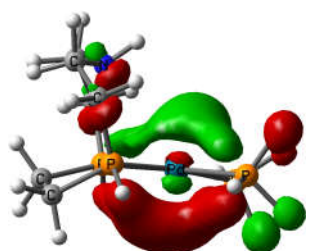
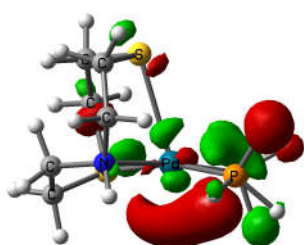


## Supplementary Information

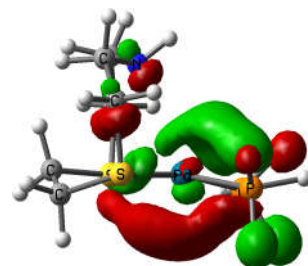
**Figure S1.** Geometric structures of Pd complexes [*endo*-Pd([9]aneA<sub>2</sub>B)L<sub>2</sub>, *endo*-Pd([9]aneABA)L<sub>2</sub> {A=P, S; B=N; L=Cl, PH<sub>3</sub>, P(CH<sub>3</sub>)<sub>3</sub>}] including the 3a<sub>1g</sub>(5s) and 2a<sub>1g</sub> orbitals optimized at the B3P86//6-311+G(d,p)/(lanl2DZ for Pd) levels. The *endo*-[Pd([9]anePNP)(PH<sub>3</sub>)<sub>2</sub>]<sup>2+</sup> and *endo*-[Pd([9]anePNP){P(CH<sub>3</sub>)<sub>3</sub>}<sub>2</sub>]<sup>2+</sup> complexes could not be optimized.



*endo*-[Pd([9]aneP<sub>2</sub>N)(PH<sub>3</sub>)<sub>2</sub>]<sup>2+</sup>

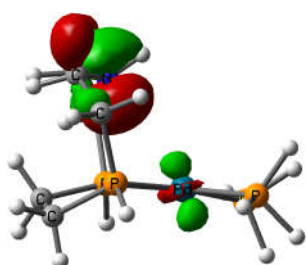


*endo*-[Pd([9]aneSNS)(PH<sub>3</sub>)<sub>2</sub>]<sup>2+</sup>

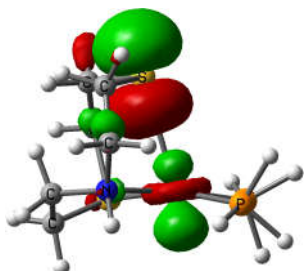


*endo*-[Pd([9]aneS<sub>2</sub>N)(PH<sub>3</sub>)<sub>2</sub>]<sup>2+</sup>

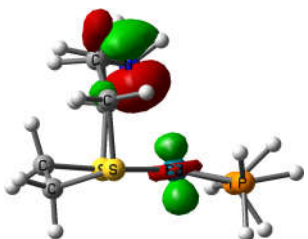
**3a<sub>1g</sub>(5s)**



*endo*-[Pd([9]aneP<sub>2</sub>N)(PH<sub>3</sub>)<sub>2</sub>]<sup>2+</sup>

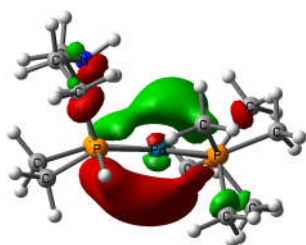


*endo*-[Pd([9]aneSNS)(PH<sub>3</sub>)<sub>2</sub>]<sup>2+</sup>

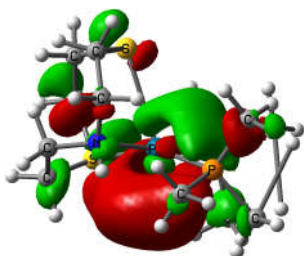


*endo*-[Pd([9]aneS<sub>2</sub>N)(PH<sub>3</sub>)<sub>2</sub>]<sup>2+</sup>

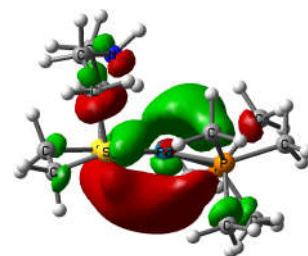
**2a<sub>1g</sub>**



*endo*-[Pd([9]aneP<sub>2</sub>N){P(CH<sub>3</sub>)<sub>3</sub>}<sub>2</sub>]<sup>2+</sup>



