

Supplementary Information for:

DFT Calculations of ^{31}P NMR Chemical Shifts in Palladium Complexes

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CALCULATIONAL DETAILS

The quantum chemical calculations have been carried out within the framework of the Kohn Sham density functional theory [1] with the Gaussian 03 [2] (Revision B.04) and the Gaussian 16 [3] (Revision A.03) software packages by using PBE0 [4] functional and a number of families of basis sets (Pople's [5-12], Ahlrichs's [13-15], Dunning's [16-17], Jensen's [18-20]). For the Pd center was used the quasi-relativistic Stuttgart-Dresden ECP28MWB with corresponding (8*s*7*p*6*d*)/[6*s*5*p*3*d*] GTO valence basis set [21] (denoted as "(SDD)"), Sapporo-ro-DKH3-DZP-diffuse [deprecated] basis set [22-23] and def2-TZVPD basis set [24-25]. Wherever possible, geometry optimization was started from X-ray structure. For most of the complexes the calculations were carried out for all possible conformers/isomers and results for the lowest energy forms were used in analysis. To take into account the medium effects, calculations were also carried out in the framework of Polarizable Continuum Model [26] (denoted as "PCM") with chloroform as solvent. To determine the energies of triplet states geometry optimizations were performed at the spin-unrestricted formalism (UPBE0/6-31+G(d)). ³¹P NMR CSs were calculated by the GIAO method [27]. All ³¹P data was referenced to H₃PO₄.

The pc-2, def2-TZVP, def2-TZVPD and Sapporo-DKH3-DZP-diffuse basis sets were downloaded from the EMSL basis set library for Gaussian package [28].

The Gaussian 03 calculations were carried out on a PC with IntelCore i7-3970X CPU, 3.5 GHz. Calculations with 6-311+G(3df), pc-2, cc-pVTZ, def2-TZVP, and Sapporo-ro-DKH3-DZP-diffuse [deprecated] basis sets were carried out with the Gaussian 16 on 20 CPUs, IntelXeon ES-2650 2.20 GHz.

