Supporting Information for:

Synthesis of Pyrrole-Based PSiP Pincer Ligands and Their Palladium, Rhodium, and Platinum Complexes

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I. Isomerization of 7A/7B

Figure S1. Time-course of reaction of $(Ph_3P)_3Rh(H)CO$ with **2** in C₆D₆ as monitored by ${}^{31}P{}^{1}H$ NMR.

Procedure: A J. Young tube was charged with $(Ph_3P)_3Rh(H)CO$ (18 mg, 0.020 mmol), proligand **2** (11 mg, 0.020 mmol), and benzene- d_6 (0.6 mL). ¹H and ³¹P{H} spectra were acquired periodically. The ratio of **7A** to **7B** was found to be 1:2.6 and 1:2.7 in two experiments over the course of 3 to 6 days



43 42 41 40 39 38 37 36 35 34 33 32 31 30 29 28 27 26 25 24 23 22 21 20 19 18 17 16 15 14 13 12 11 10 9 8 7 δ(ppm)



Figure S2. Conversion of crystals of **7B** to mixture of **7A/7B** in methylene chloride- d_2 as monitored by ³¹P{¹H} NMR (1.0:0.9 **7A:7B**). Equilibration was also observed in benzene- d_6 in which **7B** was the major species (1:3.2 of **7A:7B**).



Figure S3. Conversion of crystals of 7B to mixture of 7A/7B in methylene chloride- d_2 as monitored by ¹H NMR (1.0:0.9 7A:7B). Equilibration was also observed in benzene- d_6 in which 7B was the major species (1:3.2 of 7A:7B). Pentane is indicated with an asterisk (*).

II. Buried Volume Details for [(^{Ph}P_{Py})₂Si]PdCl (4) and [(^{Ph}P_{Ph})₂Si]PdCl

Buried volume calculations were performed by inputting the following information into the SambVca 2.1 web application¹ based on the crystal structures of $[({}^{Ph}P_{Py})_2Si]PdCl$ (4) and $[({}^{Ph}P_{Ph})_2Si]PdCl.^2$

- 1. Load file: .xyz file of crystal structure used
- 2. Select the atoms coordinated to the center of the sphere: Palladium selected
- 3. Select the atoms for z-axis definition: Silicon selected with z-negative orientation
- 4. Select the atoms for xz-plane definition: Phosphorus selected that is to the right of the methyl group oriented upward.
- 5. Select the atoms to be deleted: Palladium and chloride selected.
- 6. Check the chosen orientation:



 $[(^{Ph}P_{Py})_2Si]PdCl (4)$



[(^{Ph}P_{Ph})₂Si]PdCl

- 7. Select the atomic radii: Bond scaled by 1.17
- 8. Sphere radius: 3.5
- 9. Distance of the coordination point from the center of the sphere: 0.0
- 10. Mesh spacing for numerical integration: 0.10
- 11. Inclusion of H atoms: H atoms included



Figure S4: Steric map of $[(^{Ph}P_{Py})_2Si]PdCl; \%V_{Bur} = 71.4\%$



Figure S5: Steric map of $[(^{Ph}P_{Ph})_2Si]PdCl; \%V_{Bur} = 71.1\%$

III. Cyclic Voltammograms of [(^{Ph}P_{Py})₂Si]PdCl (4) and [(^{Ph}P_{Ph})₂Si]PdCl



Figure S6. Cyclic voltammogram of $[({}^{Ph}P_{Py})_2Si]PdCl(4)$ in THF. Conditions: scan rate: 300 mV/s; supporting electrolyte: 0.1 M [*n*-Bu₄N][PF₆]; working electrode: glassy carbon; auxiliary electrode: platinum wire; reference: silver wire referenced to the Fc^{+/0} couple.



Figure S7. Cyclic voltammogram of $[({}^{Ph}P_{Ph})_2Si]PdC1$ in THF. Conditions: scan rate: 300 mV/s; supporting electrolyte: 0.1 M [*n*-Bu₄N][PF₆]; working electrode: glassy carbon; auxiliary electrode: platinum wire; reference: silver wire referenced to the Fc^{+/0} couple.



IV. IR Spectra of [(^{Ph}P_{Py})₂Si]Rh(CO)PPh₃ (7A/7B)

Figure S8. Infrared spectrum of an equilibrium mixture of 7A/7B.



Figure S9. Infrared spectrum of crystalline 7B. Material was dissolved in DCM and immediately deposited as a thin film.