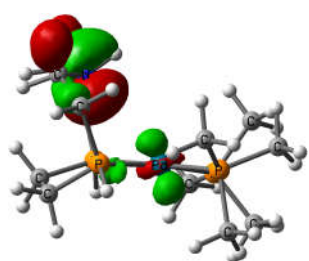
*endo*-[Pd([9]aneSNS){P(CH<sub>3</sub>)<sub>3</sub>}<sub>2</sub>]<sup>2+</sup>



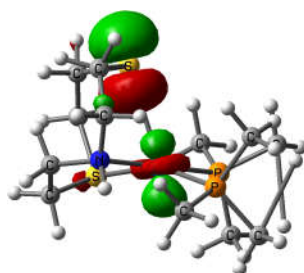
*endo*-[Pd([9]aneS<sub>2</sub>N){P(CH<sub>3</sub>)<sub>3</sub>}<sub>2</sub>]<sup>2+</sup>

**3a<sub>1g</sub>(5s)**

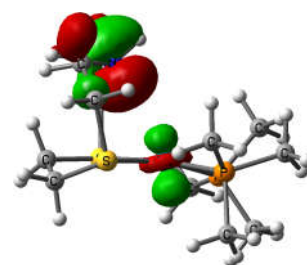
Figure S1. Cont.



*endo*-[Pd([9]aneP<sub>2</sub>N){P(CH<sub>3</sub>)<sub>3</sub>}<sub>2</sub>]<sup>2+</sup>

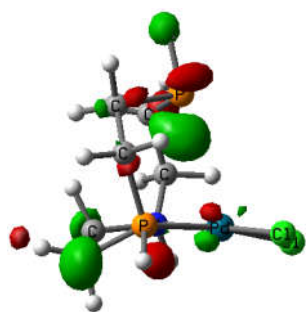


*endo*-[Pd([9]aneSNS){P(CH<sub>3</sub>)<sub>3</sub>}<sub>2</sub>]<sup>2+</sup>

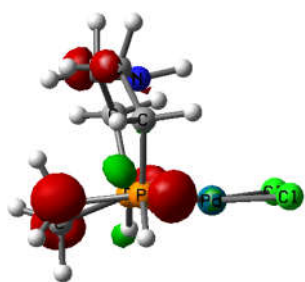


*endo*-[Pd([9]aneS<sub>2</sub>N){P(CH<sub>3</sub>)<sub>3</sub>}<sub>2</sub>]<sup>2+</sup>