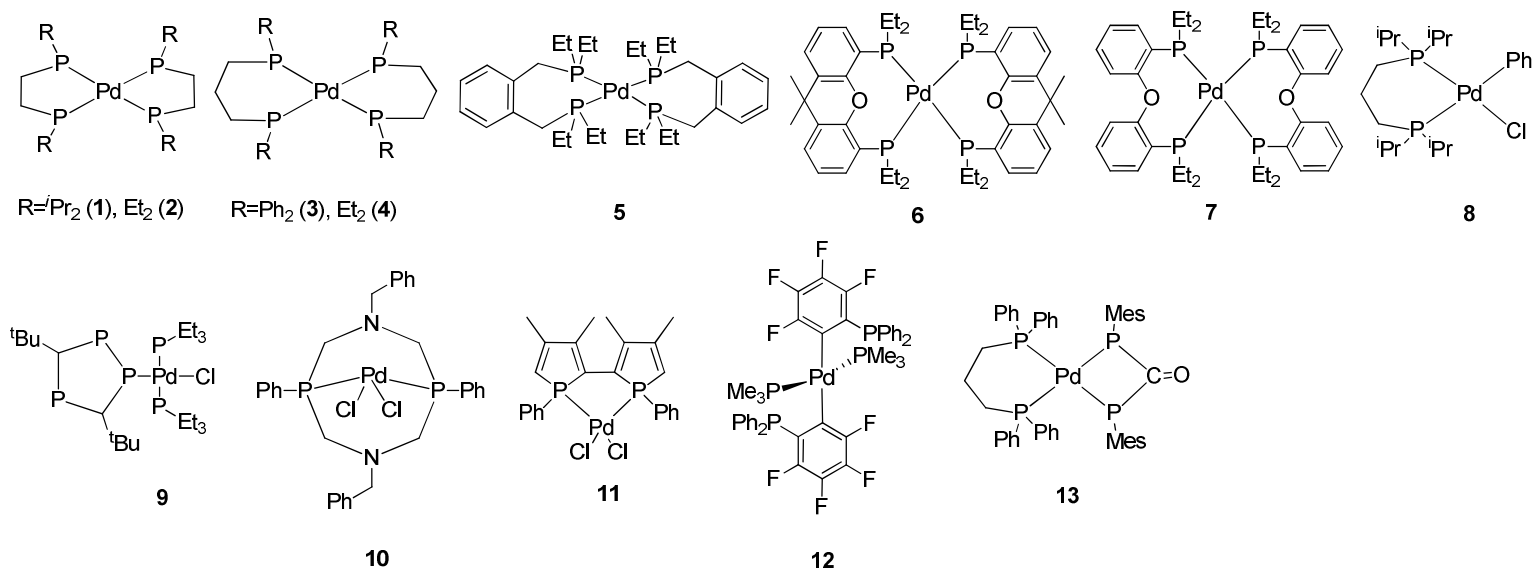


Figure S1. Complexes based on σ -donor ligands.

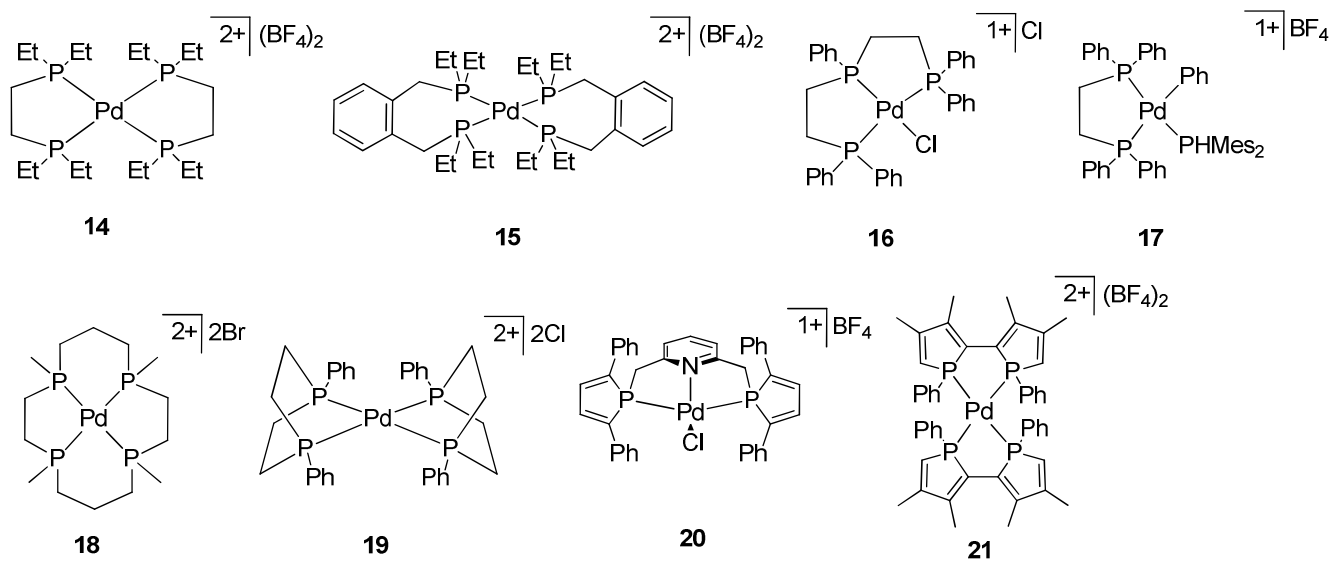


Figure S2. Charged complexes based on σ -donor ligands.