Reference	Linker	Metal	Е	Apical CO (cm ⁻¹)	$^{2}J_{P-P}$ (Hz)	Equatorial CO (cm ⁻¹)	$^{2}J_{P-P}$ (Hz)
3	Phenylene	Rh	SiMe	-	-	1897	34.9
4	Phenylene	Rh	GeMe	1968	99.1	1923	33.0
4	Phenylene	Rh	SnMe	1964	102.8	-	-
3	Phenylene	Ir	SiMe	1959	100.3	1918	25.7
5	Phenylene	Ir	SiF	1969	82.2	1937	28.0
This work	Pyrrole	Rh	SiMe	1977	88.0	1932	38.2

Table S1: Comparison of CO stretches and ${}^{2}J_{P-P}$ (PPh₂) for PEP complexes analogous to 7A/7B.

 $\begin{array}{c} Ph_2 & E \\ Ph_2 P & M - PPh_3 \\ Ph_2 P & I \\ CO \end{array} \qquad \begin{array}{c} Ph_2 & Ph_2 \\ Ph_2 P & M - CO \\ Ph_2 P & I \\ PPh_3 \end{array}$

Apical CO

Equatorial СО

V. NMR Spectra of Proligands



Figure S10. ¹H NMR spectrum of bis(2-diisopropylphosphino-pyrrole)methylsilane (1) in C_6D_6 (400 MHz).



Figure S11. ¹³C{¹H} NMR spectrum of bis(2-diisopropylphosphino-pyrrole)methylsilane (1) in C₆D₆ (100 MHz).



Figure S12. ³¹P{¹H} NMR spectrum of bis(2-diisopropylphosphino-pyrrole)methylsilane (1) in C_6D_6 (206 MHz).



Figure S13. ¹H NMR spectrum of bis(2-diphenylphosphino-pyrrole)methylsilane (2) in CD₂Cl₂ (500 MHz).



Figure S14. ${}^{13}C{}^{1}H$ NMR spectrum of bis(2-diphenylphosphino-pyrrole)methylsilane (2) in CD₂Cl₂ (126 MHz).



Figure S15. ${}^{31}P{}^{1}H$ NMR spectrum of bis(2-diphenylphosphino-pyrrole)methylsilane (2) in CD₂Cl₂ (202 MHz).



VI. NMR Spectra of Transition Metal Complexes

Figure S16. ¹H NMR spectrum of $[({}^{iPr}P_{Py})_2Si]PdCl (3)$ in C₆D₆ (400 MHz).



Figure S17. ¹³C{¹H} NMR spectrum of $[(^{iPr}P_{Py})_2Si]PdCl$ (3) in C₆D₆ (126 MHz).



Figure S18. ${}^{31}P{}^{1}H$ NMR spectrum of $[({}^{iPr}P_{Py})_2Si]PdCl(3)$ in C₆D₆ (202 MHz).



Figure S19. ¹H NMR spectrum of $[(^{Ph}P_{Py})_2Si]PdCl$ (4) in CD₂Cl₂ (500 MHz).



Figure S20. ${}^{13}C{}^{1}H$ NMR spectrum of $[({}^{Ph}P_{Py})_2Si]PdCl (4)$ in CD₂Cl₂ (126 MHz).



Figure S21. ${}^{31}P{}^{1}H$ NMR spectrum of $[({}^{Ph}P_{Py})_2Si]PdCl (4)$ in CD₂Cl₂ (126 MHz).



Figure S22. ¹H NMR spectrum of $[({}^{Ph}P_{Py})_2Si]PtCl$ (5) in CD₂Cl₂ (500 MHz).



Figure S23. ${}^{13}C{}^{1}H$ NMR spectrum of $[({}^{Ph}P_{Py})_2Si]PtCl (5)$ in $CD_2Cl_2 (126 \text{ MHz})$.



Figure S24. ²⁹Si $\{^{1}H\}$ NMR spectrum of [($^{Ph}P_{Py}$)₂Si]PtCl (5) in CD₂Cl₂ (99 MHz).



Figure S25. ${}^{31}P{}^{1}H$ NMR spectrum of [(${}^{Ph}P_{Py}$)₂Si]PtCl (5) in CD₂Cl₂ (126 MHz).



Figure S26. Crude ¹H NMR spectrum of *cis*-bis[(2-diphenylphosphino)pyrrole] platinum dichloride in CD₂Cl₂ (400 MHz).



Figure S27. ${}^{31}P{}^{1}H$ NMR spectrum of *cis*-bis[(2-diphenylphosphino)pyrrole] platinum dichloride in CD₂Cl₂ (162 MHz).



Figure S28. ¹H NMR spectrum of $[({}^{iPr}P_{Py})_2Si]Rh(H)Cl$ (6) in C₆D₆ (400 MHz).



Figure S29. ¹³C{¹H} NMR spectrum of $[({}^{iPr}P_{Py})_2Si]Rh(H)Cl$ (6) in CDCl₃ (126 MHz).



