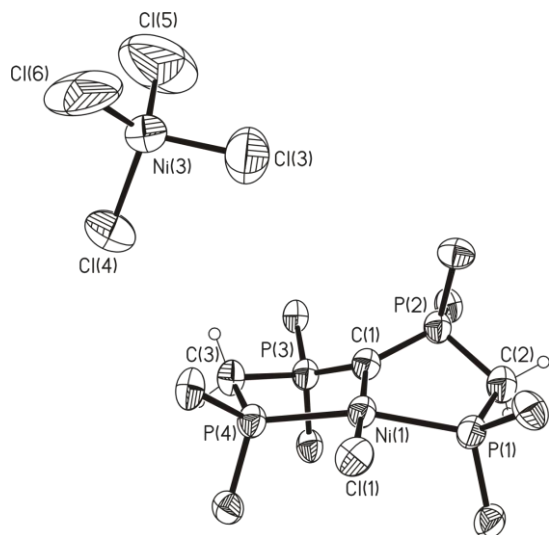
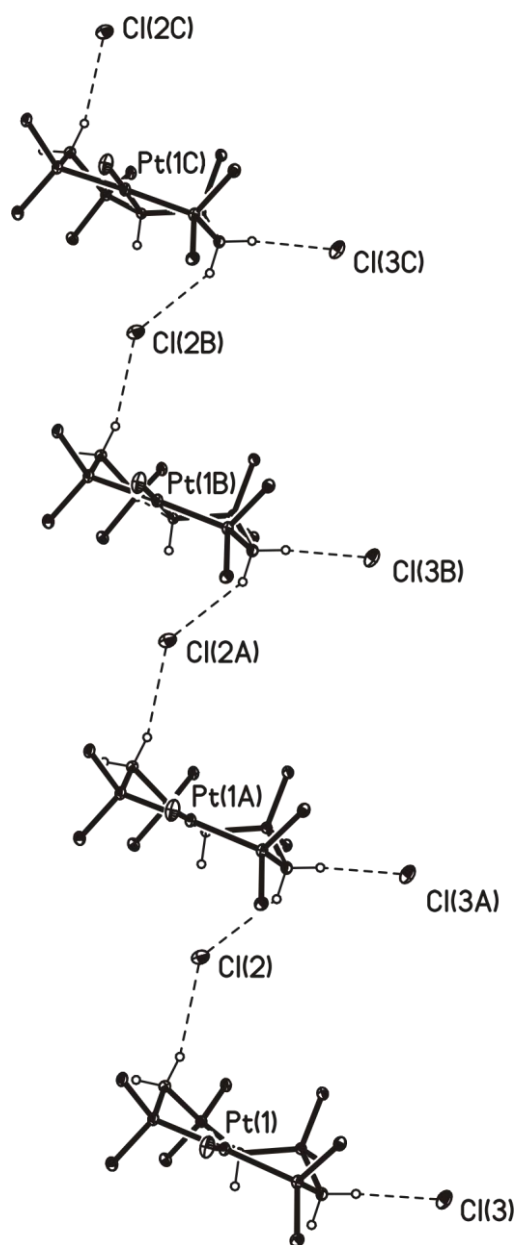


