Palladium and rhodium allyl complexes in catalysis

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Chapter 7.
An X-ray study towards the effect of the bite angle of bidentate phosphine ligands on the geometry of palladium(allyl) complexes: implications for the regioselectivity in the allylic alkylation of a disubstituted allyl moiety.

In the previous chapters we have studied mono-substituted allyl moieties. The DFT calculations described in chapter 3 predict an enhanced non-symmetry of the Pd(η^3-allyl) bond for di-substituted allyl moieties. In this chapter we study the effect of the bite angle of bidentate phosphine ligands on the structure and reactivity of (P-P)palladium(3,3-(CH_3)₂-C,H₃) complexes.

Abstract
X-ray crystal structures of a series of cationic (P-P)palladium(3,3-(CH_3)₂-C,H₃) complexes (P-P = dppe (1,2-bis(diphenylphosphino)ethane), dpdf (1,1'-bis(diphenylphosphino)ferrocene), DPEphos) and the (Xantphos)Pd(C,H₅)BF₄ complex have been determined. In the solid state structure, the phenyl rings of the ligands are oriented in the direction of the non-symmetrically bound [3,3-(CH_3)₂-C,H₃] moiety. An increase of the bite angle of the chelating ligand results in an increase of the cone angle. In complexes containing ligands having a large cone angle, the distances between the phenyl rings and the allyl moiety become small, resulting in a distortion of the symmetry of the palladium-allyl bond. In solution, two types of dynamic exchange have been observed, the η^3-η^1-η^3 rearrangement and the apparent rotation of the allyl moiety. At the same time, the folded structure of the ligand changes from an endo to an exo orientation or vice versa. The regioselectivity in the palladium catalyzed allylic alkylation of 3-methyl-but-2-enylacetate is determined by the cone angle of the bidentate phosphine ligand. Nucleophilic attack by a malonate anion takes place preferentially at the allylic carbon atom having the largest distance to palladium. Ligands with a larger cone angle direct the regioselectivity to the formation of the branched product, from 8% for dppe (1) to 61% for Xantphos (6).

The influence of the cone angle on the regioselectivity has been assigned to a sterically induced electronic effect.
7.1 Introduction

In many cases, palladium is the metal of choice in the synthetically useful catalytic allylic alkylation reaction.[1] When substrates resulting in symmetrically substituted Pd(allyl) complexes are used, such as cyclopent-2-enyl acetate or 1,3 diphenyl-prop-2-enyl-1-acetate, high enantioselectivities can be obtained.[2-4] The ligands that can be employed range from monodentate phosphorus ligands[9a] to bidentate phosphorus[2], nitrogen[3], or mixed bidentate P-N, P-S or phosphine-phosphite ligands[4].

When other types of allylic substrates are used, such as crotyl acetate or cinnamyl acetate, non-symmetrically substituted Pd(allyl) complexes are formed and regiocontrol[5] becomes an issue prior to enantiocontrol (see scheme 1). Three products can be formed: the non-chiral E and Z linear products and the chiral branched product. Excellent regio- and enantioselectivities have been obtained using P-N ligands with palladium as the metal[6]. The use of other metals, such as iridium,[7a-d] rhodium,[7c] or tungsten,[7b] can lead to analogous results, although the reaction rates are lower than those employing palladium. At this time, palladium based systems are still the most widely studied.

![Scheme 1: Regioselectivity in the palladium catalyzed allylic alkylation.](image)

Although many papers have appeared on enantioselective allylic alkylation, the subject of regioselectivity is studied less extensively. When a bidentate P-N or P-S ligand is used, the regioselectivity is determined by the difference in trans influence of the ligand donor atoms. The nucleophilic attack consequently takes place trans to the phosphorus donor (chapter 6), which in many cases exerts the strongest trans influence.[8]

For complexes containing bidentate P-P or N-N ligands, it has been suggested that the regioselectivity is determined by the bonding of the allyl moiety.[9] When the allyl group is substituted at one of its terminal positions, the symmetry of its bond to palladium is distorted. QSAR studies by Åkermark[9a] and our modeling studies (chapter 3) show that the Pd-allyl bond is changed from an η³ to an η¹–η² like structure (see scheme 2). The Pd-C1 bond is shorter than the Pd-C3 bond, the C1-C2 bond is elongated and the C2-C3 bond is shortened. In literature many examples are known of crystal structures of substituted allyl moieties that are not symmetrically bonded to palladium.[10] Based on NMR studies, it has been suggested that the electrophilicity of the substituted allylic carbon atom site is enhanced relative to the non substituted terminal site. It appears that the malonate nucleophile attacks preferentially at the allylic carbon atom with the largest Pd-C distance. This observation is supported by theoretical studies.[8a,9d]
P-P ligands, dimethyl allyl complexes.