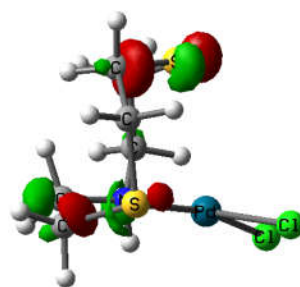
**2a<sub>1g</sub>**



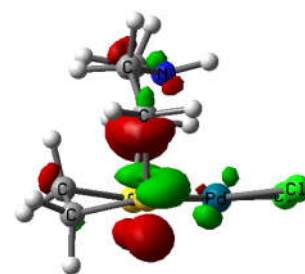
*endo*-Pd([9]anePNP)Cl<sub>2</sub>



*endo*-Pd([9]aneP<sub>2</sub>N)Cl<sub>2</sub>

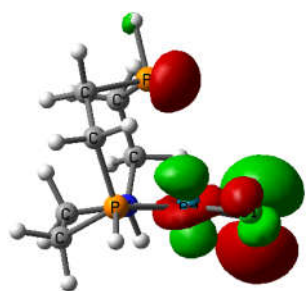


*endo*-Pd([9]aneSNS)Cl<sub>2</sub>

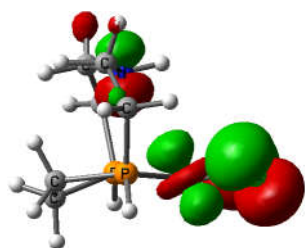


*endo*-Pd([9]aneS<sub>2</sub>N)Cl<sub>2</sub>

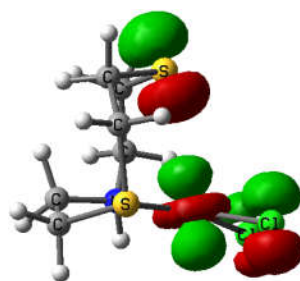
**3a<sub>1g</sub> (5s)**



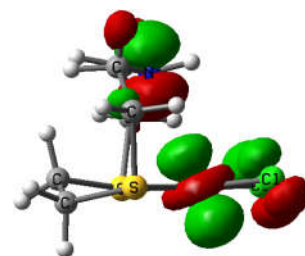
