

Supporting Information for

Complexes of 2,5-bis(α -pyridyl)pyrrolate with
Pd(II) and Pt(II): A mono-anionic iso π -electron
ligand analog of terpyridine

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Density Functional Theory. Density Functional Theory calculations (B3LYP) were carried out using Gaussian '03, Gaussian '09 or the GAMESS suite of quantum programs. In Gaussian, structures were first minimized using the 3-21G basis set, and the resulting coordinates used as the input file for higher level bases. For the calculation of H(PDP), the basis set was 6-31G*, and for the calculation of **1-Cl**, a mixed pseudopotential basis set, 6-31G* (C, H, N) and 3-21G (Pd, Cl) were used. The MIDI basis set was used for the determination of the structure of **1-Cl** using the GAMESS package for comparison to the results from the mixed basis set. Results were qualitatively similar (Fig. S1).

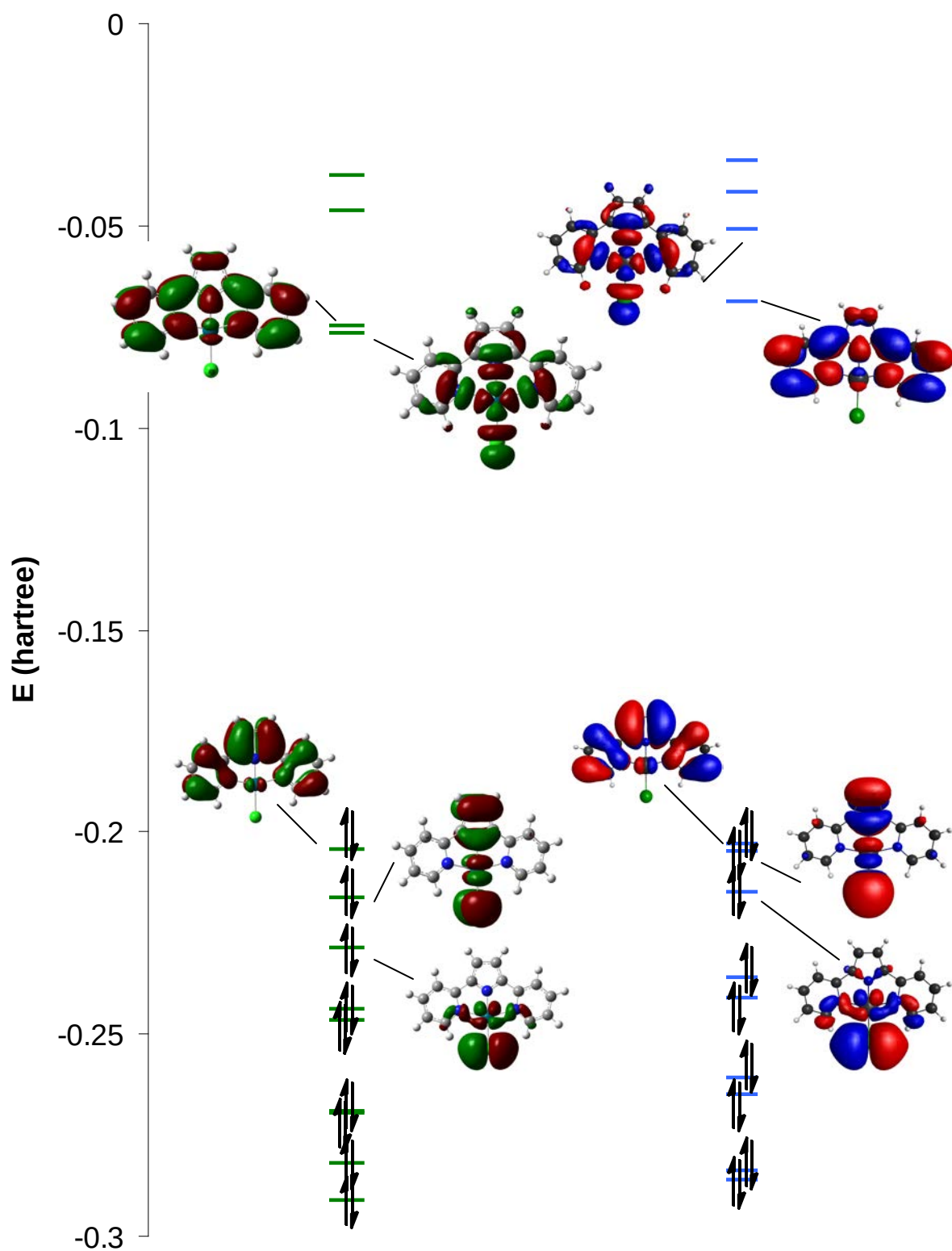


Figure S1. Frontier orbital surfaces and orbital energy diagram from DFT methods (B3LYP). Left (green and red): Mixed basis consisting of 6-31G* (CHN) and 3-21G (Pd, Cl). Right (blue and red): MIDI basis set.

Charge Decomposition Analysis-Output from QMForge⁶¹

Charge decomposition analysis of C:/Documents and Settings/MZdilla/My Documents/Gaussian/py-pyrole paper/CDA.log using the following fragments:

Frag1 (C:/Documents and Settings/MZdilla/My Documents/Gaussian/py-pyrole paper/fragm1.log)

Frag2 (C:/Documents and Settings/MZdilla/My Documents/Gaussian/py-pyrole paper/fragm2.log)

Created on Wed May 23 13:38:21 2012

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2:	0.000	-0.000	-0.000	-0.000
3:	0.000	-0.000	0.000	-0.000
4:	0.000	-0.000	-0.000	-0.000
5:	-0.000	-0.000	-0.000	-0.000
6:	-0.000	-0.000	-0.000	-0.000
7:	-0.001	-0.000	0.002	-0.000
8:	-0.001	-0.000	-0.000	-0.000
9:	-0.001	0.000	-0.000	-0.000
10:	0.000	-0.000	0.000	-0.000
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12:	-0.000	0.001	0.000	-0.000
13:	-0.000	0.003	0.000	-0.000
14:	0.001	-0.001	-0.000	-0.000
15:	0.000	-0.001	-0.000	-0.001
16:	0.000	-0.000	-0.000	-0.000
17:	-0.000	-0.000	-0.000	-0.000
18:	0.000	-0.003	-0.000	-0.001
19:	-0.000	0.001	-0.000	-0.000
20:	-0.000	0.000	-0.000	-0.000
21:	-0.000	0.001	-0.000	-0.000
22:	-0.000	0.000	-0.000	-0.000
23:	-0.000	0.000	0.000	-0.000
24:	-0.000	0.000	-0.000	-0.000
25:	-0.000	0.001	-0.000	-0.000
26:	-0.000	0.000	0.000	-0.000
27:	-0.000	0.000	0.000	-0.000
28:	-0.000	0.000	-0.000	-0.000
29:	-0.000	0.000	0.000	-0.000
30:	-0.000	0.000	-0.000	-0.000
31:	-0.000	0.000	-0.000	-0.000
32:	-0.000	0.000	0.000	-0.000
33:	-0.000	-0.001	-0.000	-0.001
34:	-0.000	-0.000	-0.000	-0.000
35:	0.000	-0.000	-0.000	-0.000
36:	-0.000	-0.000	-0.000	-0.000
37:	0.021	-0.001	0.016	-0.000
38:	0.047	-0.009	0.054	-0.002
39:	0.050	-0.002	0.025	-0.001
40:	0.000	-0.000	-0.000	0.000
41:	-0.002	-0.049	-0.012	-0.000
42:	-0.001	-0.022	-0.011	0.000

43:	0.000	-0.007	0.022	0.000
44:	-0.000	-0.001	0.005	-0.000
45:	-0.000	-0.006	-0.006	0.000
46:	0.000	0.001	0.003	0.000
47:	-0.000	-0.002	-0.000	-0.000
48:	-0.001	-0.015	0.005	0.000
49:	0.000	0.002	0.001	0.000
50:	-0.002	-0.006	0.001	-0.001
51:	-0.000	0.003	0.002	0.000
52:	0.000	0.010	0.007	0.001
53:	-0.000	0.004	0.002	0.000
54:	-0.001	-0.001	-0.005	0.000
55:	0.000	0.014	-0.004	0.001
56:	0.001	0.007	0.037	0.000
57:	-0.000	0.029	0.007	0.000
58:	-0.000	0.013	-0.003	-0.000
59:	0.001	-0.002	0.059	0.000
60:	0.001	0.012	0.010	0.001
61:	0.000	0.009	0.003	-0.000
62:	-0.001	0.002	0.009	0.000
63:	0.000	0.009	-0.002	-0.000
64:	0.005	0.067	0.044	0.000
65:	0.003	0.056	0.186	-0.000
66:	0.000	0.037	-0.012	-0.000
67:	0.005	0.000	0.048	0.000
68:	-0.001	0.040	-0.020	0.002
69:	-0.000	0.007	0.005	-0.000
70:	0.002	0.006	0.005	0.001
71:	0.005	0.003	0.035	-0.000
72:	0.017	0.030	0.201	-0.002
73:	0.001	0.014	-0.003	-0.001
74:	0.006	0.022	0.046	-0.004
75:	0.002	0.089	0.170	-0.005
76:	0.030	0.052	0.026	0.023
77:	0.027	0.000	0.095	0.000
78:	0.014	0.025	0.018	0.004
79:	0.003	-0.000	0.007	-0.000
80:	0.062	0.011	0.035	-0.004
81:	0.054	0.119	-0.194	-0.002
82:	-0.001	0.002	-0.001	-0.000
83:	0.004	0.154	-0.300	-0.008
84:	0.056	0.002	-0.165	-0.001
85:	-0.007	-0.082	-0.060	-0.105
86:	-0.004	0.020	-0.035	0.002
87:	0.009	-0.113	0.007	0.071
88:	0.012	0.012	-0.164	-0.002
89:	0.001	0.000	-0.022	-0.000
-----	HOMO	-	LUMO	gap
90:	0.000	0.000	0.000	0.000
91:	0.000	0.000	0.000	0.000
92:	0.000	0.000	0.000	0.000
.....				
307:	0.000	0.000	0.000	0.000
T:	0.418	0.568	0.175	-0.035