## Supporting Information

**Thermal ellipsoid plot of 1b.** Only the *ipso* carbon atoms of the phenyl groups are shown.



**Example for Supramolecular Structure: Compound  $7 \cdot 2\text{CHCl}_3$**



**Table 3. Summary of X-ray Data and structure refinement for 1a, 1b, 2 and 3**

	<b>1a</b> ·2C <sub>4</sub> H <sub>8</sub> O <sub>2</sub>	<b>1b</b> ·CH <sub>2</sub> Cl <sub>2</sub>	<b>2</b> ·2.5C <sub>7</sub> H <sub>8</sub> ·CH <sub>3</sub> OH	<b>3</b> ·0.5H <sub>2</sub> O·2.25C <sub>3</sub> H <sub>6</sub> O
formula	C <sub>51</sub> H <sub>44</sub> Cl <sub>2</sub> NiP <sub>4</sub> • 2 C <sub>4</sub> H <sub>8</sub> O <sub>2</sub>	C <sub>102</sub> H <sub>88</sub> Cl <sub>6</sub> Ni <sub>3</sub> P <sub>8</sub> • CH <sub>2</sub> Cl <sub>2</sub>	C <sub>51</sub> H <sub>44</sub> Cl <sub>2</sub> P <sub>4</sub> Pd • 2.5 C <sub>7</sub> H <sub>8</sub> • CH <sub>3</sub> OH	C <sub>51</sub> H <sub>44</sub> Cl <sub>2</sub> P <sub>4</sub> Pt • 0.5 H <sub>2</sub> O • 2.25 C <sub>3</sub> H <sub>6</sub> O
<i>M<sub>r</sub></i>	1086.56	2035.24	1220.42	1186.42
T (K)	233(2)	233(2)	233(2)	233(2)
cryst syst	Orthorhombic	Trigonal	Triclinic	Triclinic
space group	Iba2 (no.45)	P3 <sub>1</sub> (no.144)	P-1 (no.2)	P-1 (no.2)
a (Å)	9.6018(2)	17.4335(2)	10.2001(2)	9.7629(2)
b (Å)	23.6046(4)	17.4335(2)	12.4004(2)	23.2214(3)
c (Å)	24.3824(4)	31.0829(4)	24.4161(4)	24.4318(4)
α (deg)	90	90	84.068(1)	96.059(1)
β (deg)	90	90	82.649(1)	94.714(1)
γ (deg)	90	120	89.914(1)	92.421(1)
V (Å <sup>3</sup> )	5526.19(17)	8181.28(17)	3046.28(9)	5482.07(16)
Z	4	3	2	4
<i>D</i> <sub>calcd</sub> (Mg/m <sup>3</sup> )	1.306	1.239	1.331	1.437
μ(mm <sup>-1</sup> )	0.609	0.867	0.540	2.815
F(000)	2272	3144	1266	2396
indep rflns	4807	13390	12411	19002
data / restraints / parameters	4807 / 1 / 319	13390 / 15 / 1129	12411 / 24 / 702	19002 / 48 / 1202
R1 [I>2σ(I)]	0.0316	0.0630	0.0411	0.0405
wR2 (all data)	0.0727	0.1792	0.1098	0.1034
Largest diff. peak and hole (e.Å <sup>-3</sup> )	0.253 and -0.213	1.159 and - 0.955	0.619 and -0.726	0.930 and -0.822

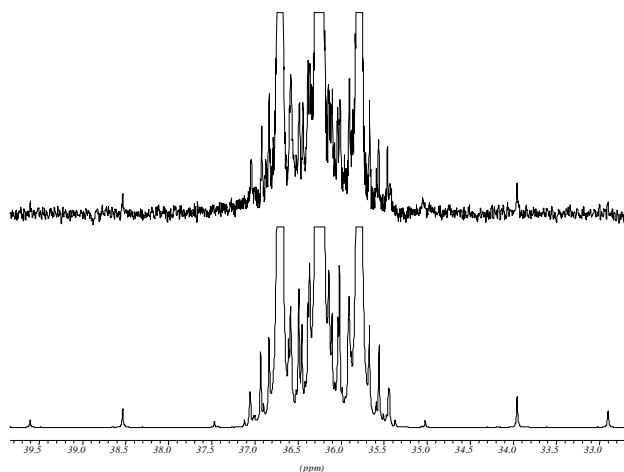
**Table 4. Summary of X-ray Data and structure refinement for 4, 5, 6 and 7**