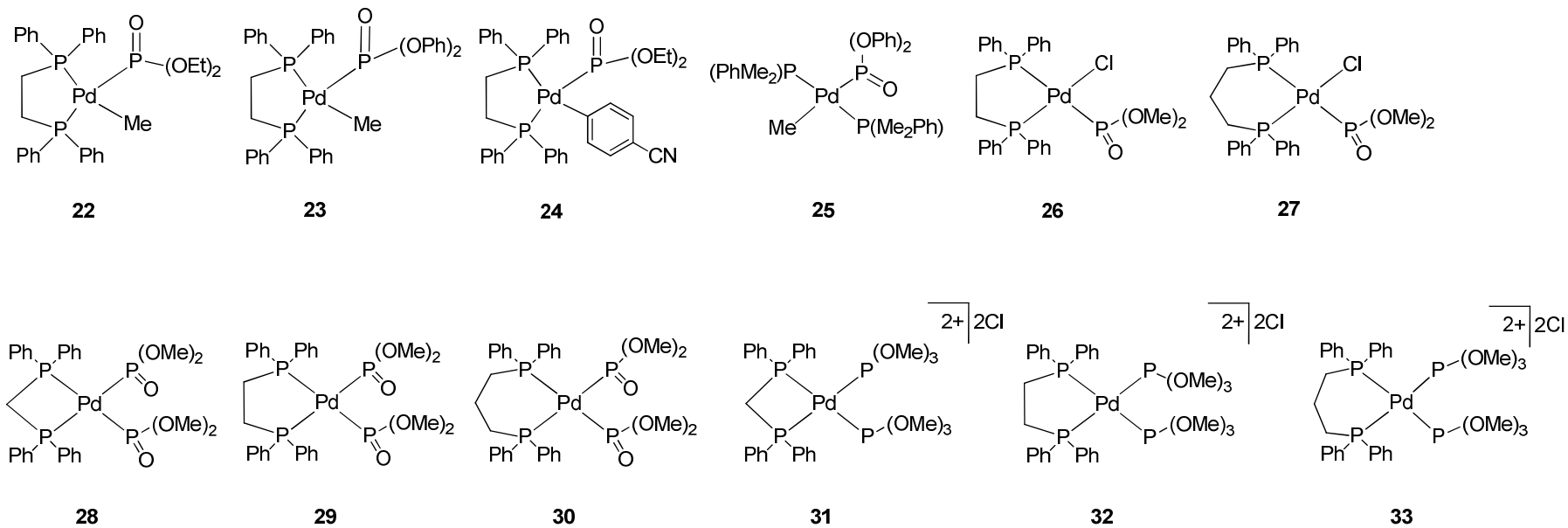


Figure S3. Complexes containing P=O groups.

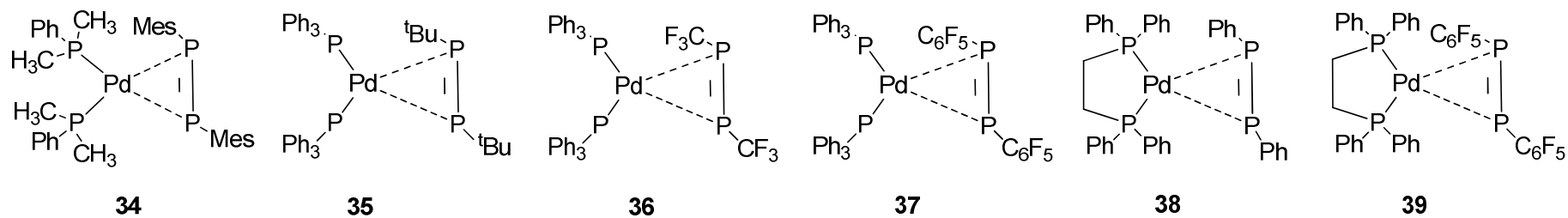


Figure S4. Complexes based on π -donating ligands.

Table S1. Experimental and calculated ³¹P NMR shifts (ppm) for all model complexes.