Scheme 2: Distorted coordination of a substituted allyl moiety to palladium.

In chapter 4 we showed the effect of the bite angle of bidentate phosphine ligands on the structure of cationic (crotyl)Pd(ligand) complexes and their performance in the regioselective allylic alkylation. The complexes were isolated as equilibrium mixtures of two isomers, a syn and an anti isomer. Molecular modeling showed an increased embracing of the allyl moiety by the phenyl rings of the ligand when the bite angle is larger. Thus, it was predicted that an increase of the bite angle of the ligand resulted in an increase of the cone angle and consequently in a decrease of the syn / anti isomer ratio. Furthermore, the stoichiometric alkylation of these complexes with sodium diethyl 2-methylmalonate showed an increased regioselectivity for the branched product when the bite angle of the ligand is larger. To gain more insight into the factors influencing the regioselectivity, the effect of the bite angle on the symmetry of the Pd-allyl bond was studied.

To this end, cationic (3,3-(CH$_3$)$_2$-C$_5$H$_3$)Pd(ligand) complexes have been prepared, using ligands enforcing different bite angles (see scheme 3). The geometry of the complexes bearing the ligands dppe, dppf, DPEphos and Xantphos was studied in detail by X-ray crystallography. The isolated complexes have been used to determine the regioselectivity in the allylic alkylation, using sodium diethyl 2-methylmalonate as the nucleophile. In order to study the relation between the observed regioselectivity and the embracing effect of the ligand (chapter 4), we have calculated two steric ligand parameters from the crystal structures, the cone angle$^{[11]}$ and the solid angle.$^{[12]}$

Scheme 3: Structures of the complexes, numbering scheme and the ligands used.
Chapter 7

7.2 Results

A series of cationic (ligand)Pd(3,3-(CH)_3-C_3H_3) complexes (see scheme 3) has been prepared following a standard procedure. The ligand was added to the [(3,3-(CH)_3-C_3H_3)PdCl]_2 dimer, after which the chloride anion is abstracted using either AgBF_4 or AgOTf. The cationic (Xantphos)Pd(C_3H_5) complex (7) has been prepared in a similar way, starting from the [(C_3H_5)PdCl]_2 dimer.

7.2.1 Allylic alkylation

The complexes 1-6 were used in the allylic alkylation using sodium diethyl 2-methylmalonate as the nucleophile. Upon nucleophilic attack of the malonate nucleophile two regio-isomeric products can be obtained (see scheme 4). Bond formation at C1 yields the linear product, while bond formation at C3 it yields the branched product. The results of the stoichiometric reaction, presented in table 1, show that the regioselectivity towards formation of the branched product increases when the bite angle of the ligand is larger, ranging from 8% for dppe (1) to 61% for Xantphos (6). In chapter 4, a discrepancy was observed between the results of stoichiometric and catalytic alkylation due to competition between syn-anti isomerization and alkylation. In this investigation, only one isomer is possible leading to similar regioselectivities for the stoichiometric and the catalytic reaction (table 2). The rate of the catalytic reaction is dependent on the ligand, being relatively slow for dppe (1) and relatively fast for dppf (4).

Scheme 4: Alkylation products of complexes 1-6, X = BF_4, OTf, OAc (stoichiometric: BF_4, OTf; catalytic: OAc).
**P-P ligands, dimethyl allyl complexes.**

Table 1: Product distribution in the stoichiometric alkylation of complexes 1-6 (reaction conditions are described in the experimental section). All experiments were carried out in triplo, standard deviation of the reported regioselectivities is 0.5% or less.