	<b>4</b> ·0.5C <sub>7</sub> H <sub>8</sub> ·1.5CHCl <sub>3</sub>	<b>5</b>	<b>6</b> ·3H <sub>2</sub> O	<b>7</b> ·2CHCl <sub>3</sub>
formula	C <sub>51</sub> H <sub>46</sub> C <sub>12</sub> P <sub>4</sub> · 0.5 C <sub>7</sub> H <sub>8</sub> · 1.5 CHCl <sub>3</sub>	C <sub>51</sub> H <sub>45</sub> ClP <sub>4</sub>	C <sub>51</sub> H <sub>45</sub> Cl <sub>3</sub> P <sub>4</sub> Pd · 3 H <sub>2</sub> O	C <sub>51</sub> H <sub>45</sub> Cl <sub>3</sub> P <sub>4</sub> Pt · 2 CHCl <sub>3</sub>
<i>M<sub>r</sub></i>	1078.78	817.20	1048.55	1321.93
T (K)	233(2)	233(2)	233(2)	233(2)
cryst syst	Orthorhombic	Monoclinic	Triclinic	Triclinic
space group	P2 <sub>1</sub> 2 <sub>1</sub> 2 (no.18)	P2 <sub>1</sub> /c (no.14)	P-1 (no.2)	P-1 (no.2)
a (Å)	15.2377(6)	16.9190(4)	9.2897(3)	9.4233(2)
b (Å)	20.5124(7)	27.1997(6)	13.5853(6)	15.0455(3)
c (Å)	8.7621(4)	18.7995(2)	19.9199(7)	19.0383(4)
α (deg)	90	90	86.390(2)	90.021(2)
β (deg)	90	92.034(2)	89.649(2)	90.644(2)
γ (deg)	90	90	72.822(2)	96.536(2)
V (Å <sup>3</sup> )	2738.70(19)	8645.9(3)	2396.84(16)	2681.50(10)
Z	2	8	2	2
<i>D</i> <sub>calcd</sub> (Mg/m <sup>3</sup> )	1.308	1.256	1.453	1.637
μ (mm <sup>-1</sup> )	0.491	0.271	0.730	3.221
F(000)	1116	3424	1076	1312
indep rflns	4284	13577	8354	9424
data / restraints / parameters	4284 / 13 / 323	13577 / 0 / 1017	8354 / 4 / 579	9424 / 13 / 636
R1 [I>2σ(I)]	0.0498	0.0477	0.0420	0.0367
wR2 (all data)	0.1283	0.1082	0.0813	0.0711
Largest diff. peak and hole	0.594 and -0.267 e.Å <sup>-3</sup>	0.283 and -0.290 e.Å <sup>-3</sup>	0.587 and -0.593 e.Å <sup>-3</sup>	0.680 and -0.780 e.Å <sup>-3</sup>

## Detailed description of WIN-DAISY simulation/iteration of NMR spectra

**<sup>31</sup>P-NMR:** Simulation and iteration of the [AX]<sub>2</sub> system for **5** and **4** was performed by keeping the P1-P4 coupling at 0 Hz. The [AX]<sub>2</sub> system of **1a** and **2** was simulated by manual fitting of <sup>2</sup>J(P1P4) and <sup>2</sup>J(P2P3) to the very weak but detectable outer lines of the respective A and X subspectra (Figure 9), both couplings were kept constant during iteration of <sup>2+3</sup>J(P1P2) and <sup>3</sup>J(P1P3).

**Figure 9.** <sup>31</sup>P NMR phosphorane region (A-part of the [AX]<sub>2</sub> <sup>31</sup>P coupling pattern) of [Ni(Cl)(C(dppm)<sub>2</sub>-κ<sup>3</sup>P,C,P)]Cl (**1a**): upper trace experimental spectrum, lower trace simulated spectrum showing the (skipped) signals of the intense inner lines (with <sup>13</sup>C satellites at the basis) and the two pairs of weak but detectable outer lines in the very left and right parts of the spectrum, respectively.



The [AX]<sub>2</sub> system of **3**, **6** and **7** was only partly accessible to simulation, since only the N-lines ( $|J(\text{AX}) + J(\text{AX}')|$ ) and one single central line of the A and X part, respectively, were resolved. In line with the NMR data for **1a** and **2**, <sup>2</sup>J(P1P4) was kept constant at 400 Hz (the spectrum is completely insensitive to variation of this coupling constant between 300 – 500 Hz) and <sup>2</sup>J(P2P3) was kept constant at 69 Hz (**3**), -3.7 Hz (**6**) or -1.6 Hz (**7**) (this coupling

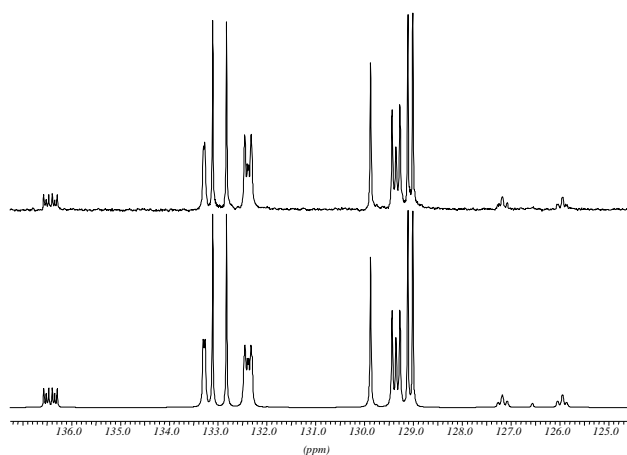
constant could be determined from the  $^{13}\text{C}$  NMR spectra).  $^{2+3}\text{J}(\text{P1P2})$  and  $^3\text{J}(\text{P1P3})$  were iterated for **3**, **6** and **7** but show a large uncertainty, due to the featureless triplet structure of the A and X signals. The Pt-P coupling constants were easily accessible from the  $^{195}\text{P}$  satellites in the  $^{31}\text{P}$  NMR spectrum of the Pt compounds **3** and **7**.