		Calculated								
Compound	Experiment	PBE0/6-311G(2d,2p)// PBE0/6-31+G(d)	PBE0/6-311G(2d,2p)// PBE0/6-31+G(d) ^a	PBE0/6-311G(2d,2p) (PCM)// PBE0/6-31+G(d) (PCM)	PBE0/6-311G(2d,2p) (PCM)// PBE0/6-31+G(d) (PCM) ^a	PBE0/6-311G(2d,2p)// PBE0/6-311+G(2d)	PBE0/6-311G(2d,2p)// PBE0/6-311+G(2d) ^a	PBE0/def2-TZVP// PBE0/6-311+G(2d)	PBE0/ def2-TZVP // PBE0/6-311+G(2d) ^a	Reference
1	45.5	22.5	29.6	29.7	33.9	27.1	36.1	43.7	38.1	29
2	31.5	15.0	23.2	14.2	20.8	12.0	22.8	28.1	24.2	30
3	3.7	-2.0	8.8	-3.0	6.4	-0.1	12.2	12.3	10.1	29
4	-2.6	-14.0	-1.4	-14.2	-3.1	-13.3	0.5	-2.2	-2.8	30
5	1.3	-17.6	-4.5	-14.4	-3.2	-17.5	-3.2	-1.4	-2.1	30
	-0.4	-18.1	-4.9	-21.5	-9.2	-24.5	-9.3	-9.0	-8.8	
6	-0.9	-19.1	-5.8	-18.6	-6.8	-21.9	-7.1	-5.6	-5.8	30
	-9.9	-27.6	-13.0	-27.3	-14.1	-30.7	-14.8	-13.1	-12.5	
7	-1.3	-20.1	-6.6	-19.1	-7.2	-23.3	-8.3	-8.4	-8.3	30
8	11.2	-6.3	5.1	-3.2	6.2	-4.4	8.4	6.1	4.6	29
	29.0	23.1	30.1	30.9	34.9	25.5	34.7	36.0	31.2	
9	123.0	121.3	113.5	143.0	129.2	120.2	118.2	133.9	118.4	31
	18.5	17.0	24.9	18.1	24.1	13.8	24.4	30.0	25.9	
10	-9.2	-9.4	2.5	-7.5	2.6	-6.9	6.2	3.7	2.5	32
11	33.4	30.7	36.5	36.9	39.9	29.1	37.9	42.0	36.6	33
12	-2.9	-17.1	-4.1	-18.0	-6.3	-17.4	-3.1	-8.4	-8.3	34
13	3.5	-17.7	-4.6	-14.9	-3.7	-20.9	-6.2	2.0	1.0	35
	172.3	183.3	166.2	166.7	149.2	180.7	171.5	200.0	177.2	
14	65.2	60.6	62.0	60.2	59.6	56.8	62.3	72.1	63.3	30
15	19.0	15.2	23.4	14.9	21.4	15.0	25.5	26.5	22.8	30
16	44.2	34.1	39.4	39.6	42.2	31.3	39.8	47.8	41.7	36
	109.3	99.2	94.8	109.5	101.1	96.3	97.1	114.0	100.6	