NMR Spectra

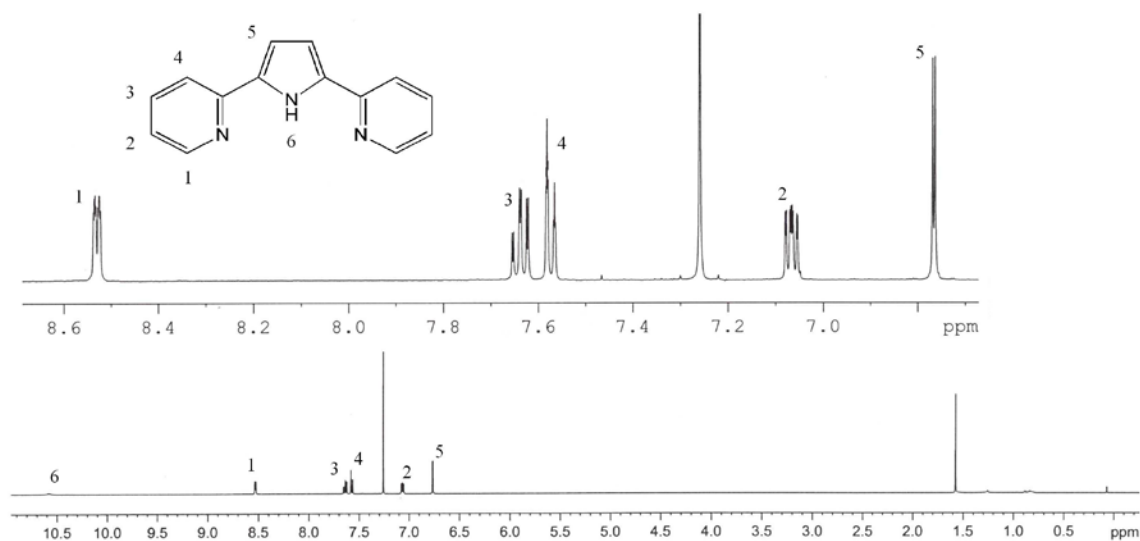


Figure S2. ^1H NMR spectrum of H(PDP) in CDCl_3

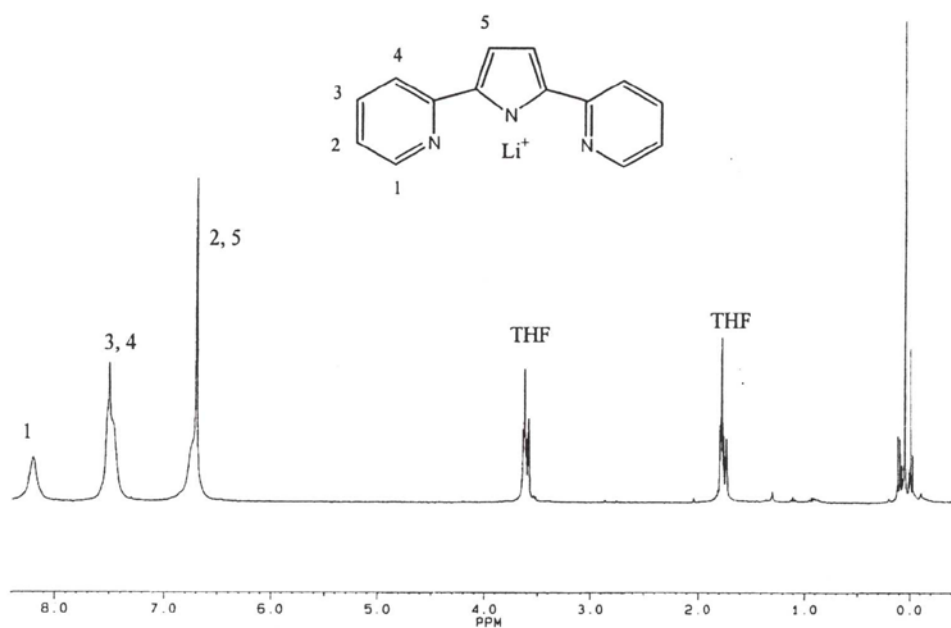


Figure S3. ^1H NMR spectrum of Li(PDP) in $\text{THF-}d_8$.

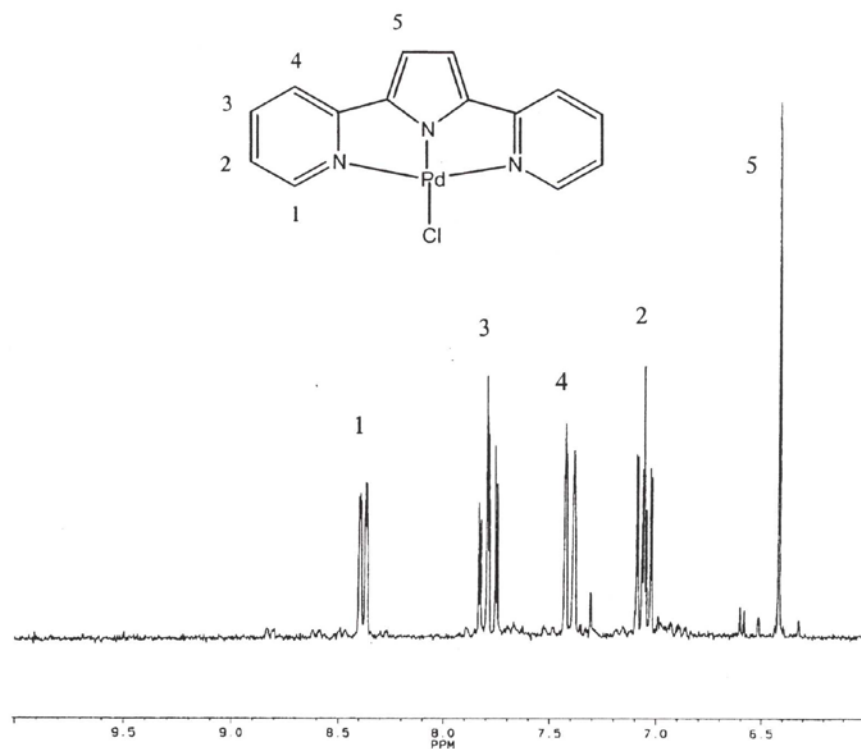


Figure S4. ^1H NMR spectrum of (PDP)Pd-Cl (**1**) in $\text{THF-}d_8$.

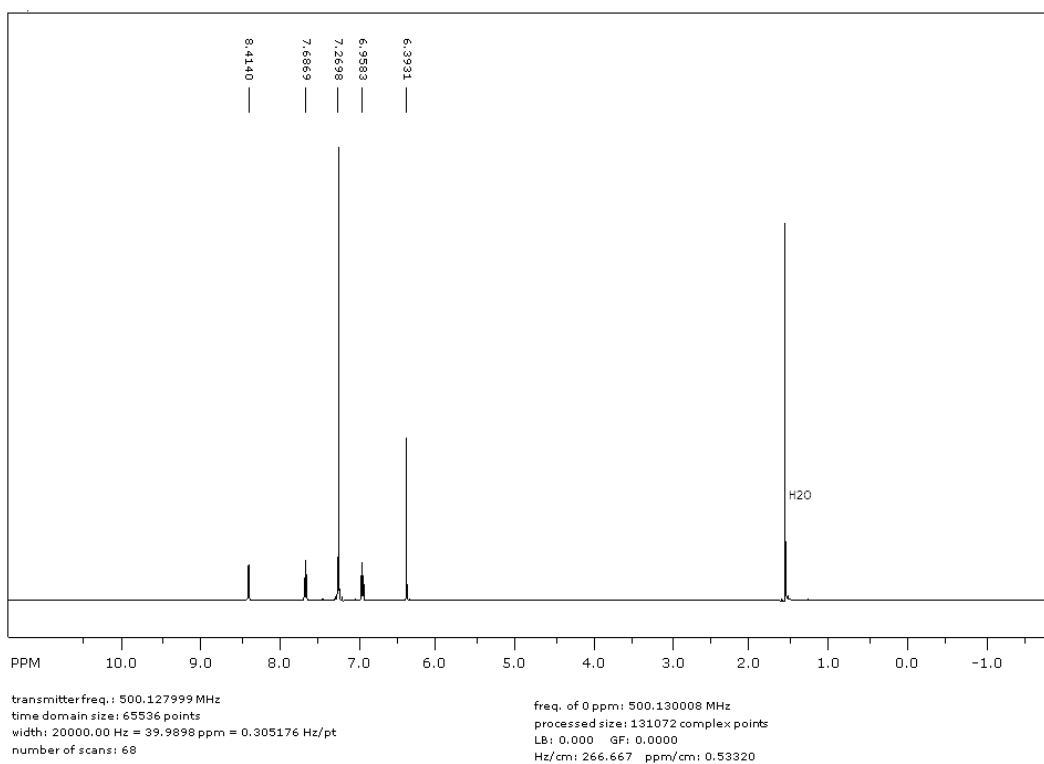


Figure S5. ^1H NMR spectrum of (PDP)Pd-Cl (**1**) in chloroform- d

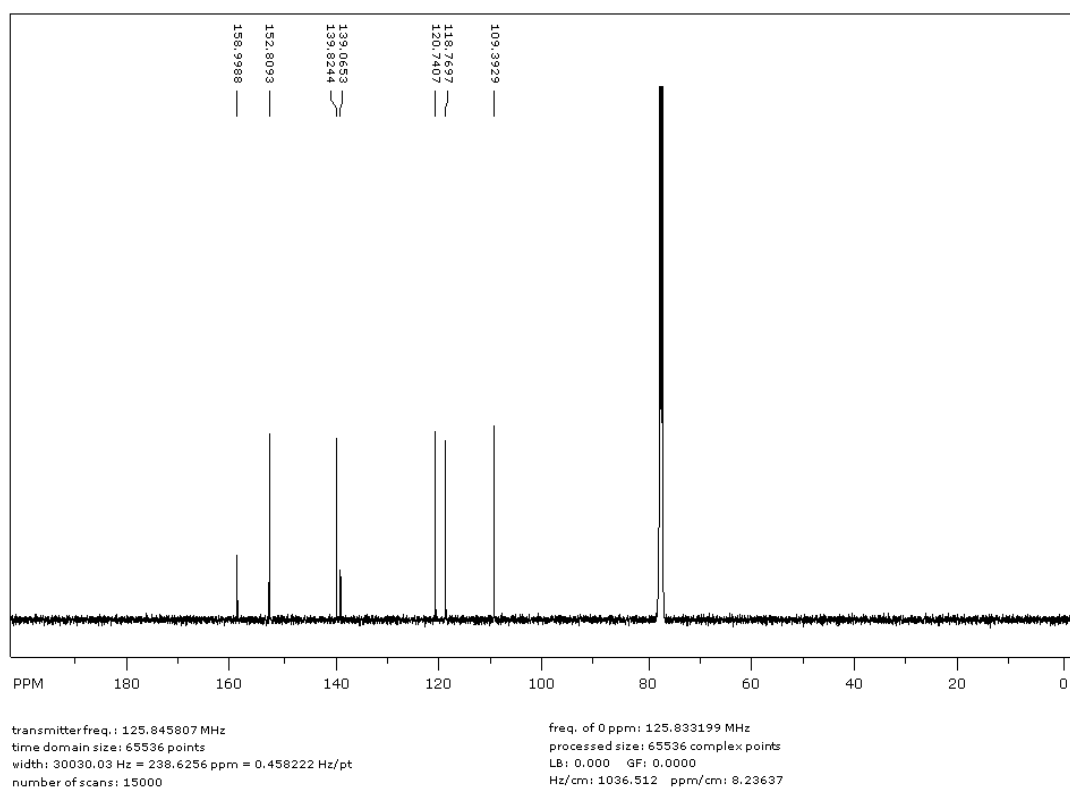


Figure S6. ^{13}C NMR spectrum of (PDP)Pd-Cl (**1**) in chloroform-*d*

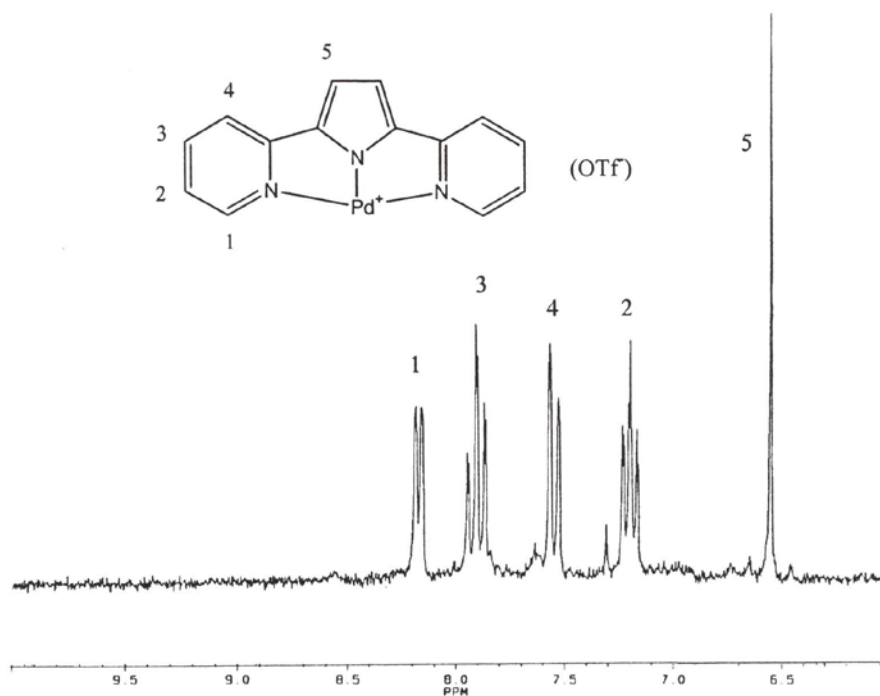


Figure S7. ^1H NMR spectrum of $(\text{PDP})\text{Pd}-\text{OH}_2]^+ [\text{OTf}]^-$ (**2**) in $\text{THF}-d_8$.