<table>
<thead>
<tr>
<th>Complex</th>
<th>Bite angle</th>
<th>% branched</th>
<th>% linear</th>
</tr>
</thead>
<tbody>
<tr>
<td>dppe (1)</td>
<td>85.77(6)</td>
<td>8</td>
<td>92</td>
</tr>
<tr>
<td>dppp (2)</td>
<td>95</td>
<td>5</td>
<td>95</td>
</tr>
<tr>
<td>dppb (3)</td>
<td>99</td>
<td>11</td>
<td>89</td>
</tr>
<tr>
<td>dppf (4)</td>
<td>101.2(3)</td>
<td>13</td>
<td>87</td>
</tr>
<tr>
<td>DPEphos (5)</td>
<td>103.93(6)</td>
<td>41</td>
<td>59</td>
</tr>
<tr>
<td>Xantphos (6)</td>
<td>108.11(7)</td>
<td>61</td>
<td>39</td>
</tr>
</tbody>
</table>

a: value taken from crystal structure  
b: value taken from reference 10

Table 2: Results of the catalytic alkylation of 3-methyl-but-2-enyl acetate (reaction conditions are described in the experimental section). All experiments were carried out in triplo, standard deviation of the reported regioselectivities is 0.5% or less, the standard deviation of the reported rates are 5% or less.

<table>
<thead>
<tr>
<th>Complex</th>
<th>Bite angle</th>
<th>TOF (mole/mole/h)</th>
<th>% branched</th>
<th>% linear</th>
</tr>
</thead>
<tbody>
<tr>
<td>dppe (1)</td>
<td>85.77(6)</td>
<td>400</td>
<td>6</td>
<td>94</td>
</tr>
<tr>
<td>dppf (4)</td>
<td>101.2(3)</td>
<td>12000</td>
<td>13</td>
<td>87</td>
</tr>
<tr>
<td>DPEphos (5)</td>
<td>103.93(6)</td>
<td>6200</td>
<td>45</td>
<td>55</td>
</tr>
<tr>
<td>Xantphos (6)</td>
<td>108.11(7)</td>
<td>2500</td>
<td>63</td>
<td>37</td>
</tr>
</tbody>
</table>

a: initial turn over frequency, determined after 2 minutes  
b: determined after complete conversion  
c: value taken form crystal structure

7.2.2 **NMR experiments**

In the complexes, the effect of the ligand on the allyl moiety is reflected in the NMR spectra. In the $^1$H-NMR of complexes 1-6, the signals of the two methyl groups of the allyl moiety appear as separate signals. The chemical shift is dependent on the ligand and is found at higher field when the bite angle of the ligand is larger (table 3).

At room temperature, the signals of Ha and Hb are broadened in all cases, especially for complexes bearing ligands with a large bite angle. The signal of Hc appears as a triplet due to coupling with the Ha and Hb
The phosphorus spectra show two different signals, which are broad for the Xantphos modified complex (Table 3).

Table 3: Selected NMR data of the complexes 1-7, data determined at 298 K, δ in ppm.

<table>
<thead>
<tr>
<th>Complex (ligand)</th>
<th>Bite angle(°)</th>
<th>line width (P (Hz))</th>
<th>δ Me syn</th>
<th>δ Me anti</th>
</tr>
</thead>
<tbody>
<tr>
<td>1 (dppe)</td>
<td>85.77(6)°</td>
<td>8</td>
<td>1.90</td>
<td>1.45</td>
</tr>
<tr>
<td>2 (dppp)</td>
<td>95°</td>
<td>9</td>
<td>1.28</td>
<td>1.13</td>
</tr>
<tr>
<td>3 (dppb)</td>
<td>99°</td>
<td>6</td>
<td>1.36</td>
<td>0.82</td>
</tr>
<tr>
<td>4 (dpf)</td>
<td>101.2(3)°</td>
<td>3</td>
<td>1.11</td>
<td>1.11</td>
</tr>
<tr>
<td>5 (DPEphos)</td>
<td>103.93(6)°</td>
<td>10</td>
<td>1.23</td>
<td>1.06</td>
</tr>
<tr>
<td>6 (Xantphos)</td>
<td>108.11(7)°</td>
<td>91</td>
<td>0.92</td>
<td>1.04</td>
</tr>
<tr>
<td>7 (Xantphos)</td>
<td>108.11(7)°</td>
<td>3</td>
<td>--</td>
<td>--</td>
</tr>
</tbody>
</table>

a: value taken from crystal structure
b: value taken from reference 10

The broadened signals indicate the occurrence of dynamic exchange of the allyl moiety, which appears to proceed at a higher rate when the bite angle of the ligand is larger. This dynamic exchange of the allyl moiety of complex 6 has been studied in detail by means of variable temperature NMR. At -40°C the slow exchange limit is reached. Upon raising the temperature, apart from Ha and Hb, also the two methyl groups of the ligand backbone, which appear as separate signals due to its folded structure (see below), and the two phosphorus atoms exhibit exchange behavior. Rate data, obtained from simulation of the spectra show that the observed dynamic behavior is the result of two different processes. The exchange of the methyl groups of the ligand and the phosphorus atoms occur with the same rate (k = 450 s⁻¹ at 303 K, ΔHᵣ₂₉₈ = +71 kJ/mole, ΔSᵣ₂₉₈ = -88 J/mole.K), which is significantly higher than that of the Ha-Hb exchange (k = 100 s⁻¹ at 303 K, ΔHᵣ₂₉₈ = +56 kJ/mole, ΔSᵣ₂₉₈ = -30 J/mole.K).