17	50.6	38.0	42.7	48.1	49.4	44.1	51.1	61.8	54.2	37
	44.6	26.4	32.9	30.1	34.2	26.1	35.3	41.4	36.0	
	-49.9	-58.2	-39.0	-60.2	-41.8	-61.2	-41.7	-48.6	-44.1	
18	38.5	33.6	39.0	33.5	37.1	29.5	38.2	43.8	38.2	38
19	55.6	46.2	49.7	47.8	49.1	42.8	50.0	58.2	51.0	39
20	33.2	25.1	31.8	35.5	38.8	34.6	42.7	47.0	41.0	40
21	36.6	29.4	35.4	28.1	32.5	16.6	26.9	40.0	34.8	33
22	83.1	95.2	91.4	103.2	95.8	84.5	86.7	96.0	84.6	41
	36.5	14.9	23.1	20.0	25.7	13.4	24.1	31.2	26.9	
	42.6	37.0	41.9	41.5	43.8	34.2	42.4	49.9	43.6	
23	92.9	101.9	97.1	116.4	106.9	95.8	96.7	107.0	94.4	42
	48.3	44.6	48.4	49.0	50.1	40.9	48.3	56.4	49.4	
	40.1	29.8	35.8	30.4	34.5	24.3	33.7	40.1	34.9	
24	74.7	106.5	101.0	99.6	92.7	83.4	85.8	94.5	83.3	41
	38.0	23.9	30.8	30.6	34.6	23.7	33.1	37.1	32.2	
	39.0	36.3	41.3	36.0	39.2	20.7	30.5	40.8	35.5	
25	99.5	111.5	105.2	115.4	106.0	100.1	100.5	110.1	97.2	42
	-0.2	-1.3	9.3	-0.9	8.1	-4.8	8.0	10.6	8.6	
26	39.2	13.6	22.0	24.1	29.2	9.5	20.6	24.9	21.3	43
	57.3	50.9	53.7	62.7	61.7	48.8	55.3	65.3	57.3	
	67.7	87.3	84.6	91.5	85.9	77.4	80.5	88.2	77.7	
27	-5.1	-19.9	-6.5	-12.2	-1.4	-22.7	-7.8	-7.1	-7.1	43
	16.7	16.7	24.6	23.5	28.7	14.4	24.9	30.6	26.4	
	64.8	90.4	87.3	95.5	89.3	81.3	83.9	91.8	80.9	
28	-29.7	-47.6	-30.0	-46.8	-30.5	-48.8	-30.8	-32.5	-29.8	43
	75.0	94.2	90.5	99.1	92.3	84.3	86.5	94.2	83.0	
29	43.1	22.9	29.9	38.6	41.4	25.7	34.9	43.2	37.6	43
	73.9	85.2	82.9	96.5	90.1	80.3	83.0	90.3	79.5	
30	0.5	-8.9	2.9	-4.7	4.9	-11.0	2.6	4.6	3.3	43
	70.5	89.7	86.7	94.9	88.8	80.6	83.3	91.2	80.3	
31	-32.4	-58.7	-39.4	-56.2	-38.4	-60.4	-41.0	-45.8	-41.6	43
	104.5	114.7	107.9	115.8	106.4	106.6	106.2	117.5	103.8	
32	54.9	66.9	67.3	64.9	63.5	62.7	67.5	78.4	69.0	43
	103.0	112.7	106.2	117.1	107.4	108.2	107.6	118.8	104.9	
33	4.2	10.7	19.5	11.5	18.6	8.1	19.4	24.1	20.6	43
	100.3	113.0	106.5	113.7	104.6	105.2	105.0	116.1	102.5	
34	49.7	53.1	55.6	38.2	41.0	48.3	54.8	62.2	54.5	44

	-12.8	-26.1	-11.7	-23.6	-11.0	-28.7	-13.0	-13.6	-12.9	
35	123.7	121.0	113.3	106.8	98.8	115.8	114.3	128.4	113.5	45
	21.4	9.0	18.1	9.8	17.1	7.3	18.7	22.9	19.6	
36	50.5	49.0	52.1	35.6	38.9	45.1	52.0	57.4	50.3	46
	22.7	12.4	21.0	13.6	20.3	10.7	21.7	25.1	21.5	
37	32.0	22.1	29.2	9.3	16.7	17.3	27.5	30.7	26.5	47
	24.8	15.2	23.4	15.9	22.3	13.7	24.3	29.3	25.3	
38	34.3	20.5	27.9	31.5	35.4	20.4	30.2	34.6	30.0	47
	41.0	25.9	32.5	27.3	31.9	23.1	32.6	39.2	34.1	
39	1.7	-4.0	7.1	-24.5	-11.7	1.5	13.6	14.4	12.0	47
	52.1	33.4	38.8	36.6	39.7	32.1	40.5	47.9	41.8	
R^2		0.9535	0.9535	0.9447	0.9447	0.9627	0.9627	0.9720	0.9720	
$RMSE$		13.8	8.9	14.2	9.8	13.6	8.0	10.8	6.9	

^a corrected by using the factors for empirical linear scaling obtained by the linear regression procedure for the title Pd-complexes.