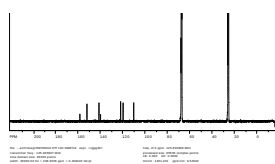


Figure S8. ^{13}C NMR spectrum of $(\text{PDP})\text{Pd}-\text{OH}_2]^+ [\text{OTf}]^-$ (**2**) in $\text{THF}-d_8$.

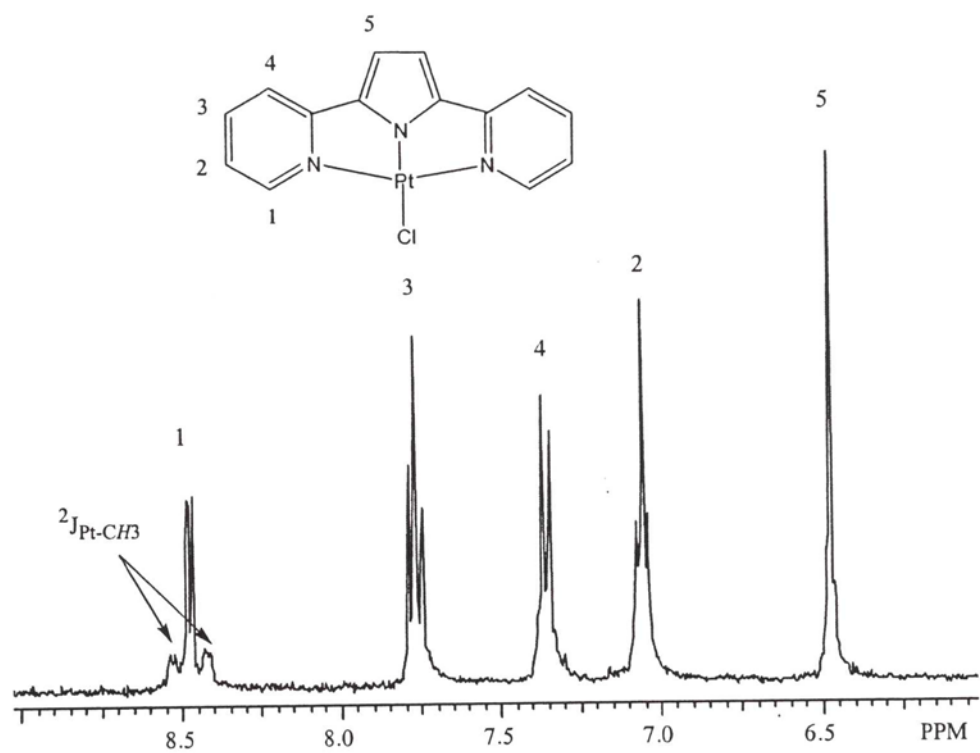


Figure S9. ^1H NMR spectrum of (PDP)Pt-Cl (**3**) in $\text{THF-}d_8$.

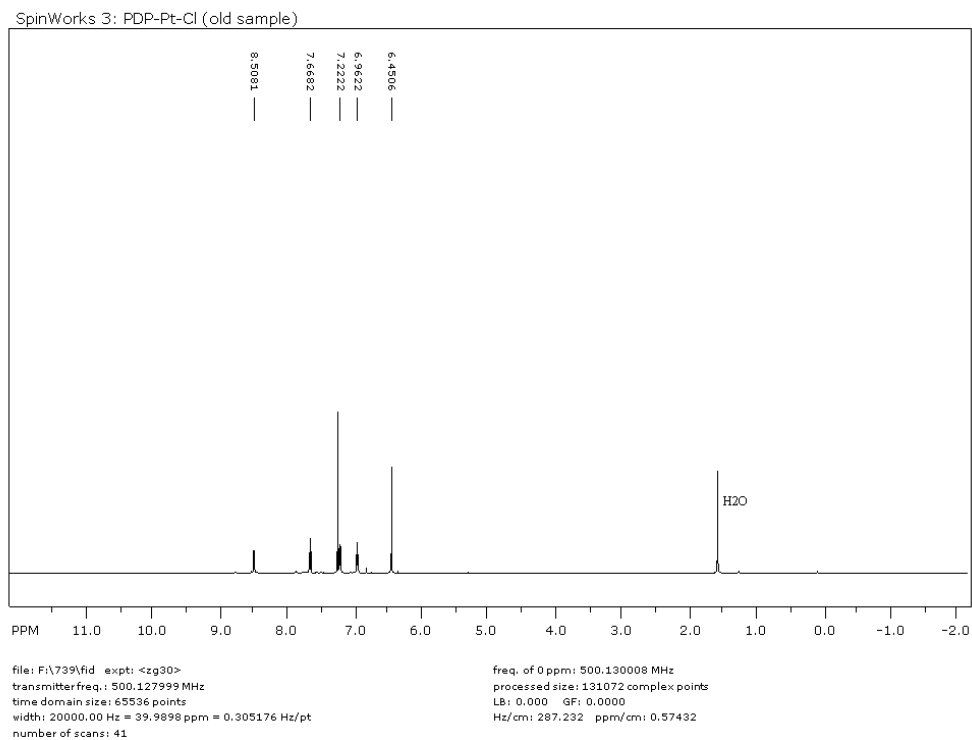


Figure S10. ^1H NMR spectrum of (PDP)Pt-Cl (**3**) in chloroform- d

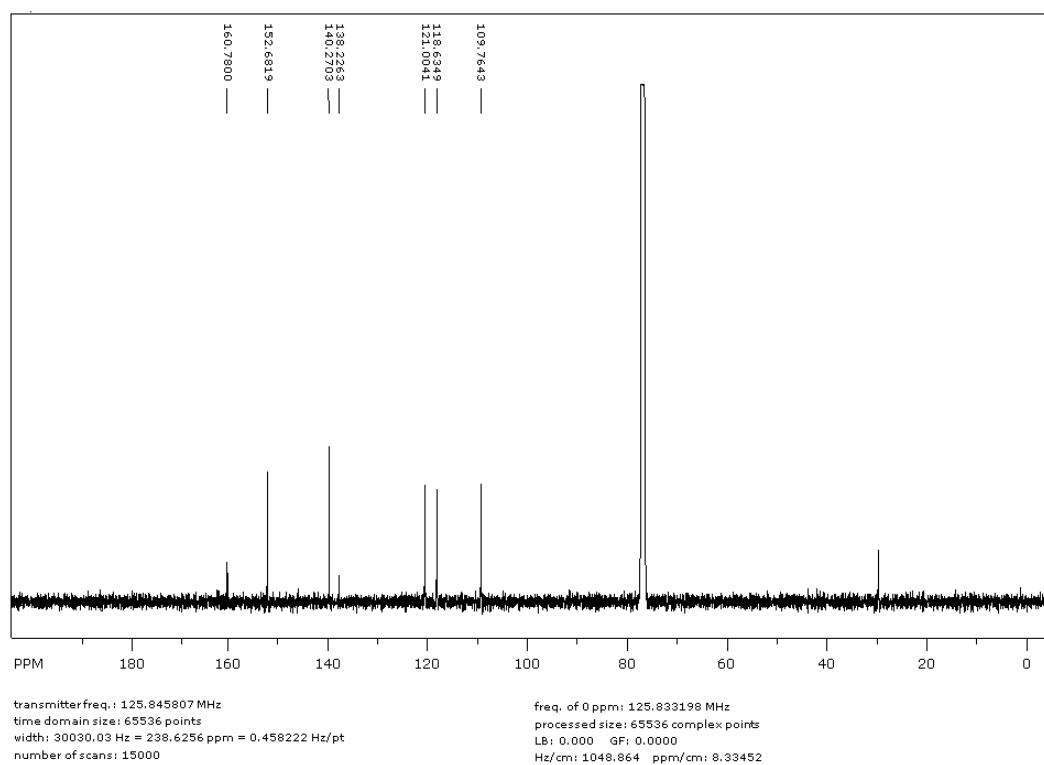


Figure S11. ^{13}C NMR spectrum of (PDP)Pt-Cl (**3**) in chloroform-*d*

UV-Visible Absorption Spectra

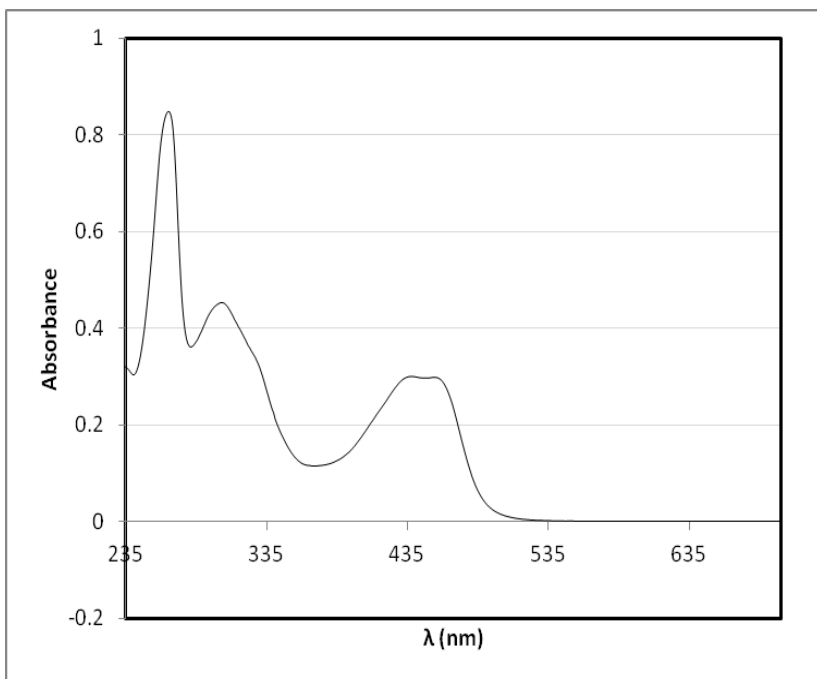


Figure S12. UV-Vis Absorption spectrum of (PDP)Pd-Cl (**1**) in dichloromethane. λ_{max} = 265 (ϵ 36849), 303 (19662), 437 (13027), 454 (12945)

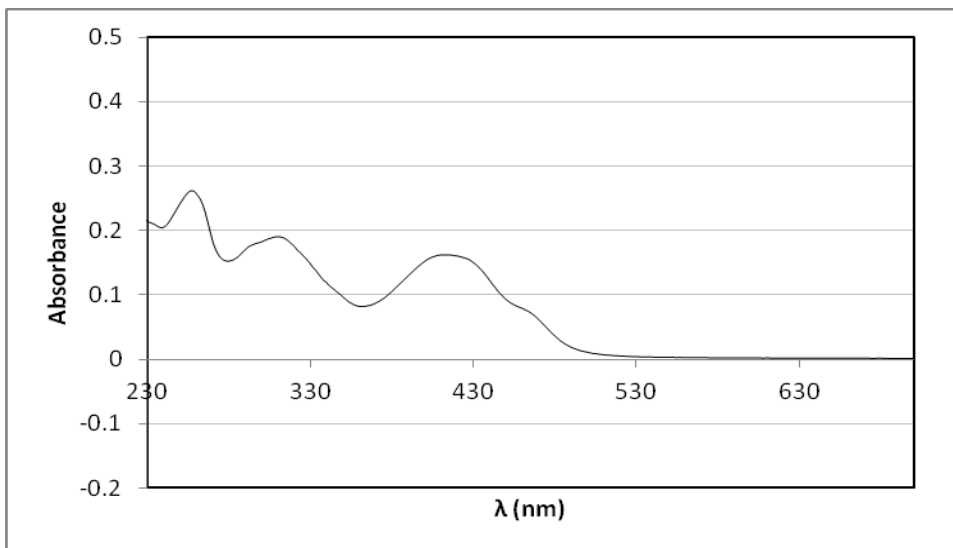


Figure S12. UV-Vis Absorption spectrum of (PDP)Pd-Cl (**1**) in ethanol. λ_{max} = 258 nm (7816), 311 nm (5672), 414 nm (4839).