*endo*-Pd([9]anePNP)Cl<sub>2</sub>



*endo*-Pd([9]aneP<sub>2</sub>N)Cl<sub>2</sub>



*endo*-Pd([9]aneSNS)Cl<sub>2</sub>



*endo*-Pd([9]aneS<sub>2</sub>N)Cl<sub>2</sub>

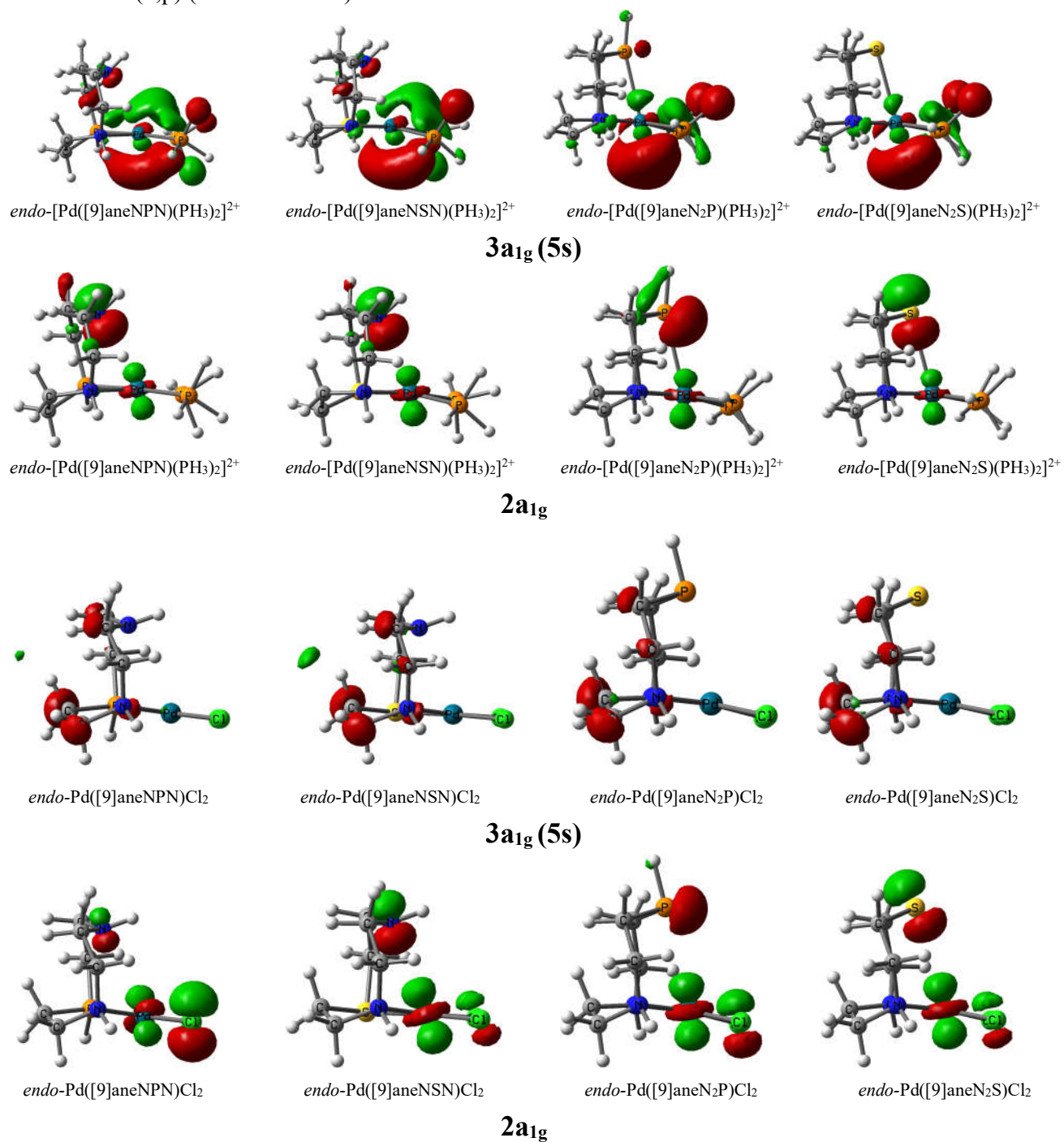
**2a<sub>1g</sub>**

**Table S1.** Optimized average bond distances (Å), average atomic charges (CHelpG, au), and relative energies (eV) of the equilibrium structures {Pd([9]aneA<sub>2</sub>B)L<sub>2</sub>} and [Pd([9]aneABA)L<sub>2</sub>] at the B3P86//6-311+G(d,p) (lanl2DZ for Pd) levels.