Table S2. Empirical scaling factors obtained by the linear regression analysis of calculated and experimental $\delta^{31}\text{P}$ for the title Pd-complexes.

Level of theory	R^2	Slope	Intercept	$RMSE$
PBE0/6-311G(2d,2p)//PBE0/6-31+G(d)	0.954	1.1766	-12.297	8.9
PBE0/6-311G(2d,2p) (PCM)//PBE0/6-31+G(d) (PCM)	0.945	1.1881	-10.561	9.8
PBE0/6-311G(2d,2p)//PBE0/6-311+G(2d)	0.963	1.1346	-13.8962	8.0
PBE0/def2-TZVP//PBE0/6-311+G(2d)	0.972	0.9297	1.1235	6.9

Table S3. Triplet-singlet states energy gap (UPBE0/6-31+G(d)) for the "training" set of Pd complexes (**2**, **9**, **17**, **22-23**, **26**, **28-30**, **32**, **38**).

Compound	E, kcal/mol
38	38.9
9	38.9
17	46.6
2	51.7
22	46.1
23	44.9
26	40.9
28	- ^a
29	35.8
30	35.6
32	67.6

^a triplet state does not correspond to energy minima.

Table S4. Experimental and calculated (by different combinations) ³¹P NMR shifts (ppm) for the "training" set of Pd complexes (**2**, **9**, **17**, **22-23**, **26**, **28-30**, **32**, **38**).