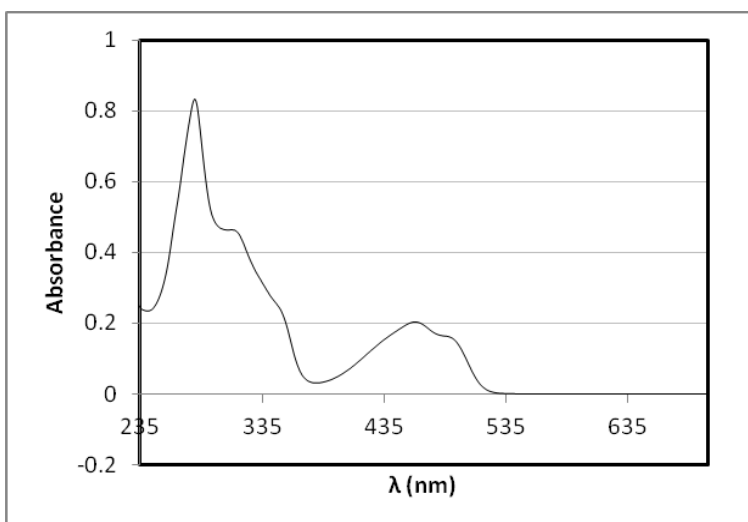


Figure S14. UV-Vis Absorption spectrum of (PDP)Pt-Cl (**3**) in dichloromethane. λ_{max} = 280 (ϵ 25287), 310 (14093), 459 (6171), 486 (4960)

Cyclic Voltammetry

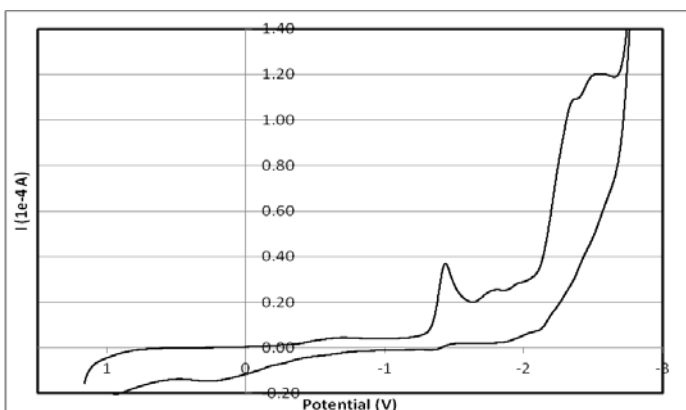


Figure S15. Cyclic voltammogram of (PDP)Pd-Cl (**1**) in acetonitrile. Scan rate = 0.1 V/s. Potentials are vs. NHE.

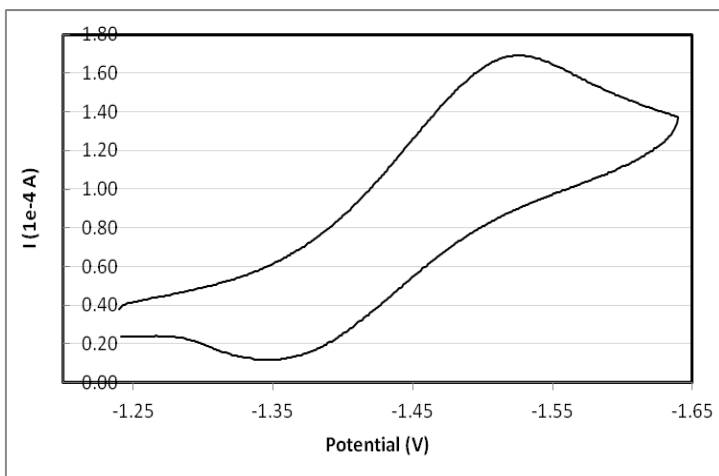


Figure S16. Rapid scan cyclic voltammogram of (PDP)Pd-Cl (**1**) in acetonitrile in the region of -1.45 V vs. NHE. Scan rate = 1 V/s

Infrared Spectra

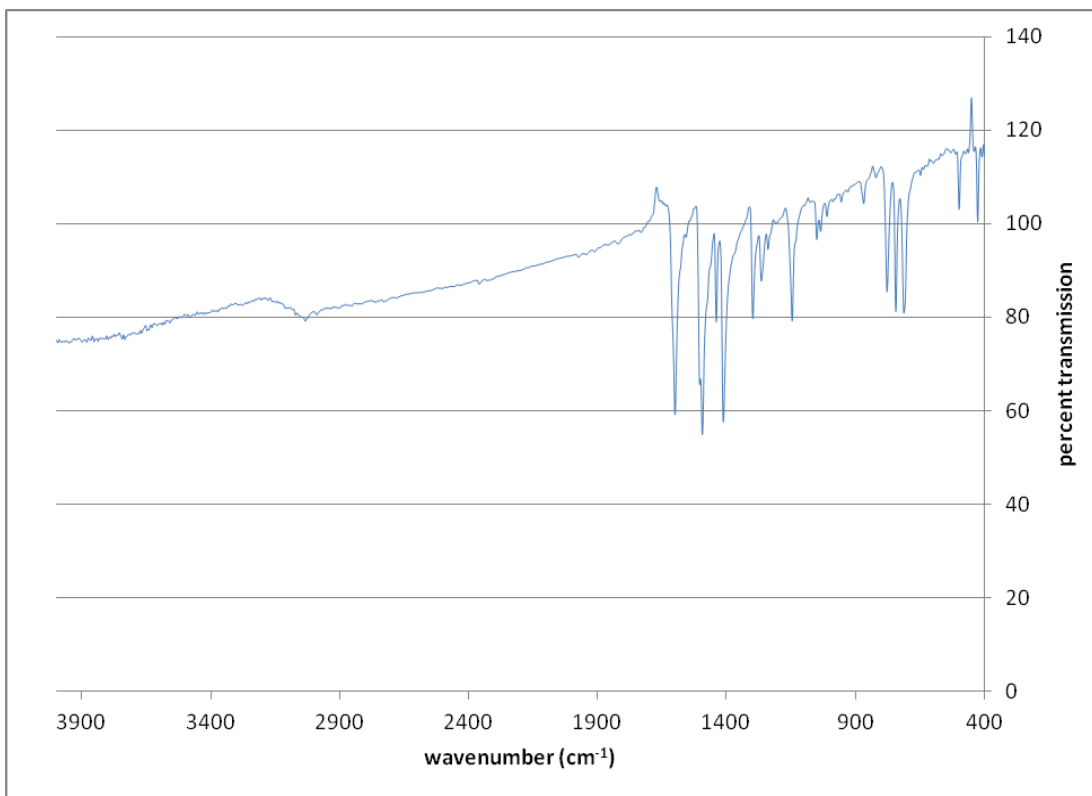


Figure S17. IR Spectrum of PDPPd-Cl (1) (KBr Pellet)

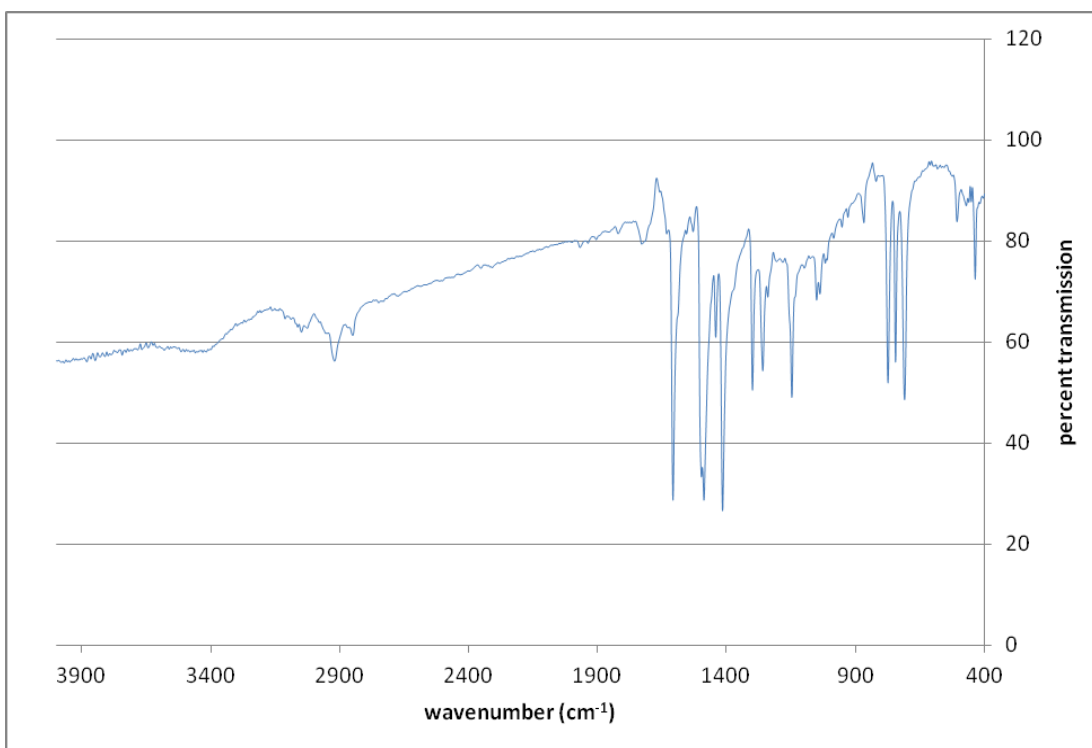


Figure S18. IR Spectrum of PDPPt-Cl (3) (KBr Pellet)

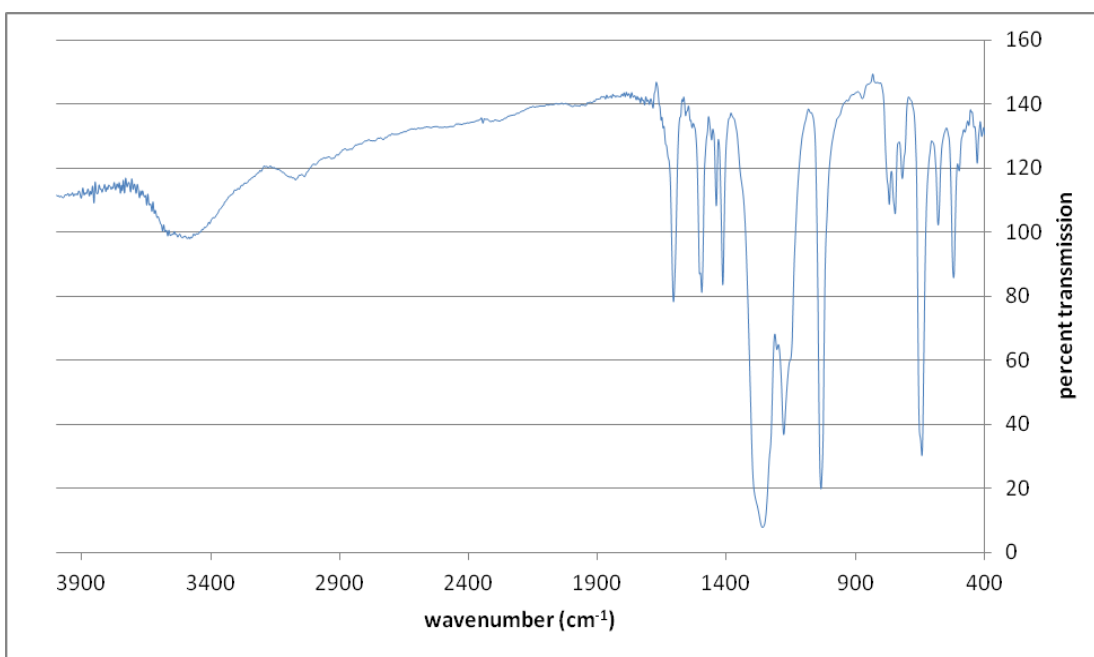


Figure S19. IR Spectrum of (PDP)Pd-OH₂]⁺ [OTf]⁻ (2) (KBr Pellet)