Compound	Average distance				Average atomic charge				Relative energy	
	R <sup>a</sup> <sub>Pd-P</sub>	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>b</sup> <sub>Pd...P</sub>	R <sup>b</sup> <sub>Pd...N</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>PH3</sub>	Q <sup>d</sup> <sub>P</sub>	Q <sup>d</sup> <sub>N</sub>	ΔE <sup>e</sup> <sub>H-L</sub>	ΔE <sup>f</sup> <sub>BAB-B2A</sub>
<i>endo</i> -[Pd([9]aneP <sub>2</sub> N)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.342			3.217	0.448	0.343		-0.185	3.82	
	R <sup>a</sup> <sub>Pd-S</sub>	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>b</sup> <sub>Pd...S</sub>	R <sup>b</sup> <sub>Pd...N</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>PH3</sub>	Q <sup>d</sup> <sub>S</sub>	Q <sup>d</sup> <sub>N</sub>		
<i>endo</i> -[Pd([9]aneSNS)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.369	2.161	2.867		0.377	0.359	-0.32 2		2.96	0.01
<i>endo</i> -[Pd([9]aneS <sub>2</sub> N)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.373			2.895	0.396	0.363		-0.239	3.38	0.00
	R <sup>a</sup> <sub>Pd-P</sub>	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>b</sup> <sub>Pd...P</sub>	R <sup>b</sup> <sub>Pd...N</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>P(CH3)3</sub>	Q <sup>d</sup> <sub>P</sub>	Q <sup>d</sup> <sub>N</sub>		
<i>endo</i> - [Pd([9]aneP <sub>2</sub> N){P(CH <sub>3</sub> ) <sub>3</sub> ] <sub>2</sub> ] <sup>2+</sup>	2.364			3.290	0.633	-0.461		0.180	4.31	
	R <sup>a</sup> <sub>Pd-S</sub>	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>b</sup> <sub>Pd...S</sub>	R <sup>b</sup> <sub>Pd...N</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>P(CH3)3</sub>	Q <sup>d</sup> <sub>S</sub>	Q <sup>d</sup> <sub>N</sub>		
<i>endo</i> - [Pd([9]aneSNS){P(CH <sub>3</sub> ) <sub>3</sub> ] <sub>2</sub> ] <sup>2+</sup>	2.424	2.193	2.916		0.456	0.398	- 0.063		3.51	0.02
<i>endo</i> - [Pd([9]aneS <sub>2</sub> N){P(CH <sub>3</sub> ) <sub>3</sub> ] <sub>2</sub> ] <sup>2+</sup>	2.410			3.136	0.482	0.392		-0.133	3.84	0.00
	R <sup>a</sup> <sub>Pd-P</sub>	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>b</sup> <sub>Pd...P</sub>	R <sup>b</sup> <sub>Pd...N</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>Cl</sub>	Q <sup>d</sup> <sub>P</sub>	Q <sup>d</sup> <sub>N</sub>	ΔE <sup>e</sup> <sub>H-L</sub>	ΔE <sup>f</sup> <sub>BAB-B2A</sub>
<i>endo</i> -Pd([9]anePNP)Cl <sub>2</sub>	2.209	2.130	3.175		0.929	-0.795	-0.28 5		3.92	0.55
<i>endo</i> -Pd([9]aneP <sub>2</sub> N)Cl <sub>2</sub>	2.247			3.072	0.945	-0.820		-0.191	4.14	0.00
	R <sup>a</sup> <sub>Pd-S</sub>	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>b</sup> <sub>Pd...S</sub>	R <sup>b</sup> <sub>Pd...N</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>Cl</sub>	Q <sup>d</sup> <sub>S</sub>	Q <sup>d</sup> <sub>N</sub>		
<i>endo</i> -Pd([9]aneSNS)Cl <sub>2</sub>	2.282	2.132	3.093		0.788	-0.755	-0.24 6		3.38	0.23
<i>exptl</i>	2.263 <sup>g</sup>	2.087 <sup>g</sup>	2.928 <sup>g</sup> 3.011 <sup>h</sup>							
<i>endo</i> -Pd([9]aneS <sub>2</sub> N)Cl <sub>2</sub>	2.297			2.904	0.884	-0.770		-0.205	3.50	0.00
<i>exptl</i> <sup>g</sup>	2.269			2.722						

<sup>a</sup>Bond length between the Pd<sup>II</sup> center and equatorial binding site of the tridentate ligand. <sup>b</sup>Bond length between the Pd<sup>II</sup> center and axial binding site of the tridentate ligand. <sup>c</sup>Atomic charges of the Pd<sup>II</sup> center and the binding atom of *trans* L-ligand. <sup>d</sup>Atomic charge of an apical binding atom. <sup>e</sup>Energy gap between HOMO and LUMO. <sup>f</sup>Relative energy gap between Pd([9]aneA<sub>2</sub>B)L<sub>2</sub> and Pd([9]aneABA)L<sub>2</sub>. <sup>g</sup>Ref. [39]. <sup>h</sup>Cited from Ref. [39].

**Figure S2.** The  $3a_{1g}(5s)$  and  $2a_{1g}$  orbitals of  $endo$ -Pd([9]ane $B_2A$ ) $L_2$  and  $endo$ -Pd([9]ane $BAB$ ) $L_2$  (A=P, S; B=N; L=PH $_3$ , Cl $^-$ ) calculated at the CAM-B3LYP/6-311+G(d,p) (lan12DZ for Pd) level.