Figure S30. ${}^{31}P{}^{1}H$ NMR spectrum of $[({}^{iPr}P_{Py})_2Si]Rh(H)Cl$ (6) in C₆D₆ (206 MHz).



Figure S31. ¹H NMR spectrum of [(^{Ph}P_{Py})₂Si]Rh(CO)PPh₃ (7A/7B) in CD₂Cl₂ (500 MHz).



Figure S32. ¹³C{¹H} NMR spectrum of $[({}^{Ph}P_{Py})_{2}Si]Rh(CO)PPh_{3}$ (7A/7B) in CD₂Cl₂ (126 MHz).



Figure S33. ${}^{31}P{}^{1}H$ NMR spectrum of $[({}^{Ph}P_{Py})_2Si]Rh(CO)PPh_3$ (7A/7B) in CD₂Cl₂ (126 MHz).

VII. X-Ray Crystallographic Data

X-Ray data collection, solution, and refinement for [(^{Ph}P_{Py})₂Si]PdCl (4; CCDC 2143700).

A Leica MZ 75 microscope was used to identify a suitable colorless block with very well defined faces with dimensions (max, intermediate, and min) 0.478 x 0.452 x 0.207 mm³ from a representative sample of crystals of the same habit. The crystal mounted on a nylon loop was then placed in a cold nitrogen stream (Oxford) maintained at 110 K. A BRUKER Quest X-ray (fixed-Chi geometry) diffractometer with a PHOTON III detector was employed for crystal screening, unit cell determination, and data collection. The goniometer was controlled using the APEX3 software suite.⁶ The sample was optically centered with the aid of a video camera such that no translations were observed as the crystal was rotated through all positions. The X-ray radiation employed was generated from a Mo-Iµs X-ray tube ($K_{\alpha} = 0.71073$ Å). 45 data frames were taken at widths of 1°. These reflections were used to determine the unit cell. The unit cell was verified by examination of the h k l overlays on several frames of data. No super-cell or erroneous reflections were observed. After careful examination of the unit cell, an extended data collection procedure (8 sets) was initiated using omega and phi scans. Integrated intensity information for each reflection was obtained by reduction of the data frames with the program APEX3. The integration method employed a three dimensional profiling algorithm and all data were corrected for Lorentz and polarization factors, as well as for crystal decay effects. Finally the data was merged and scaled to produce a suitable data set. The absorption correction program SADABS⁷ was employed to correct the data for absorption effects. Systematic reflection conditions and statistical tests of the data suggested the space group $P2_1/c$. A solution was obtained readily using XT/XS in APEX3.^{6,8} Hydrogen atoms were placed in idealized positions and were set riding on the respective parent atoms. All non-hydrogen atoms were refined with anisotropic thermal parameters. Absence of additional symmetry and voids were confirmed using PLATON (ADDSYM).⁹ The structure was refined (weighted least squares refinement on F^2) to convergence.^{8,10} Olex2 was employed for the final data presentation and structure plots.¹⁰

X-Ray data collection, solution, and refinement for [(^{Ph}P_{Py})₂Si]PtCl (5; CCDC 2143692).

A Leica MZ 75 microscope was used to identify a suitable colorless block with very well defined faces with dimensions (max, intermediate, and min) 0.402 x 0.381 x 0.217 mm³ from a representative sample of crystals of the same habit. The crystal mounted on a nylon loop was then placed in a cold nitrogen stream (Oxford) maintained at 110 K. A BRUKER Quest X-ray (fixed-Chi geometry) diffractometer with a PHOTON III detector was employed for crystal screening, unit cell determination, and data collection. The goniometer was controlled using the APEX3 software suite.⁶ The sample was optically centered with the aid of a video camera such that no translations were observed as the crystal was rotated through all positions. The X-ray radiation employed was generated from a Mo-Iµs X-ray tube ($K_{\alpha} = 0.71073$ Å). 45 data frames were taken at widths of 1°. These reflections were used to determine the unit cell. The unit cell was verified by examination of the *h k l* overlays on several frames of data. No super-cell or erroneous

reflections were observed. After careful examination of the unit cell, an extended data collection procedure (8 sets) was initiated using omega and phi scans. Integrated intensity information for each reflection was obtained by reduction of the data frames with the program APEX3. The integration method employed a three dimensional profiling algorithm and all data were corrected for Lorentz and polarization factors, as well as for crystal decay effects. Finally the data was merged and scaled to produce a suitable data set. The absorption correction program SADABS⁷ was employed to correct the data for absorption effects. Systematic reflection conditions and statistical tests of the data suggested the space group $P2_1/c$. A solution was obtained readily using XT/XS in APEX3.^{6,8} Hydrogen atoms were placed in idealized positions and were set riding on the respective parent atoms. All non-hydrogen atoms were refined with anisotropic thermal parameters. Absence of additional symmetry and voids were confirmed using PLATON (ADDSYM).⁹ The structure was refined (weighted least squares refinement on F^2) to convergence.^{8,10} Olex2 was employed for the final data presentation and structure plots.¹⁰

X-Ray data collection, solution, and refinement for *cis*-bis[(2-diphenylphosphino)pyrrole] platinum dichloride (CCDC 2143693).