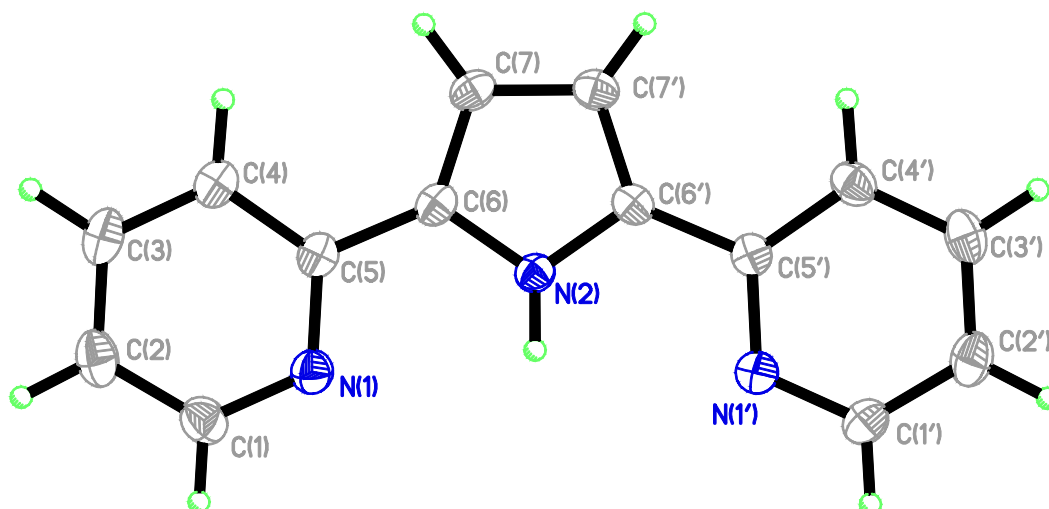


Figure X1. Structure of PDP(H) ligand with thermal ellipsoids (50%), and hydrogens shown as open circles.

Table X1.1. Sample and crystal data for (PDP)H.

Chemical formula	$C_{14}H_{11}N_3$	
Formula weight	221.26	
Temperature	100(2) K	
Wavelength	0.71073 Å	
Crystal size	0.200 x 0.200 x 0.330 mm	
Crystal system	orthorhombic	
Space group	$P 2_1 2_1 2_1$	
Unit cell dimensions	$a = 5.9045(7)$ Å	$\alpha = 90^\circ$
	$b = 12.7669(16)$ Å	$\beta = 90^\circ$
	$c = 15.3114(18)$ Å	$\gamma = 90^\circ$
Volume	$1154.2(2)$ Å ³	
Z	4	
Density (calculated)	1.273 Mg/cm ³	
Absorption coefficient	0.079 mm ⁻¹	
F(000)	464	

Table X1.2. Data collection and structure refinement for (PDP)H.

Theta range for data collection	2.08 to 27.91°
Index ranges	-7<=h<=7, -16<=k<=14, -19<=l<=20
Reflections collected	6749
Independent reflections	2769 [R(int) = 0.0193]
Max. and min. transmission	0.9845 and 0.9745
Structure solution technique	direct methods
Structure solution program	SHELXS-97 (Sheldrick, 2008)
Refinement method	Full-matrix least-squares on F ²
Refinement program	SHELXL-97 (Sheldrick, 2008)
Function minimized	$\Sigma w(F_o^2 - F_c^2)^2$
Data / restraints / parameters	2769 / 0 / 154
Goodness-of-fit on F²	1.027 ^a
Final R indices^b	2510 data; I>2σ(I) R1 = 0.0344, wR2 = 0.0775 all data R1 = 0.0405, wR2 = 0.0807
Weighting scheme	w=1/[σ ² (F _o ²)+(0.0382P) ² +0.1960P] where P=(F _o ² +2F _c ²)/3
Absolute structure parameter	0.0(20)
Largest diff. peak and hole	0.154 and -0.181 eÅ ⁻³
R.M.S. deviation from mean	0.034 eÅ ⁻³

^aS = $[\Sigma w(F_{obs}^2 - F_{calc}^2)^2 / (m - n)]^{1/2}$ ^bR₁ = $(\Sigma ||F_{obs}| - |F_{calc}||) / \Sigma |F_{obs}|$; wR₂ = $[(\Sigma w(F_{obs}^2 - F_{calc}^2)^2) / \Sigma wF_{obs}^2]^{1/2}$

Table X1.3. Atomic coordinates and equivalent isotropic atomic displacement parameters (Å²) for (PDP)H.

U(eq) is defined as one third of the trace of the orthogonalized U_{ij} tensor.

	x/a	y/b	z/c	U(eq)
C1	0.4255(3)	0.88864(11)	0.51074(10)	0.0305(3)
C1'	0.4266(2)	0.37018(10)	0.22845(8)	0.0257(3)
C2	0.3277(3)	0.83685(11)	0.58101(10)	0.0327(3)
C2'	0.3236(3)	0.46404(11)	0.20706(9)	0.0298(3)
C3	0.1219(3)	0.87284(11)	0.61196(9)	0.0322(3)
C3'	0.1134(2)	0.48559(11)	0.24287(9)	0.0301(3)
C4	0.0214(2)	0.95803(11)	0.57164(8)	0.0269(3)
C4'	0.0142(3)	0.41296(10)	0.29780(9)	0.0255(3)
C5	0.1308(2)	0.00520(10)	0.50078(8)	0.0215(3)
C5'	0.1283(2)	0.31905(10)	0.31475(7)	0.0199(3)

	x/a	y/b	z/c	U(eq)
C6	0.0333(2)	0.09630(10)	0.45723(8)	0.0211(3)
C6'	0.0312(2)	0.23905(9)	0.37173(8)	0.0201(3)
C7'	0.8179(2)	0.23005(10)	0.40863(8)	0.0235(3)
C7	0.8205(2)	0.14041(10)	0.46260(8)	0.0232(3)
N1	0.3324(2)	0.97165(8)	0.47077(7)	0.0258(2)
N1'	0.3346(2)	0.29764(8)	0.28130(6)	0.0227(2)
N2	0.15837(19)	0.15673(8)	0.40114(6)	0.0199(2)

Table X1.4. Bond lengths (Å) for (PDP)H.

C1-N1	1.3417(18)	C1-C2	1.389(2)
C1-H1A	0.95	C1'-N1'	1.3447(16)
C1'-C2'	1.3832(19)	C1'-H1'A	0.95
C2-C3	1.383(2)	C2-H2A	0.95
C2'-C3'	1.385(2)	C2'-H2'A	0.95
C3-C4	1.384(2)	C3-H3A	0.95
C3'-C4'	1.3819(19)	C3'-H3'A	0.95
C4-C5	1.3990(18)	C4-H4A	0.95
C4'-C5'	1.3996(18)	C4'-H4'A	0.95
C5-N1	1.3460(17)	C5-C6	1.4590(18)
C5'-N1'	1.3493(17)	C5'-C6'	1.4605(17)
C6-N2	1.3704(16)	C6-C7	1.3792(19)
C6'-N2	1.3677(16)	C6'-C7'	1.3854(18)
C7-C7	1.4116(18)	C7'-H7'A	0.95
C7-H7A	0.95	N2-H2B	0.88

Table X1.5. Bond angles (°) for (PDP)H.

N1-C1-C2	123.99(14)	N1-C1-H1A	118.0
C2-C1-H1A	118.0	N1'-C1'-C2'	124.15(13)
N1'-C1'-H1'A	117.9	C2'-C1'-H1'A	117.9
C3-C2-C1	118.22(13)	C3-C2-H2A	120.9
C1-C2-H2A	120.9	C1'-C2'-C3'	118.21(13)
C1'-C2'-H2'A	120.9	C3'-C2'-H2'A	120.9
C2-C3-C4	119.02(13)	C2-C3-H3A	120.5
C4-C3-H3A	120.5	C4'-C3'-C2'	119.18(13)
C4'-C3'-H3'A	120.4	C2'-C3'-H3'A	120.4

Table X1.5 (cont)

C3-C4-C5	119.08(13)	C3-C4-H4A	120.5
C5-C4-H4A	120.5	C3'-C4'-C5'	118.92(13)
C3'-C4'-H4'A	120.5	C5'-C4'-H4'A	120.5
N1-C5-C4	122.43(12)	N1-C5-C6	116.52(11)
C4-C5-C6	121.02(12)	N1'-C5'-C4'	122.53(12)
N1'-C5'-C6'	116.06(11)	C4'-C5'-C6'	121.40(12)
N2-C6-C7	107.36(11)	N2-C6-C5	121.51(12)
C7-C6-C5	131.12(12)	N2-C6'-C7'	107.52(11)
N2-C6'-C5'	121.24(11)	C7'-C6'-C5'	131.19(12)
C6'-C7'-C7	107.21(12)	C6'-C7'-H7'A	126.4
C7-C7'-H7'A	126.4	C6-C7-C7'	107.83(12)
C6-C7-H7A	126.1	C7'-C7-H7A	126.1
C1-N1-C5	117.26(12)	C1'-N1'-C5'	116.98(11)
C6'-N2-C6	110.07(11)	C6'-N2-H2B	125.0
C6-N2-H2B	125.0		

Table X1.6. Anisotropic atomic displacement parameters ($\text{\AA}^2$) for (PDP)H.