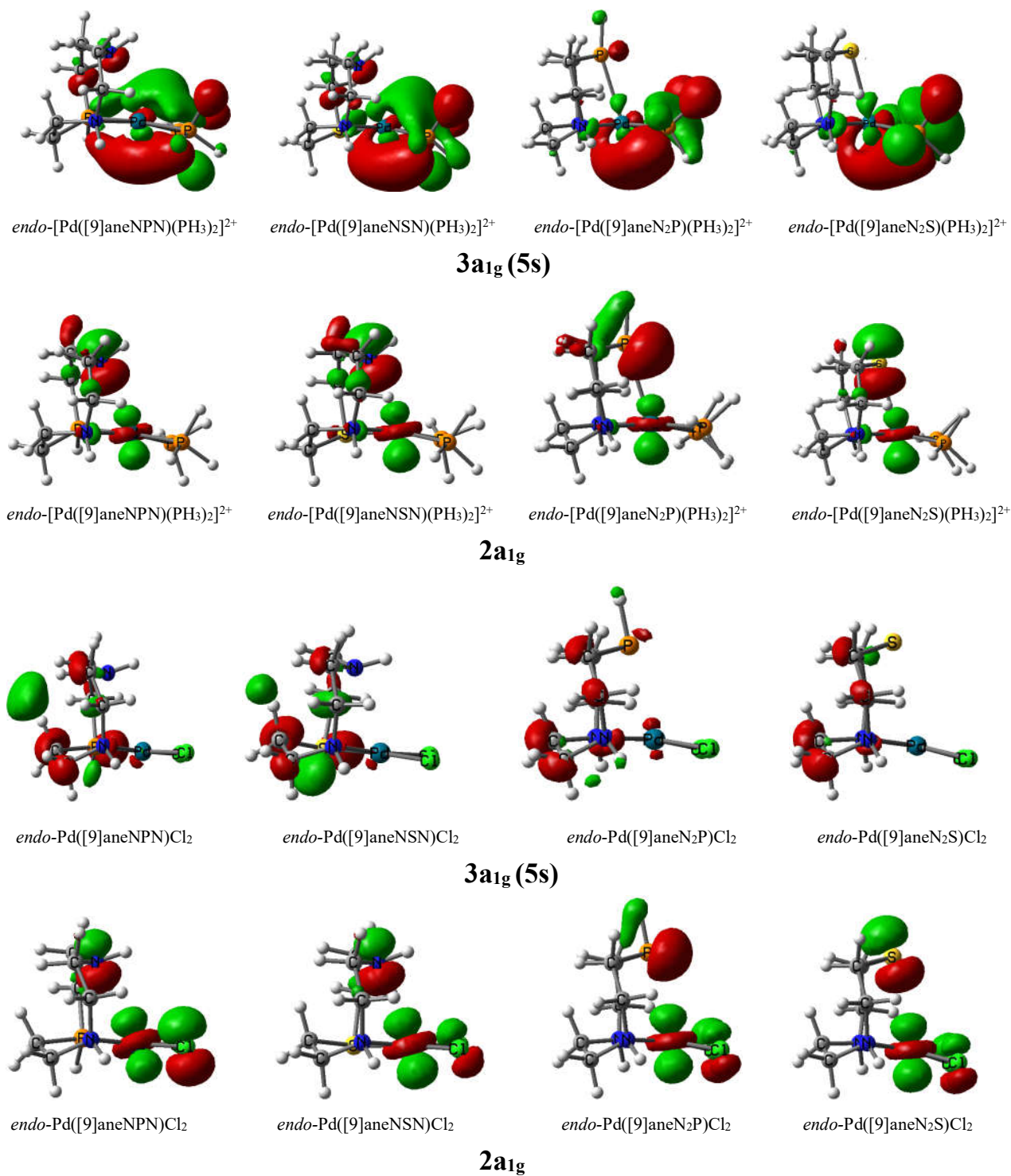


**Table S2.** Optimized average bond distances (Å), average atomic charges (NBO, au), and relative energies (eV) of the equilibrium structures {Pd([9]aneB<sub>2</sub>A)L<sub>2</sub>} and {Pd([9]aneBAB)L<sub>2</sub>} at the CAM-B3LYP/6-311+G(d,p) (lanl2DZ for Pd) levels.