To gain more insight in the different dynamic processes, a NOESY spectrum was recorded for complex 6 bearing the Xantphos ligand, that enforces a large bite angle. The methyl groups of the allyl appeared to have a steric interaction with the phenyl rings of the ligand. This interaction was studied in more detail by recording NOE difference spectra, in which the methyl groups were irradiated. Irradiation of the signal at 0.92 ppm, which was assigned to the syn CH₃ based on the value of ΔJ(P-CH₃), showed that this CH₃ group has a strong interaction with He. In contrast, the irradiation of the CH₃ signal at 1.04 ppm shows a much weaker interaction. These observations confirm the assignment of the signals to respectively the syn and anti CH₃ groups. The two CH₃ groups differ in interaction with the phenyl groups of the ligand. The interaction is stronger for the syn methyl than for the anti methyl group. This results in a shielding effect of the syn-CH₃ group in complexes 6, which is reflected in its resonance at unusual high field.
7.2.3 X-ray structures

The influence of the bite angle on the steric interaction of the ligand with the allyl moiety was also studied by X-ray crystallography. After recrystallization from CH\textsubscript{2}Cl\textsubscript{2} / hexane we obtained crystals suitable for X-ray structure determination of the cationic (ligand)palladium(3,3-(CH\textsubscript{3})\textsubscript{2}-C\textsubscript{3}H\textsubscript{5}) complexes with the ligands dppe (1), dppf (4) and DPEphos (5), whereas no suitable crystals were obtained of complexes bearing ligands with a bite angle larger than DPEphos. To study the coordination mode of the Xantphos ligand, the cationic (Xantphos)Pd(C\textsubscript{3}H\textsubscript{5}) complex was prepared and crystallized.

Selected structural data obtained from the X-ray structures (figures 1-4) are presented in tables 4 and 5.

Table 4: Selected bond distances obtained from the crystal structures (Å).

<table>
<thead>
<tr>
<th>Parameter</th>
<th>1 (dppe)</th>
<th>4 (dppf)</th>
<th>5 (DPEphos)*</th>
<th>7 (Xantphos)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Pd-C1</td>
<td>2.185(9)</td>
<td>2.192(15)</td>
<td>2.161(7)</td>
<td>2.17(1)</td>
</tr>
<tr>
<td>Pd-C2/C2a</td>
<td>2.174(7)</td>
<td>2.152(16)</td>
<td>2.21(1)</td>
<td>2.16(1)</td>
</tr>
<tr>
<td>Pd-C2b</td>
<td></td>
<td></td>
<td>2.21(1)</td>
<td></td>
</tr>
<tr>
<td>Pd-C3</td>
<td>2.253(7)</td>
<td>2.291(18)</td>
<td>2.37(1)</td>
<td>2.17(1)</td>
</tr>
<tr>
<td>C1-C2</td>
<td>1.42(1)</td>
<td>1.42(2)</td>
<td>1.39(1)</td>
<td>1.34(2)</td>
</tr>
<tr>
<td>C1-C2</td>
<td></td>
<td></td>
<td>1.28(1)</td>
<td></td>
</tr>
<tr>
<td>C2/C2a-C3</td>
<td>1.41(1)</td>
<td>1.41(2)</td>
<td>1.41(2)</td>
<td>1.34(2)</td>
</tr>
<tr>
<td>C2b-C3</td>
<td></td>
<td></td>
<td>1.59(2)</td>
<td></td>
</tr>
<tr>
<td>C3-C4 (anti)</td>
<td>1.51(1)</td>
<td>1.58(2)</td>
<td>1.48(1)</td>
<td></td>
</tr>
<tr>
<td>C3-C5 (syn)</td>
<td>1.50(1)</td>
<td>1.44(2)</td>
<td>1.48(1)</td>
<td></td>
</tr>
<tr>
<td>Pd-P (cis to C1)</td>
<td>2.296(2)</td>
<td>2.313(6)</td>
<td>2.336(2)</td>
<td>2.372(2)</td>
</tr>
<tr>
<td>Pd-P (cis to C3)</td>
<td>2.293(2)</td>
<td>2.33(9)</td>
<td>2.379(2)</td>
<td>2.372(2)</td>
</tr>
</tbody>
</table>