The anisotropic atomic displacement factor exponent takes the form: $-2\pi^2 [h^2 a^{*2} U_{11} + \dots + 2 h k a^* b^* U_{12}]$

	U_{11}	U_{22}	U_{33}	U_{23}	U_{13}	U_{12}
C1	0.0309(8)	0.0255(7)	0.0350(7)	-0.0001(6)	-0.0011(6)	0.0033(6)
C1'	0.0269(6)	0.0296(7)	0.0207(6)	-0.0008(5)	0.0009(5)	-0.0033(6)
C2	0.0403(9)	0.0230(6)	0.0347(7)	0.0052(6)	-0.0065(7)	-0.0014(7)
C2'	0.0367(8)	0.0309(7)	0.0218(6)	0.0067(5)	-0.0031(6)	-0.0046(7)
C3	0.0410(9)	0.0290(7)	0.0265(6)	0.0065(6)	0.0003(6)	-0.0073(7)
C3'	0.0377(8)	0.0259(7)	0.0266(6)	0.0054(5)	-0.0067(6)	0.0044(6)
C4	0.0294(8)	0.0268(6)	0.0244(6)	0.0008(5)	0.0022(6)	-0.0034(6)
C4'	0.0262(7)	0.0271(6)	0.0231(6)	-0.0003(5)	-0.0025(6)	0.0044(6)
C5	0.0252(7)	0.0195(6)	0.0198(5)	-0.0029(5)	-0.0015(5)	-0.0047(5)
C5'	0.0231(7)	0.0216(6)	0.0150(5)	-0.0022(4)	-0.0034(5)	-0.0003(5)
C6	0.0251(7)	0.0215(6)	0.0167(5)	-0.0034(5)	0.0014(5)	-0.0038(5)
C6'	0.0225(7)	0.0195(6)	0.0184(5)	-0.0045(4)	-0.0024(5)	0.0011(5)
C7'	0.0234(7)	0.0253(6)	0.0219(6)	-0.0042(5)	-0.0006(6)	0.0019(6)
C7	0.0238(7)	0.0260(6)	0.0197(6)	-0.0044(5)	0.0027(6)	-0.0035(6)
N1	0.0266(6)	0.0230(5)	0.0277(5)	0.0011(5)	0.0014(5)	-0.0007(5)
N1'	0.0236(5)	0.0243(5)	0.0201(5)	-0.0014(4)	-0.0012(5)	0.0000(5)
N2	0.0210(5)	0.0202(5)	0.0184(5)	-0.0015(4)	0.0012(5)	-0.0001(5)

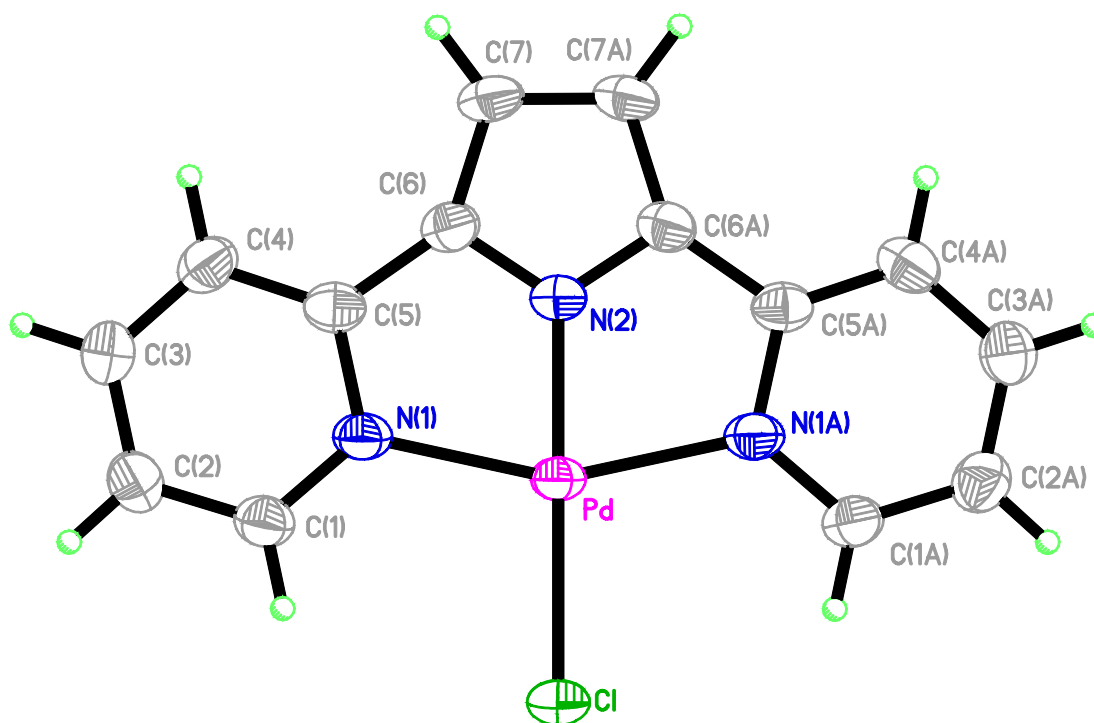


Figure X2. Structure of (PDP)PdCl (1) with thermal ellipsoids (50%), and hydrogens shown as open circles.

Table X2.1. Sample and crystal data for PdCl (1).

Chemical formula	C ₁₄ H ₁₀ ClN ₃ Pd	
Formula weight	362.10	
Temperature	200(2) K	
Wavelength	0.71069 Å	
Crystal size	0.020 x 0.040 x 0.380 mm	
Crystal system	monoclinic	
Space group	C2/c	
Unit cell dimensions	a = 10.6112(9) Å	α = 90°
	b = 13.4902(15) Å	β = 99.556(6)°
	c = 8.8782(9) Å	γ = 90°
Volume	1253.3(2) Å ³	
Z	4	
Density (calculated)	1.919 Mg/cm ³	
Absorption coefficient	1.680 mm ⁻¹	
F(000)	712	

Table X2.2. Data collection and structure refinement for (PDP)PdCl (1).

Theta range for data collection	2.46 to 25.34°
Index ranges	-12<=h<=12, -16<=k<=16, -10<=l<=9
Reflections collected	4789
Independent reflections	1143 [R(int) = 0.0467]
Structure solution technique	direct methods
Structure solution program	SIR92 (Altomare, 1993)
Refinement method	Full-matrix least-squares on F ²
Refinement program	SHELXL-93 (Sheldrick, 1993)
Function minimized	$\Sigma w(F_o^2 - F_c^2)^2$
Data / restraints / parameters	1143 / 0 / 89
Goodness-of-fit on F²	1.106 ^a
Final R indices^b	1143 data; I>2σ(I) R1 = 0.0459, wR2 = 0.1036 all data R1 = 0.0490, wR2 = 0.1060
Weighting scheme	w=1/[σ ² (F _o ²)+(0.0496P) ² +9.3146P] where P=(F _o ² +2F _c ²)/3
Extinction coefficient	0.0022(5)
Largest diff. peak and hole	0.715 and -1.438 eÅ ⁻³
R.M.S. deviation from mean	0.140 eÅ ⁻³

$$^a S = [\Sigma w(F_{obs}^2 - F_{calc}^2)^2 / (m - n)]^{1/2} \quad ^b R_1 = (\Sigma ||F_{obs}| - |F_{calc}||) / \Sigma |F_{obs}|; \quad wR_2 = [(\Sigma w(F_{obs}^2 - F_{calc}^2)^2) / \Sigma wF_{obs}^2]^{1/2}$$

Table X2.3. Atomic coordinates and equivalent isotropic atomic displacement parameters (Å²) for PdCl (1).

U(eq) is defined as one third of the trace of the orthogonalized U_{ij} tensor.

	x/a	y/b	z/c	U(eq)
Pd	0.5	0.07807(4)	0.75	0.0304(3)
Cl	0.5	0.25081(13)	0.75	0.0430(5)
N1	0.6445(5)	0.0453(3)	0.6268(6)	0.0346(11)
N2	0.5	0.9383(4)	0.75	0.0329(14)
C1	0.7171(6)	0.1110(4)	0.5645(7)	0.0375(13)
C2	0.8132(6)	0.0804(4)	0.4857(8)	0.0421(14)
C3	0.8343(6)	0.9809(5)	0.4699(8)	0.0426(14)
C4	0.7609(6)	0.9128(4)	0.5329(7)	0.0400(14)
C5	0.6650(6)	0.9455(4)	0.6089(6)	0.0336(12)
C6	0.5794(6)	0.8834(4)	0.6815(7)	0.0347(12)
C7	0.5492(6)	0.7842(4)	0.7064(7)	0.0413(14)

Table X2.4. Bond lengths (Å) for PdCl (1).

Pd-N2	1.885(6)	Pd-N1#1	2.073(5)
Pd-N1	2.073(5)	Pd-Cl	2.330(2)
N1-C1	1.351(7)	N1-C5	1.377(7)
N2-C6#1	1.341(6)	N2-C6	1.341(6)
C1-C2	1.391(9)	C2-C3	1.372(8)
C3-C4	1.382(9)	C4-C5	1.384(8)
C5-C6	1.461(8)	C6-C7	1.402(8)
C7-C7#1	1.400(13)		

Symmetry transformations used to generate equivalent atoms:

#1 -x+1, y, -z+3/2

Table X2.5. Bond angles (°) for PdCl (1).

N2-Pd-N1#1	77.71(12)	N2-Pd-N1	77.70(12)
N1#1-Pd-N1	155.4(2)	N2-Pd-Cl	180.0
N1#1-Pd-Cl	102.29(12)	N1-Pd-Cl	102.30(12)
C1-N1-C5	118.8(5)	C1-N1-Pd	126.8(4)
C5-N1-Pd	114.4(4)	C6#1-N2-C6	113.0(6)
C6#1-N2-Pd	123.5(3)	C6-N2-Pd	123.5(3)
N1-C1-C2	121.8(5)	C3-C2-C1	119.2(5)
C2-C3-C4	119.8(5)	C3-C4-C5	119.7(5)
N1-C5-C4	120.8(5)	N1-C5-C6	112.8(5)
C4-C5-C6	126.4(5)	N2-C6-C7	106.2(5)
N2-C6-C5	111.5(5)	C7-C6-C5	142.3(5)
C7#1-C7-C6	107.3(3)		

Symmetry transformations used to generate equivalent atoms:

#1 -x+1, y, -z+3/2

Table X2.6. Anisotropic atomic displacement parameters ($\text{\AA}^2$) for PdCl (1).

The anisotropic atomic displacement factor exponent takes the form: $-2\pi^2 [h^2 a^{*2} U_{11} + \dots + 2 h k a^* b^* U_{12}]$						
	U_{11}	U_{22}	U_{33}	U_{23}	U_{13}	U_{12}
Pd	0.0331(4)	0.0208(3)	0.0376(4)	0	0.0066(3)	0
Cl	0.0530(13)	0.0220(8)	0.0571(13)	0	0.0181(10)	0
N1	0.037(3)	0.024(2)	0.043(3)	0.002(2)	0.006(2)	0.001(2)
N2	0.038(4)	0.023(3)	0.038(4)	0	0.009(3)	0
C1	0.039(3)	0.029(3)	0.043(3)	0.003(2)	0.006(3)	-0.001(2)
C2	0.037(3)	0.042(3)	0.049(4)	0.003(3)	0.011(3)	-0.004(2)
C3	0.034(3)	0.045(3)	0.051(4)	0.000(3)	0.013(3)	0.002(3)
C4	0.037(3)	0.035(3)	0.047(3)	-0.008(3)	0.006(3)	0.004(2)
C5	0.038(3)	0.028(3)	0.032(3)	-0.002(2)	0.000(2)	-0.002(2)
C6	0.037(3)	0.028(3)	0.039(3)	-0.001(2)	0.006(2)	0.003(2)
C7	0.052(4)	0.023(2)	0.050(4)	-0.003(2)	0.011(3)	0.004(2)

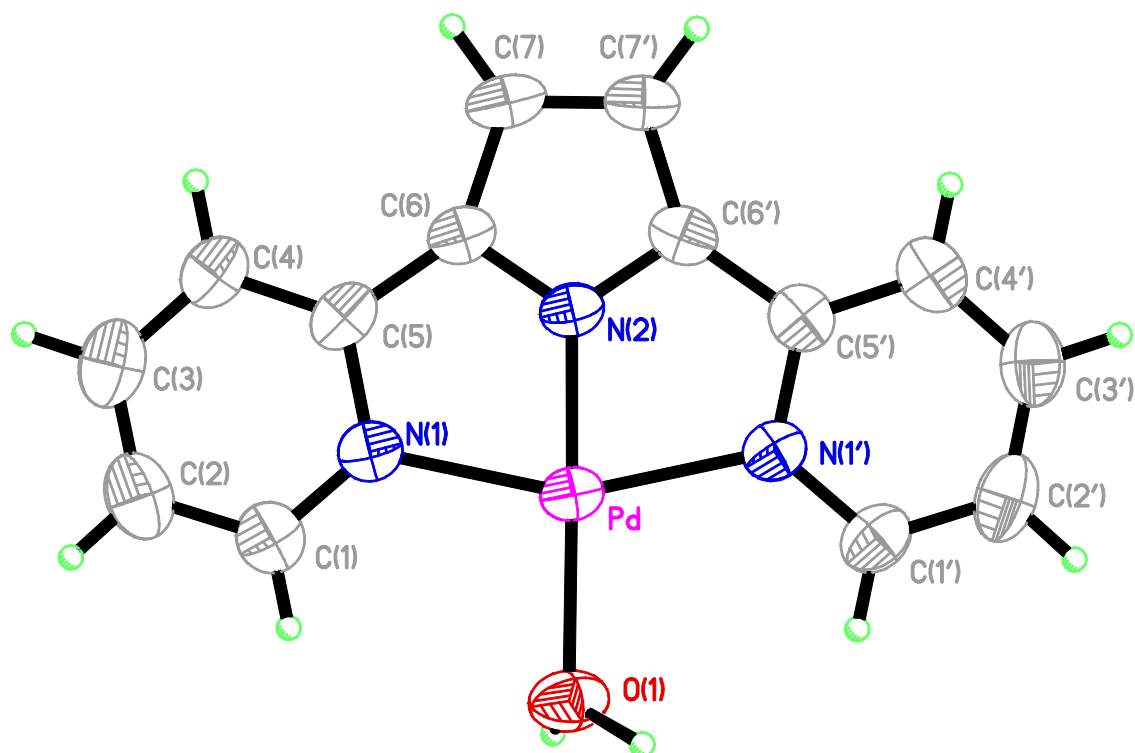


Figure X3. Structure of (PDP)Pd(OH)₂⁺[OTf]⁻·H₂O (**2**) with thermal ellipsoids (30%), and hydrogens shown as open circles. Triflate counterion and solvent water not shown.