		Calculated																					
		Combination																					
Compound	Experimental	6-311G(2d,2p)// 6-31+G(d)	6-311G(2d,2p)// 6-31+G(2d)	6-311G(2d,2p)// 6-311+G(d)	6-311G(2d,2p)// 6-311+G(2d)	6-311+G(2d,2p)// 6-311+G(2d)	6-311G(2d,2p)// 6-311+G(3df)	6-311G(2d,2p)//6-311+G(2d) Pd - def2-TZVPPD//def2-TZVPPD	6-311G(2d,2p)//6-311+G(2d) Pd - SDD//DKH3-DZP	6-311G(2d,2p)//6-311+G(2d) Pd - DKH3-DZP//DKH3-DZP	cc-pVTZ// 6-311+G(3df)	pc-2// 6-311+G(3df)	def2-TZVP// 6-311+G(3df)	def2-TZVP// 6-311+G(2d)	TZV// 6-311+G(3df)	def2-TZVP// def2-TZVP	6-311G(2d,2p)// cc-pVDZ	6-311G(2d,2p)// cc-pVTZ	6-311G(2d,2p)// pc-2	6-311G(2d,2p)// def2-TZVP	TZV// TZV	cc-pVTZ// 6-31+G(d)	cc-pVTZ// cc-pVDZ
2	31.5	15.0	22.3	25.8	12.0	21.8	25.3	14.8	1.3	-4.0	13.7	27.2	31.3	28.1	-37.8	30.3	12.4	22.2	25.0	24.2	-58.2	3.2	-3.9
9	123.0	121.3	129.4	133.5	120.2	131.8	132.0	124.6	111.7	112.3	123.1	131.9	135.2	133.9	75.4	133.8	119.2	128.9	138.0	130.5	68.9	112.5	109.0
	18.5	17.0	24.3	27.8	13.8	24.0	25.8	17.6	22.8	21.7	7.4	33.4	31.9	30.0	-46.8	31.0	14.4	23.5	25.4	24.9	-66.4	-1.5	-6.9
17	50.6	38.0	53.4	57.8	44.1	56.2	56.4	47.9	49.1	47.2	37.3	64.2	64.2	61.8	-8.9	62.9	44.7	54.2	55.8	55.2	-32.8	28.5	22.6
	44.6	26.4	35.4	39.8	26.1	36.7	38.4	29.3	28.2	27.9	27.5	41.3	43.7	41.4	-33.9	42.6	25.7	35.8	37.7	37.3	-58.0	17.6	10.8
	-49.9	-58.2	-52.0	-47.3	-61.2	-51.7	-49.0	-57.6	-62.8	-68.3	-52.2	-44.7	-46.6	-48.6	-122.8	-48.6	-63.6	-53.6	-50.1	-51.0	-150.8	-64.2	-71.3
38	34.3	20.5	31.5	34.4	20.4	33.0	24.8	25.2	49.0	92.4	23.0	28.0	28.5	34.6	-33.4	28.7	22.6	25.9	27.0	24.6	52.4	11.4	15.7
	41.0	25.9	32.0	36.6	23.1	34.8	35.9	26.3	17.8	13.4	26.0	39.1	41.9	39.2	-0.8	40.9	23.3	33.5	35.2	34.6	-55.4	38.7	8.5
22	83.1 ^a	95.2	96.7	104.4	84.5	94.4	93.7	88.9	90.0	93.2	85.4	92.2	95.0	96.0	76.7	94.6	104.7	95.9	94.3	93.4	119.8	85.6	89.1
	36.5	14.9	23.0	26.2	13.4	25.6	26.3	16.8	12.0	9.5	13.4	28.1	34.1	31.2	-39.5	32.8	12.3	23.3	25.3	24.9	-40.2	2.5	-3.9
	42.6	37.0	43.6	48.4	34.2	46.0	46.6	37.9	34.3	32.0	34.3	49.4	52.3	49.9	-19.9	51.3	36.1	44.5	46.0	45.5	-66.0	25.1	19.9
23	92.9	101.9	106.3	117.4	95.8	106.9	104.5	99.5	100.6	105.6	96.8	120.5	105.9	107.0	94.7	104.4	119.4	106.6	100.9	102.9	150.3	98.1	104.5
	48.3	44.6	50.6	55.2	40.9	52.9	53.2	44.4	38.5	21.9	41.9	72.9	58.7	56.4	-16.5	58.3	42.1	51.3	50.0	52.3	-37.7	33.3	29.1
	40.1	29.8	32.4	38.0	24.3	34.1	36.4	27.6	23.0	35.6	24.6	55.6	42.1	40.1	-32.7	40.6	23.4	33.7	32.5	34.7	-52.6	14.6	6.4
26	39.2	13.6	20.2	23.6	9.5	21.3	21.1	13.1	11.8	12.2	10.8	21.5	26.4	24.9	-45.8	26.3	13.4	20.7	21.7	20.6	-53.9	1.4	-5.0
	57.3	50.9	58.2	63.0	48.8	59.9	60.9	53.1	51.0	47.1	47.2	62.3	67.3	65.3	-14.4	66.3	47.8	58.4	60.2	59.8	-36.3	38.2	31.3
	67.7	87.3	89.6	95.6	77.4	87.0	86.1	81.4	83.0	85.4	78.1	85.5	87.1	88.2	77.1	87.4	96.2	89.1	87.5	86.4	115.0	79.1	82.6
28	-29.7	-47.6	-38.5	-35.9	-48.8	-36.6	-36.7	-45.1	-41.6	-50.4	-45.8	-34.2	-30.7	-32.5	-101.8	-31.6	-54.3	-40.2	-37.7	-37.9	-133.1	-55.1	-64.8
	75.0	94.2	95.3	103.2	84.3	93.5	93.2	88.7	88.5	90.6	84.9	91.9	93.7	94.2	80.8	93.6	106.0	95.5	93.9	93.0	117.4	85.2	92.1
29	43.1	22.9	35.3	39.4	25.7	38.4	37.9	29.0	20.5	11.5	21.8	43.0	45.6	43.2	-25.1	39.8	25.3	34.7	36.4	34.1	-48.7	15.6	8.1