*: C2a corresponds to C85 in figure 3 and C2b corresponds to C77 in figure 3.
Table 5: Selected geometrical data obtained from the crystal structures, distances in Å, angles in °.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>1 (dppe)</th>
<th>4 (dppf)</th>
<th>5 (DPEphos)*</th>
<th>7 (Xantphos)</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\angle (P-Pd-P)$</td>
<td>85.77(6)</td>
<td>101.2(3)</td>
<td>103.93(6)</td>
<td>108.11(7)</td>
</tr>
<tr>
<td>$\angle (C-C-C)$</td>
<td>121.2(7)</td>
<td>124(1)</td>
<td>120.2(7)(a)</td>
<td>115(1)(b)</td>
</tr>
<tr>
<td>d(C5(syn)-closest Ph-plane)</td>
<td>3.44(2)</td>
<td>3.54(3)</td>
<td>3.20(1)$^a$</td>
<td></td>
</tr>
<tr>
<td>d(C1)-(PPdP-plane)</td>
<td>+0.044(10)</td>
<td>-0.17(2)</td>
<td>+0.319(8)</td>
<td>-0.35(1)</td>
</tr>
<tr>
<td>d(C2)-(PPdP-plane)</td>
<td>-0.440(10)</td>
<td>-0.35(3)</td>
<td>+0.91(1)$^c$</td>
<td>-0.47(1)$^d$</td>
</tr>
<tr>
<td>d(C3)-(PPdP-plane)</td>
<td>+0.329(9)</td>
<td>+0.34(3)</td>
<td>-0.13(1)</td>
<td>-0.348(3)</td>
</tr>
<tr>
<td>$\angle ((CCC-plane)-(PPdP-plane))$</td>
<td>114.4(9)</td>
<td>114(2)</td>
<td>111(1)$^e$</td>
<td>99(1)$^d$</td>
</tr>
<tr>
<td>$\angle ((C4-C3-C5-plane)-closest-Ph-plane)$</td>
<td>7.9(5)</td>
<td>5(1)</td>
<td>10.2(7)</td>
<td></td>
</tr>
</tbody>
</table>

*: C2a corresponds to C85 in figure 3 and C2b corresponds to C77 in figure 3.

a: C81 in figure 3
b: C65 in figure 3
c: C2 corresponds to C77 in figure 3
d: C2 corresponds to C85 in figure 3

Figure 1: Two views of the crystal structure of complex 1 (dppe).
Figure 2: Two views of the crystal structure of complex 4 (dpf).

Figure 3: Two views of the crystal structure of complex 5 (DPEphos).
All structures are cationic, since the counterion is located at a large, non-bonding distance from palladium. In all complexes two phenyl rings of the ligand, located in the P-Pd-P plane, point in the direction of the allyl moiety. The planes of the -C(CH₃)₂ fragment and the closest phenyl ring are almost parallel (table 5). The two other phenyl ring of the ligand point in the same direction as C2.

The DPEphos ligand (β = 103.93(6)°) shows an interaction between one of the aromatic rings (C--C) of the backbone and one of the phenyl rings bound to phosphorus (C--C) (figure 3). The angle between the two planes is 23.8(2)° and the minimum distance is 3.37(3) Å (d(C54-C58)). This type of orientation for DPEphos has also been observed in the two other crystal structures known of palladium complexes of this ligand, the zerovalent (tce)Pd(DPEphos) complex[14] and the (DPEphos)Pd(para-C₆H₄-CN)Br complex[15].
With respect to C2, the ferrocene unit of the dppe ligand is located under the P-Pd-P plane, and is twisted as a result of its staggered conformation (figure 2). The two Cp rings are staggered and almost parallel (angle = 3.6(5)°). Also in this complex, two phenyl rings of the ligand point in the direction of the allyl moiety.

The Xantphos complex (figure 4) has C₃ symmetry, two phenyl rings of the ligand point in the direction of the allyl moiety and two are directed upwards with respect to C₂. The latter phenyl groups are almost parallel (13.4(2)°) and the distance between the ipso carbon atoms is 3.87(2) Å.