Table X3.1. Sample and crystal data for (PDP)Pd(OH)₂⁺[OTf]⁻·H₂O (2**).**

Chemical formula	C ₁₅ H ₁₄ F ₃ N ₃ O ₅ PdS	
Formula weight	511.75	
Temperature	295(1) K	
Wavelength	0.71069 Å	
Crystal size	0.07 x 0.10 x 0.27 mm	
Crystal system	triclinic	
Space group	P -1	
Unit cell dimensions	a = 11.8162(5) Å	α = 94.8250(10)°
	b = 12.8499(7) Å	β = 106.811(4)°
	c = 6.9992(4) Å	γ = 110.907(3)°
Volume	929.26(8) Å ³	
Z	2	
Density (calculated)	1.829 Mg/cm ³	
Absorption coefficient	1.173 mm ⁻¹	
F(000)	508	

Table X3.2. Data collection and structure refinement for (PDP)Pd(OH)₂⁺[OTf]⁻·H₂O (2).

Theta range for data collection	3.06 to 25.35°		
Reflections collected	7770		
Independent reflections	3128 [R(int) = 0.0264]		
Max. and min. transmission	1.0000 and 0.6832		
Structure solution technique	direct methods		
Structure solution program	SIR92 (Altomare, 1993)		
Refinement method	Full-matrix least-squares on F ²		
Refinement program	SHELXL-93 (Sheldrick, 1993)		
Function minimized	$\Sigma w(F_o^2 - F_c^2)^2$		
Data / restraints / parameters	3128 / 0 / 254		
Goodness-of-fit on F²	1.111 ^a		
Final R indices^b	2830 data; I>2σ(I)	R1 = 0.0496, wR2 = 0.1175	
	all data	R1 = 0.0560, wR2 = 0.1221	
Weighting scheme	$w=1/[\sigma^2(F_o^2)+(0.0548P)^2+1.5735P]$		
	where $P=(F_o^2+2F_c^2)/3$		
Largest diff. peak and hole	0.485 and -0.860 eÅ ⁻³		
R.M.S. deviation from mean	0.082 eÅ ⁻³		

$$^a S = [\Sigma w(F_{obs}^2 - F_{calc}^2)^2 / (m - n)]^{1/2} \quad ^b R_1 = (\Sigma ||F_{obs}| - |F_{calc}||) / \Sigma |F_{obs}|; \quad wR_2 = [(\Sigma w(F_{obs}^2 - F_{calc}^2)^2) / \Sigma wF_{obs}^2]^{1/2}$$

Table X3.3. Atomic coordinates and equivalent isotropic atomic displacement parameters (Å²) for (PDP)Pd(OH)₂⁺[OTf]⁻·H₂O (2).

U(eq) is defined as one third of the trace of the orthogonalized U_{ij} tensor.

	x/a	y/b	z/c	U(eq)
Pd1	0.17016(4)	0.56567(4)	0.34130(6)	0.05674(18)
N1	0.0902(5)	0.3921(4)	0.3340(7)	0.0621(11)
N1'	0.1688(5)	0.7242(4)	0.3021(7)	0.0619(11)
N2	0.9952(4)	0.5325(4)	0.2397(7)	0.0592(11)
C1	0.1521(8)	0.3257(5)	0.3915(10)	0.0788(18)
C1'	0.2719(7)	0.8233(5)	0.3440(10)	0.0787(18)
C2	0.0873(10)	0.2102(7)	0.3792(11)	0.096(2)
C2'	0.2580(10)	0.9227(5)	0.3062(11)	0.097(3)
C3	0.9559(10)	0.1645(7)	0.3101(11)	0.093(2)
C3'	0.1368(10)	0.9205(7)	0.2254(11)	0.092(2)

	x/a	y/b	z/c	U(eq)
C4	0.8890(7)	0.2322(5)	0.2507(9)	0.0763(18)
C4'	0.0317(8)	0.8212(5)	0.1841(10)	0.080(2)
C5	0.9564(5)	0.3463(5)	0.2634(8)	0.0655(15)
C5'	0.0474(7)	0.7220(5)	0.2225(8)	0.0669(15)
C6	0.9026(5)	0.4284(5)	0.2086(8)	0.0626(14)
C6'	0.9461(5)	0.6101(5)	0.1873(8)	0.0644(15)
C7	0.7839(5)	0.4405(5)	0.1295(9)	0.0750(18)
C7'	0.8115(5)	0.5520(5)	0.1161(9)	0.0735(17)
O1	0.3666(5)	0.6004(4)	0.4640(7)	0.0863(13)
O2	0.5142(5)	0.5602(4)	0.2721(7)	0.0827(13)
S1	0.61630(17)	0.78765(13)	0.9263(2)	0.0728(4)
F1	0.6290(5)	0.9768(4)	0.1086(11)	0.145(2)
F2	0.6573(14)	0.8604(8)	0.2973(11)	0.254(6)
F3	0.4726(9)	0.8327(5)	0.1009(17)	0.205(4)
C8	0.5884(11)	0.8687(8)	0.1125(18)	0.114(3)
O3	0.5709(5)	0.6752(4)	0.9568(9)	0.1011(17)
O4	0.5330(7)	0.8006(5)	0.7420(9)	0.136(3)
O5	0.7475(5)	0.8439(5)	0.9603(13)	0.142(3)

Table X3.4. Bond lengths (Å) for (PDP)Pd(OH)₂]⁺[OTf]⁻·H₂O (2).

Pd1-N2	1.853(5)	Pd1-N1	2.078(5)
Pd1-N1'	2.084(5)	Pd1-O1	2.090(5)
N1-C1	1.326(9)	N1-C5	1.387(8)
N1'-C1'	1.346(8)	N1'-C5'	1.369(8)
N2-C6	1.342(7)	N2-C6'	1.348(8)
C1-C2	1.389(11)	C1'-C2'	1.381(11)
C2-C3	1.363(12)	C2'-C3'	1.368(12)
C3-C4	1.384(11)	C3'-C4'	1.361(11)
C4-C5	1.377(9)	C4'-C5'	1.389(9)
C5-C6	1.440(9)	C5'-C6'	1.450(9)
C6-C7	1.421(9)	C6'-C7'	1.403(9)
C7-C7'	1.370(10)	S1-O5	1.393(6)
S1-O3	1.409(5)	S1-O4	1.442(6)
S1-C8	1.777(10)	F1-C8	1.303(10)
F2-C8	1.350(13)	F3-C8	1.253(12)

Table X3.5. Bond angles (°) for (PDP)Pd(OH)₂]⁺[OTf]⁻·H₂O (2).

N2-Pd1-N1	78.1(2)	N2-Pd1-N1'	78.0(2)
N1-Pd1-N1'	156.1(2)	N2-Pd1-O1	178.1(2)
N1-Pd1-O1	100.6(2)	N1'-Pd1-O1	103.3(2)
C1-N1-C5	119.6(6)	C1-N1-Pd1	127.5(5)
C5-N1-Pd1	112.9(4)	C1'-N1'-C5'	119.4(6)
C1'-N1'-Pd1	127.1(5)	C5'-N1'-Pd1	113.5(4)
C6-N2-C6'	112.2(5)	C6-N2-Pd1	124.0(4)
C6'-N2-Pd1	123.8(4)	N1-C1-C2	122.1(8)
N1'-C1'-C2'	121.5(8)	C3-C2-C1	118.7(8)
C3'-C2'-C1'	119.2(8)	C2-C3-C4	120.4(8)
C4'-C3'-C2'	120.0(8)	C5-C4-C3	119.4(7)
C3'-C4'-C5'	120.0(8)	C4-C5-N1	119.9(6)
C4-C5-C6	126.6(7)	N1-C5-C6	113.4(5)
N1'-C5'-C4'	119.9(6)	N1'-C5'-C6'	113.3(5)
C4'-C5'-C6'	126.8(7)	N2-C6-C7	105.7(6)
N2-C6-C5	111.5(5)	C7-C6-C5	142.8(6)
N2-C6'-C7'	106.6(6)	N2-C6'-C5'	111.4(5)
C7'-C6'-C5'	142.0(6)	C7'-C7-C6	107.9(6)
C7-C7'-C6'	107.7(6)	O5-S1-O3	118.3(4)
O5-S1-O4	114.4(5)	O3-S1-O4	112.0(4)
O5-S1-C8	104.0(5)	O3-S1-C8	105.2(4)
O4-S1-C8	100.2(5)	F3-C8-F1	109.0(10)
F3-C8-F2	107.4(11)	F1-C8-F2	106.6(9)
F3-C8-S1	113.3(7)	F1-C8-S1	113.0(8)
F2-C8-S1	107.1(8)		

Table X3.6. Anisotropic atomic displacement parameters (Å²) for (PDP)Pd(OH)₂]⁺[OTf]⁻·H₂O (2).

