2—C11—C12	115.7 (2)
N1—C1—C2—C3	179.8 (2)	C10—N2—C11—C16	131.1 (3)
C6—C1—C2—C3	2.2 (4)	Pd1—N2—C11—C16	-65.5 (3)
C1—C2—C3—C4	0.4 (4)	C16—C11—C12—C13	0.0 (4)

C2—C3—C4—C5	-2.2 (4)	N2—C11—C12—C13	178.8 (2)
C3—C4—C5—C6	1.3 (4)	C11—C12—C13—C14	-2.1 (4)
C4—C5—C6—C7	-177.2 (3)	C12—C13—C14—C15	2.2 (4)
C4—C5—C6—C1	1.3 (4)	C13—C14—C15—C16	-0.1 (4)
N1—C1—C6—C7	-2.1 (4)	C14—C15—C16—C11	-1.9 (4)
C2—C1—C6—C7	175.5 (2)	C12—C11—C16—C15	1.9 (4)
N1—C1—C6—C5	179.4 (2)	N2—C11—C16—C15	-176.8 (2)