The P-C coupling constants extracted from the  $^{13}\text{C}$  NMR spectrum could be verified in the  $^{13}\text{C}$  satellites of the  $^{31}\text{P}$  signals for all compounds and were found appropriate; only peaks due to the coupling constants larger than ca. 30 Hz were sufficiently separated from the main signals for verification.

$^{13}\text{C}$ -NMR: The P-P coupling constants from the  $^{31}\text{P}$  spectra were used for simulation of the  $^{13}\text{C}$  NMR spectra, P-C coupling constants were fitted manually to the  $[\text{AX}]_2\text{Z}$  (central carbon atom, C1) and  $\text{AA}'\text{XX}'\text{Z}$  pattern (all other carbon atoms), respectively, of the  $^{13}\text{C}$  signals. Assignment of aromatic carbon resonances was performed by comparison with the literature data for  $\text{PPh}_3$ ,  $[\text{MePPh}_3]\text{I}$  and phosphine complexes.<sup>40b,42</sup> Moreover the assignment of the aromatic carbon signals to Ph1 and Ph2 was unambiguously possible due to the different P-C one bond couplings for  $C_i$  and in general by the significantly different coupling pattern to P1 and P2, respectively, which is largely dependent on the relative magnitudes of  $|\text{J}(\text{PC}) - \text{J}(\text{P}'\text{C})|$  and  $\text{J}(\text{PP}')$  towards the directly phenyl-bound phosphorus (see also Figure 10):<sup>40</sup> In the free ligands **4** and **5** the  $C_o/C_m$  signals at P1 show pure doublets, whereas pure triplets are observed for these carbon atoms in all metal complexes. The  $C_o/C_m$  signals at P2 show pure triplets in **1** - **3** “filled-in doublets” in **6** and **7** and higher order spectra with four to six resolved transitions for **4** and **5**. The  $C_i$  signals show a pattern related to  $C_o/C_m$  but with further splitting due to substantial  $^3\text{J}(\text{PC})$  coupling along the ligand backbone, the  $C_p$  signals are either singlets or show only partially resolved splitting. As already mentioned in the  $^{31}\text{P}$  NMR section the  $^2\text{J}(\text{P2P3})$  could be determined unambiguously from the  $^{13}\text{C}$  pattern of Ph2-

*Ci* for **3** (position of the outer lines of the quintett pattern) or all Ph1a-*Ci/o/m* and Ph1b-*Ci/o/m* signals (intensities of the inner lines of the “filled-in doublets”).

**Figure 10.**  $^{13}\text{C}$  NMR aromatic region of  $[\text{CH}(\text{dppm})_2]\text{Cl}$  (**5**): upper trace experimental spectrum, lower trace simulated spectrum showing different  $^{31}\text{P}$  coupling pattern of C atoms depending of relative magnitudes of  $|\text{J}(\text{PC}) - \text{J}(\text{P}'\text{C})|$  and  $\text{J}(\text{PP}')$ : Singlet (129.9 ppm), doublet (129.1, 133.0 ppm), filled-in doublet (129.4, 133.3 ppm), doublet of filled-in doublets (136.4 ppm), higher order pattern  $\geq 4$  lines (126.6, 132.4 ppm).



**$^1\text{H}$  NMR:** The P-P coupling constants from the  $^{31}\text{P}$  NMR spectra were used for simulation of the  $^1\text{H}$  NMR spectra, H-P coupling constants were fitted manually to the pattern of the  $^1\text{H}$  signal of H1 and H2. The H2 signals were not accessible for simulation of coupling constants in the free ligands due to an ambiguous pseudo-doublet pattern (**5**) or a featureless, broad appearance ( $W_{1/2} = 36$  Hz; **4**). Only the non-aromatic proton signals were simulated.