Compound	Average distance				Average atomic charge				Relative energy	
	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>a</sup> <sub>Pd-P</sub>	R <sup>b</sup> <sub>Pd...N</sub>	R <sup>b</sup> <sub>Pd...P</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>PH3</sub>	Q <sup>d</sup> <sub>N</sub>	Q <sup>d</sup> <sub>P</sub>	ΔE <sup>e</sup> <sub>H-L</sub>	ΔE <sup>f</sup> <sub>BAB-B2A</sub>
<i>endo</i> -[Pd([9]aneNPN)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.191	2.296	2.898		0.186	0.301	-0.691		6.37	0.35
<i>endo</i> -[Pd([9]aneN <sub>2</sub> P)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.137			2.879	0.044	0.340		0.608	5.68	0.00
	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>a</sup> <sub>Pd-S</sub>	R <sup>b</sup> <sub>Pd...N</sub>	R <sup>b</sup> <sub>Pd...S</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>PH3</sub>	Q <sup>d</sup> <sub>N</sub>	Q <sup>d</sup> <sub>S</sub>		
<i>endo</i> -[Pd([9]aneNSN)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.156	2.364	2.771		0.115	0.337	-0.686		6.13	0.11
<i>endo</i> -[Pd([9]aneN <sub>2</sub> S)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.136			2.931	0.019	0.339		0.314	5.67	0.00
	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>a</sup> <sub>Pd-P</sub>	R <sup>b</sup> <sub>Pd...N</sub>	R <sup>b</sup> <sub>Pd...P</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>Cl</sub>	Q <sup>d</sup> <sub>N</sub>	Q <sup>d</sup> <sub>P</sub>		
<i>endo</i> -Pd([9]aneNPN)Cl <sub>2</sub>	2.119	2.241	3.065		0.101	-0.480	-0.710		6.81	-0.22
<i>endo</i> -Pd([9]aneN <sub>2</sub> P)Cl <sub>2</sub>	2.094			3.162	0.299	-0.451		0.627	6.80	0.00
	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>a</sup> <sub>Pd-S</sub>	R <sup>b</sup> <sub>Pd...N</sub>	R <sup>b</sup> <sub>Pd...S</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>Cl</sub>	Q <sup>d</sup> <sub>N</sub>	Q <sup>d</sup> <sub>S</sub>		
<i>endo</i> -Pd([9]aneNSN)Cl <sub>2</sub>	2.102	2.348	2.935		0.177	-0.450	-0.702		6.48	-0.16
<i>endo</i> -Pd([9]aneN <sub>2</sub> S)Cl <sub>2</sub>	2.097			3.238	0.309	-0.458		0.269	6.68	0.00

<sup>a</sup>Bond length between the Pd<sup>II</sup> center and equatorial binding atom of the tridentate ligand. <sup>b</sup>Bond length between the Pd<sup>II</sup> center and axial binding site of the tridentate ligand. <sup>c</sup>Atomic charges of the Pd<sup>II</sup> center and the binding atom of *trans* L-ligand. <sup>d</sup>Atomic charge of an apical binding atom. <sup>e</sup>Relative energy gap between HOMO and LUMO. <sup>f</sup>Energy gap between Pd([9]aneB<sub>2</sub>A)L<sub>2</sub> and Pd([9]aneBAB)L<sub>2</sub> types. <sup>g</sup>Energy gap between *exo*- and *endo*-type structures.

**Figure S3.** The  $3a_{1g}(5s)$  and  $2a_{1g}$  orbitals of *endo*-Pd([9]aneB<sub>2</sub>A)L<sub>2</sub> and *endo*-Pd([9]aneBAB)L<sub>2</sub> (A=P, S; B=N; L= PH<sub>3</sub>, Cl<sup>-</sup>) calculated at the B3P86/6-311+G(d,p) (3-21G(d) for Pd) level.