The anisotropic atomic displacement factor exponent takes the form: $-2\pi^2 [h^2 a^{*2} U_{11} + \dots + 2 h k a^* b^* U_{12}]$

	U ₁₁	U ₂₂	U ₃₃	U ₂₃	U ₁₃	U ₁₂
Pd1	0.0496(2)	0.0668(2)	0.0494(2)	0.00502(18)	0.01690(18)	0.0200(2)
N1	0.062(3)	0.070(3)	0.050(2)	0.007(2)	0.018(2)	0.024(2)
N1'	0.061(3)	0.068(3)	0.056(3)	0.009(2)	0.025(2)	0.021(2)
N2	0.050(3)	0.068(3)	0.056(3)	0.009(2)	0.019(2)	0.020(2)
C1	0.092(5)	0.080(5)	0.064(4)	0.008(3)	0.026(3)	0.037(4)
C1'	0.078(5)	0.079(5)	0.071(4)	0.009(3)	0.033(3)	0.018(3)

	U_{11}	U_{22}	U_{33}	U_{23}	U_{13}	U_{12}
C2	0.135(8)	0.088(5)	0.079(5)	0.015(4)	0.040(5)	0.059(5)
C2'	0.129(8)	0.067(5)	0.090(5)	0.019(4)	0.054(5)	0.020(4)
C3	0.128(7)	0.070(4)	0.075(4)	0.006(4)	0.044(5)	0.027(5)
C3'	0.125(7)	0.076(5)	0.087(5)	0.022(4)	0.050(5)	0.041(5)
C4	0.081(5)	0.072(4)	0.060(4)	0.004(3)	0.025(3)	0.015(3)
C4'	0.104(6)	0.092(5)	0.060(4)	0.023(3)	0.034(4)	0.052(4)
C5	0.072(4)	0.066(4)	0.045(3)	0.000(2)	0.025(3)	0.011(3)
C5'	0.082(5)	0.078(4)	0.050(3)	0.013(3)	0.030(3)	0.037(3)
C6	0.056(4)	0.073(4)	0.050(3)	0.003(3)	0.018(2)	0.020(3)
C6'	0.067(4)	0.085(4)	0.049(3)	0.015(3)	0.021(3)	0.037(3)
C7	0.048(4)	0.101(5)	0.062(3)	0.000(3)	0.019(3)	0.017(3)
C7'	0.056(4)	0.096(5)	0.068(4)	0.010(3)	0.021(3)	0.032(3)
O1	0.070(3)	0.100(3)	0.085(3)	0.003(3)	0.024(2)	0.036(3)
O2	0.088(3)	0.076(3)	0.098(3)	0.026(2)	0.043(3)	0.037(2)
S1	0.0658(10)	0.0602(9)	0.0878(11)	0.0132(8)	0.0269(8)	0.0201(7)
F1	0.144(5)	0.061(3)	0.205(6)	-0.010(3)	0.054(4)	0.027(3)
F2	0.437(18)	0.204(9)	0.106(5)	0.009(5)	0.058(8)	0.142(10)
F3	0.184(8)	0.129(5)	0.330(12)	-0.010(6)	0.185(9)	0.031(5)
C8	0.110(8)	0.078(6)	0.137(8)	0.003(5)	0.040(6)	0.023(5)
O3	0.130(5)	0.063(3)	0.125(4)	0.031(3)	0.061(4)	0.039(3)
O4	0.171(7)	0.096(4)	0.091(4)	0.016(3)	-0.019(4)	0.049(4)
O5	0.078(4)	0.119(5)	0.240(8)	0.055(5)	0.069(5)	0.038(3)

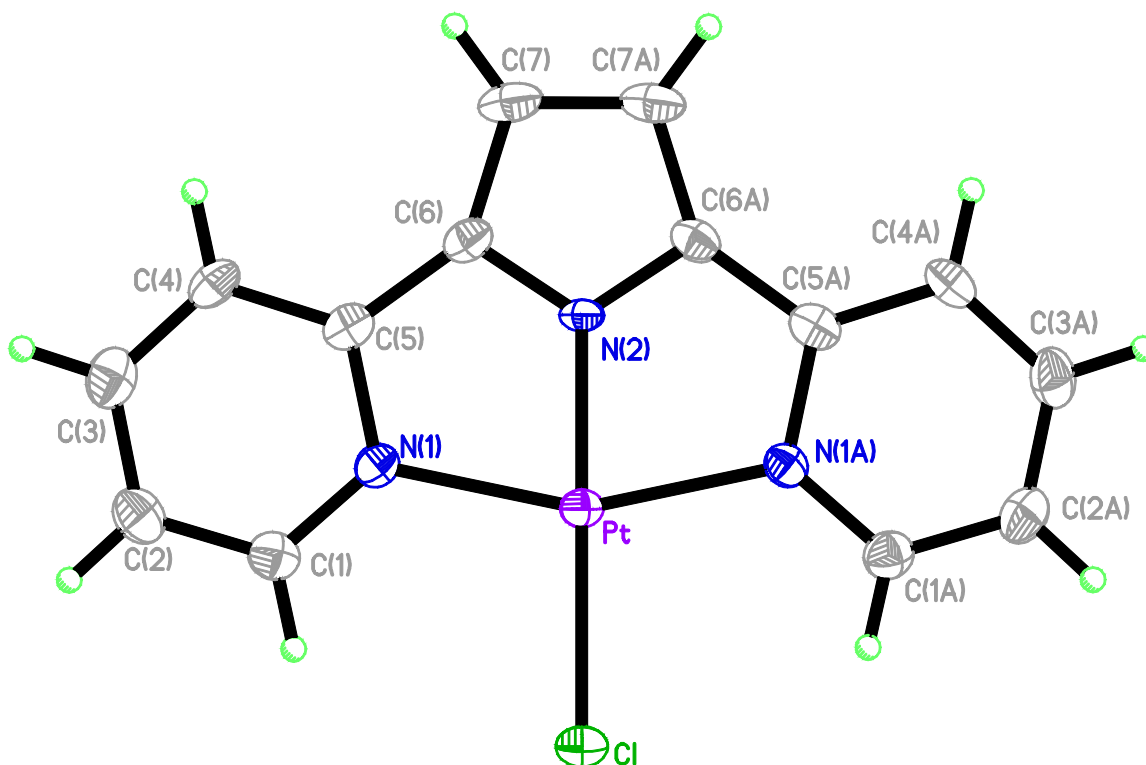


Figure X4. Structure of (PDP)PdCl (**3**) with thermal ellipsoids (50%), and hydrogens shown as open circles.

Table X4.1. Sample and crystal data for (PDP)PtCl.

Chemical formula	C ₁₄ H ₁₀ ClN ₃ Pt	
Formula weight	450.79	
Temperature	143(2) K	
Wavelength	0.71069 Å	
Crystal size	0.020 x 0.020 x 0.480 mm	
Crystal system	monoclinic	
Space group	C2/c	
Unit cell dimensions	a = 10.769(3) Å	α = 90°
	b = 13.456(4) Å	β = 98.944(6)°
	c = 8.751(2) Å	γ = 90°
Volume	1252.7(6) Å ³	
Z	4	
Density (calculated)	2.390 Mg/cm ³	
Absorption coefficient	11.401 mm ⁻¹	
F(000)	840	

Table X4.2. Data collection and structure refinement for (PDP)PtCl.

Theta range for data collection	3.03 to 27.47°
Index ranges	-13<=h<=11, -13<=k<=16, -8<=l<=10
Reflections collected	4100
Independent reflections	1281 [R(int) = 0.0231]
Max. and min. transmission	1.0000 and 0.7353
Structure solution technique	direct methods
Structure solution program	SIR92 (Altomare, 1993)
Refinement method	Full-matrix least-squares on F ²
Refinement program	SHELXL-93 (Sheldrick, 1993)
Function minimized	$\Sigma w(F_o^2 - F_c^2)^2$
Data / restraints / parameters	1281 / 0 / 88
Goodness-of-fit on F²	1.144 ^a
Final R indices^b	1281 data; I>2σ(I) R1 = 0.0209, wR2 = 0.0497 all data R1 = 0.0224, wR2 = 0.0500
Weighting scheme	w=1/[σ ² (F _o ²)+(0.0205P) ² +8.4294P] where P=(F _o ² +2F _c ²)/3
Extinction coefficient	0.0022(5)
Largest diff. peak and hole	0.684 and -1.500 eÅ ⁻³
R.M.S. deviation from mean	0.149 eÅ ⁻³

$$^a S = [\Sigma w(F_{obs}^2 - F_{calc}^2)^2 / (m - n)]^{1/2} \quad ^b R_1 = (\Sigma ||F_{obs}| - |F_{calc}||) / \Sigma |F_{obs}|; \quad wR_2 = [(\Sigma w(F_{obs}^2 - F_{calc}^2)^2) / \Sigma wF_{obs}^2]^{1/2}$$

Table X4.3. Atomic coordinates and equivalent isotropic atomic displacement parameters ($\text{\AA}^2$) for (PDP)PtCl.

U(eq) is defined as one third of the trace of the orthogonalized U_{ij} tensor.

	x/a	y/b	z/c	U(eq)
Pt	0.5	0.42436(2)	0.75	0.01686(9)
Cl	0.5	0.25091(11)	0.75	0.0281(4)
N1	0.6445(3)	0.4556(3)	0.6290(4)	0.0184(7)
N2	0.5	0.5650(3)	0.75	0.0177(11)
C1	0.7164(4)	0.3900(4)	0.5668(5)	0.0240(10)
C2	0.8138(5)	0.4191(4)	0.4900(6)	0.0275(11)
C3	0.8360(4)	0.5194(4)	0.4769(6)	0.0279(11)
C4	0.7638(4)	0.5881(3)	0.5381(6)	0.0233(10)
C5	0.6659(4)	0.5567(3)	0.6138(5)	0.0196(9)
C6	0.5810(4)	0.6196(3)	0.6836(5)	0.0211(9)
C7	0.5501(5)	0.7198(3)	0.7077(6)	0.0286(11)

Table X4.4. Bond lengths ($\text{\AA}$) for (PDP)PtCl.

Pt-N2	1.893(5)	Pt-N1#1	2.057(4)
Pt-N1	2.057(4)	Pt-Cl	2.334(2)
N1-C1	1.344(6)	N1-C5	1.390(6)
N2-C6#1	1.340(5)	N2-C6	1.340(5)
C1-C2	1.387(7)	C2-C3	1.379(7)
C3-C4	1.370(7)	C4-C5	1.395(6)
C5-C6	1.449(6)	C6-C7	1.412(7)
C7-C7#1	1.400(10)		

Symmetry transformations used to generate equivalent atoms:

#1 -x+1, y, -z+3/2

Table X4.5. Bond angles (°) for (PDP)PtCl.

N2-Pt-N1#1	78.22(11)	N2-Pt-N1	78.22(11)
N1#1-Pt-N1	156.4(2)	N2-Pt-Cl	180.0
N1#1-Pt-Cl	101.78(11)	N1-Pt-Cl	101.78(11)
C1-N1-C5	119.3(4)	C1-N1-Pt	127.2(3)
C5-N1-Pt	113.5(3)	C6#1-N2-C6	113.5(5)
C6#1-N2-Pt	123.2(3)	C6-N2-Pt	123.2(3)
N1-C1-C2	122.5(5)	C3-C2-C1	118.1(5)
C4-C3-C2	120.7(4)	C3-C4-C5	119.9(4)
N1-C5-C4	119.4(4)	N1-C5-C6	114.0(4)
C4-C5-C6	126.6(4)	N2-C6-C7	105.9(4)
N2-C6-C5	111.0(4)	C7-C6-C5	143.1(4)
C7#1-C7-C6	107.3(3)		

Symmetry transformations used to generate equivalent atoms:

#1 -x+1, y, -z+3/2

Table 6. Anisotropic atomic displacement parameters (Å²) for (PDP)PtCl.

The anisotropic atomic displacement factor exponent takes the form: $-2\pi^2 [h^2 a^{*2} U_{11} + \dots + 2 h k a^* b^* U_{12}]$						
	U ₁₁	U ₂₂	U ₃₃	U ₂₃	U ₁₃	U ₁₂
Pt	0.01706(13)	0.01443(14)	0.0186(2)	0	0.00127(9)	0
Cl	0.0332(9)	0.0158(8)	0.0373(10)	0	0.0117(7)	0
N1	0.017(2)	0.019(2)	0.018(2)	0.0004(14)	0.0008(14)	-0.0020(15)
N2	0.020(3)	0.010(2)	0.023(3)	0	0.004(2)	0
C1	0.022(2)	0.023(2)	0.026(3)	-0.001(2)	0.001(2)	0.001(2)
C2	0.022(2)	0.031(3)	0.029(3)	-0.003(2)	0.002(2)	0.005(2)
C3	0.017(2)	0.037(3)	0.030(3)	0.005(2)	0.002(2)	0.000(2)
C4	0.023(2)	0.023(2)	0.023(3)	0.003(2)	0.001(2)	-0.008(2)
C5	0.021(2)	0.021(2)	0.014(2)	0.001(2)	-0.004(2)	-0.004(2)
C6	0.025(2)	0.019(2)	0.018(2)	0.002(2)	0.002(2)	-0.004(2)
C7	0.044(3)	0.015(2)	0.026(3)	0.001(2)	0.001(2)	-0.004(2)

References