A Leica MZ 75 microscope was used to identify a suitable colorless block with very well defined faces with dimensions (max, intermediate, and min) 0.207 x 0.124 x 0.081 mm³ from a representative sample of crystals of the same habit. The crystal mounted on a nylon loop was then placed in a cold nitrogen stream (Oxford) maintained at 110 K. A BRUKER Quest X-ray (fixed-Chi geometry) diffractometer with a PHOTON III detector was employed for crystal screening, unit cell determination, and data collection. The goniometer was controlled using the APEX3 software suite.⁶ The sample was optically centered with the aid of a video camera such that no translations were observed as the crystal was rotated through all positions. The X-ray radiation employed was generated from a Mo-Iµs X-ray tube ($K_{\alpha} = 0.71073$ Å). 45 data frames were taken at widths of 1°. These reflections were used to determine the unit cell. The unit cell was verified by examination of the h k l overlays on several frames of data. No super-cell or erroneous reflections were observed. After careful examination of the unit cell, an extended data collection procedure (5 sets) was initiated using omega and phi scans. Integrated intensity information for each reflection was obtained by reduction of the data frames with the program APEX3. The integration method employed a three dimensional profiling algorithm and all data were corrected for Lorentz and polarization factors, as well as for crystal decay effects. Finally the data was merged and scaled to produce a suitable data set. The absorption correction program SADABS⁷ was employed to correct the data for absorption effects. Systematic reflection conditions and statistical tests of the data suggested the space group P-1. A solution was obtained readily using XT/XS in APEX3.^{6,8} Hydrogen atoms were placed in idealized positions and were set riding on the respective parent atoms (except hydrogen atoms bonded to N1 and N2). All non-hydrogen atoms were refined with anisotropic thermal parameters. Absence of additional symmetry and voids were confirmed using PLATON (ADDSYM).9 The structure was refined (weighted least squares refinement on F^2) to convergence.^{8,10} Olex2 was employed for the final data presentation and structure plots.¹⁰

X-Ray data collection, solution, and refinement for [(^{Ph}P_{Py})₂Si]Rh(CO)PPh₃ (7; CCDC 2143691).

A Leica MZ 75 microscope was used to identify a suitable yellow block with very well defined faces with dimensions (max, intermediate, and min) 0.421 x 0.387 x 0.204 mm³ from a representative sample of crystals of the same habit. The crystal mounted on a nylon loop was then placed in a cold nitrogen stream (Oxford) maintained at 110 K. A BRUKER Quest X-ray (fixed-Chi geometry) diffractometer with a PHOTON III detector was employed for crystal screening, unit cell determination, and data collection. The goniometer was controlled using the APEX3 software suite.⁶ The sample was optically centered with the aid of a video camera such that no translations were observed as the crystal was rotated through all positions. The X-ray radiation employed was generated from a Mo-Iµs X-ray tube ($K_{\alpha} = 0.71073$ Å). 45 data frames were taken at widths of 1°. These reflections were used to determine the unit cell. The unit cell was verified by examination of the h k l overlays on several frames of data. No super-cell or erroneous reflections were observed. After careful examination of the unit cell, an extended data collection procedure (10 sets) was initiated using omega and phi scans. Integrated intensity information for each reflection was obtained by reduction of the data frames with the program APEX3. The integration method employed a three dimensional profiling algorithm and all data were corrected for Lorentz and polarization factors, as well as for crystal decay effects. Finally the data was merged and scaled to produce a suitable data set. The absorption correction program SADABS⁷ was employed to correct the data for absorption effects. Systematic reflection conditions and statistical tests of the data suggested the space group P-1. A solution was obtained readily using XT/XS in APEX3.^{6,8} Hydrogen atoms were placed in idealized positions and were set riding on the respective parent atoms. All non-hydrogen atoms were refined with anisotropic thermal parameters. Absence of additional symmetry and voids were confirmed using PLATON (ADDSYM).⁹ The structure was refined (weighted least squares refinement on F^2) to convergence.^{8,10} Olex2 was employed for the final data presentation and structure plots.¹⁰

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