**Table S3.** Optimized average bond distances (Å), average atomic charges (NBO, au), and relative energies (eV) of the equilibrium structures {Pd([9]aneB<sub>2</sub>A)L<sub>2</sub>} and [Pd([9]aneBAB)L<sub>2</sub>] at the B3P86//6-311+G(d,p) (3-21G(d) for Pd) levels.

Compound	Average distance				Average atomic charge				Relative energy	
	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>a</sup> <sub>Pd-P</sub>	R <sup>b</sup> <sub>Pd...N</sub>	R <sup>b</sup> <sub>Pd... P</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>PH<sub>3</sub></sub>	Q <sup>d</sup> <sub>N</sub>	Q <sup>d</sup> <sub>P</sub>	ΔE <sup>e</sup> <sub>H-L</sub>	ΔE <sup>f</sup> <sub>BAB- B<sub>2</sub>A</sub>
<i>endo</i> - [Pd([9]aneNPN)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.216	2.292	2.762		0.003	0.207	-0.68 7		3.46	0.29
<i>endo</i> - [Pd([9]aneN <sub>2</sub> P)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.153			2.789	0.099	0.249		0.576	2.76	0.00
	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>a</sup> <sub>Pd-S</sub>	R <sup>b</sup> <sub>Pd...N</sub>	R <sup>b</sup> <sub>Pd... S</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>PH<sub>3</sub></sub>	Q <sup>d</sup> <sub>N</sub>	Q <sup>d</sup> <sub>S</sub>		
<i>endo</i> - [Pd([9]aneNSN)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.162	2.376	2.748		0.048	0.242	-0.68 6		3.16	0.12
<i>endo</i> - [Pd([9]aneN <sub>2</sub> S)(PH <sub>3</sub> ) <sub>2</sub> ] <sup>2+</sup>	2.148			2.867	0.129	0.244		0.307	2.73	0.00
	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>a</sup> <sub>Pd-P</sub>	R <sup>b</sup> <sub>Pd...N</sub>	R <sup>b</sup> <sub>Pd... P</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>Cl</sub>	Q <sup>d</sup> <sub>N</sub>	Q <sup>d</sup> <sub>P</sub>		
<i>endo</i> -Pd([9]aneNPN)Cl <sub>2</sub>	2.132	2.240	2.932		0.646	-0.61 6	-0.65 7		3.79	-0.18
<i>endo</i> -Pd([9]aneN <sub>2</sub> P)Cl <sub>2</sub>	2.107			2.959	0.341	-0.45 4		0.597	3.54	0.00
	R <sup>a</sup> <sub>Pd-N</sub>	R <sup>a</sup> <sub>Pd-S</sub>	R <sup>b</sup> <sub>Pd...N</sub>	R <sup>b</sup> <sub>Pd... S</sub>	Q <sup>c</sup> <sub>Pd</sub>	Q <sup>c</sup> <sub>Cl</sub>	Q <sup>d</sup> <sub>N</sub>	Q <sup>d</sup> <sub>S</sub>		
<i>endo</i> -Pd([9]aneNSN)Cl <sub>2</sub>	2.110	2.357	2.786		0.254	-0.45 9	-0.70 1		3.35	-0.14
<i>endo</i> -Pd([9]aneN <sub>2</sub> S)Cl <sub>2</sub>	2.111			3.103	0.752	-0.58 3		0.225	3.49	0.00

<sup>a</sup>Bond length between the Pd<sup>II</sup> center and equatorial binding atom of the tridentate ligand. <sup>b</sup>Bond length between the Pd<sup>II</sup> center and axial binding site of the tridentate ligand. <sup>c</sup>Atomic charges of the Pd<sup>II</sup> center and an binding atom of *trans* L-ligand. <sup>d</sup>Atomic charge of an apical binding atom. <sup>e</sup>Relative energy gap between HOMO and LUMO. <sup>f</sup>Energy gap between Pd([9]aneB<sub>2</sub>A)L<sub>2</sub> and Pd([9]aneBAB)L<sub>2</sub> types. <sup>g</sup>Energy gap between *exo*- and *endo*-type structures.