	73.9	85.2	91.8	99.0	80.3	89.8	89.4	84.5	85.9	87.5	81.1	87.5	89.6	90.3	75.6	90.3	103.9	91.5	90.1	89.9	120.6	80.8	87.8
30	0.5	-8.9	-1.9	2.3	-11.0	-0.1	0.4	-7.8	-7.6	-11.5	-13.6	5.6	6.5	4.6	-67.3	5.5	-12.1	-2.3	0.0	-0.8	-94.8	-22.8	-30.0
	70.5	89.7	92.1	99.8	80.6	91.8	89.9	85.2	86.8	87.1	81.5	88.7	90.3	91.2	80.6	90.3	98.9	92.2	90.7	89.9	120.3	81.1	84.4
32	54.9	66.9	72.1	77.1	62.7	72.9	73.9	66.5	74.0	78.3	58.8	77.7	79.6	78.4	8.3	78.7	65.6	73.0	73.9	73.0	-4.7	51.8	47.0
	103.0	112.7	119.1	124.5	108.2	118.9	116.4	112.1	115.0	116.3	103.9	113.6	117.0	118.8	89.4	116.5	124.0	117.7	116.6	115.9	112.4	102.7	105.4
	R^2	0.9359	0.9512	0.9439	0.9547	0.9593	0.9586	0.9530	0.9042	0.8287	0.9498	0.9428	0.9654	0.9644	0.8333	0.9638	0.9113	0.9529	0.9582	0.9564	0.6889	0.9271	0.9011
	$\Delta\delta^b$	24	19	22	18	18	14	19	20	28 (65) ^c	15	19	9	13	68	11	36	17	11	15	118 (101) ^d	22	37

^a ³¹P NMR shifts of P=O phosphorus atoms colored in red; ^b average deviation (ppm) from the general correlation line for P=O phosphorus atoms; ^c in brackets average deviation from the general correlation line for P=P phosphorus atoms.

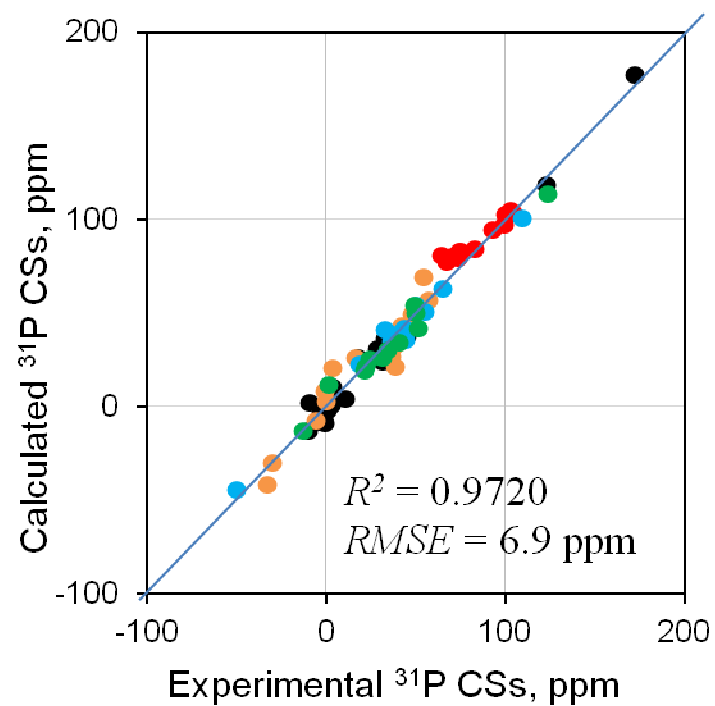


Figure S5. Correlation of calculated (PBE0/{def2-TZVP; Pd(SDD)}/{PBE0/{6-311+G(2d); Pd(SDD)}}, scaled) vs experimental ^{31}P CSs for palladium complexes **1-39**. Complexes based on σ -donor ligands (●), charged palladium complexes based on σ -donor ligands (●), complexes containing P=O groups (● are the P = O phosphorus, ● are other phosphorus atoms in this type of complexes) and complexes based on π -donor ligands (●).

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