The two aromatic rings in the backbone of the Xantphos ligand are not coplanar as a result of the sp³ carbon atom and the oxygen atom in the bridge. The value found for the angle between the planes (27.4(5)°) is similar to that found for the zerovalent (tcne)Pd(Xantphos) complex reported previously[14]. As a result of this folded structure, one of the methyl groups of the backbone is located equatorially and one axially, resulting in two different signals in the NMR spectrum (see above).

As expected, the C1-C2-C₃ plane is not perpendicular to the P-Pd-P plane. The angle between these planes appears to be dependent on the ligand, as it ranges from 114.4(9)° for dppe to 100(2)° for Xantphos (table 5).

In all C₅H₉-complexes, the allyl moiety is not symmetrically bound to the palladium center. The Pd-C₁ bond is shorter than the Pd-C₃ bond and the C₁-C₂ bond is longer than the C₂-C₃ bond (table 4). Considering these differences between the bond lengths, the non-symmetry of the allyl moiety is more pronounced when the bite angle of the ligand is larger.

In the complex of dppe, the bond between C₃ and the syn methyl substituent on the allyl moiety is slightly shorter (1.50(1) Å) than the bond from C₃ to the anti substituent (1.51(1) Å). When the bite angle is larger, as in the case of dppe, the C₃-C₅(syn) bond becomes shorter (1.44(2) Å) and the C₃-C₄(anti) bond is longer (1.58(2) Å).

It has been reported, that in some cases the allyl moiety is slightly rotated around the Pd-allyl axis[4c, 4d, 9]. This rotation towards a product-like state could be important for determining the regioselectivity. In the crystal structures we present here, no significant rotated orientation of the allyl group is observed and no correlation with the bite angle of the ligand is found (table 5).

The palladium phosphorus distances are in the same range as found previously for zerovalent (P-P)Pd(tcne) complexes.[14] In general, the bond between palladium and phosphorus cis to C₁ is shorter than that to phosphorus cis to C₃. This difference is more pronounced when the bite angle of the ligand is larger; it is approximately zero for the dppe ligated complex and amounts to 0.02 Å for dppe (4) and to 0.04 Å for DPEphos (5). The distance between palladium and the phosphorus atoms is also dependent on the bite angle of the ligand. A smaller bite angle results in a smaller Pd-P distance. It ranges from 2.293(2) Å for dppe to 2.372(2) Å for Xantphos.
7.2.4 Cone angle, solid angle

From the above observations it is clear, that the bite angle of the ligand influences the symmetry of the Pd-allyl bond via steric interaction. The top views of the crystal structures show that the phenyl rings are closer to the allyl moiety when the bite angle of the ligand is larger, as was already suggested by the results of the semi-empirical molecular modeling in chapter 4. The bite angle, however, is an indirect parameter for determining the amount of steric interaction a ligand induces. Therefore we decided to determine the values of two direct parameters from the X-ray structures, the cone angle $\theta$ introduced by Tolman, and the solid angle $\Omega$ as introduced by Bagnall and Xing-Fu.

The results in table 6 show in general the same trend for each of these geometrical parameters. Going from dppe to Xantphos, the bite angle, the cone angle and the solid angle increase.

Table 6: Values of the bite angle, the cone angle and the solid angle, as determined from the X-ray structures.

<table>
<thead>
<tr>
<th>Complex</th>
<th>Bite angle (°)</th>
<th>Cone angle (°)</th>
<th>Solid angle (sterrad)</th>
<th>% branched product</th>
</tr>
</thead>
<tbody>
<tr>
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<td>7 (Xantphos)</td>
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<td>246.9</td>
<td>5.56</td>
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</tr>
</tbody>
</table>

7.3 Discussion

7.3.1 Structure of the complexes in the solid state

From the X-ray data, it can be concluded that the presence of substituents on the allyl moiety distorts the symmetry of the bond to palladium (see scheme 5). This effect has been observed before, both in other crystal structures and in modeling studies [4-9]. In all cases, it is observed that the bonding distance between C3 and palladium is longer than between palladium and C1. The weakened bonding between the substituted allylic site and palladium corresponds to an enhanced bond strength of the allylic C2-C3 bond. In the present case, the distortion is dependent on the ligand used and, specifically, on the bite angle of the ligand used. A larger bite angle results in an increase of the cone angle of the ligand and therefore in a more pronounced embracing of the allyl moiety by the phenyl rings of the ligand (see table 6). The increase in solid angle with larger bite angle indicates that there is less space for the allyl ligand to coordinate to palladium. In figure 5, a space-filling model of the crystal structure containing the Xantphos ligand is presented. There is enough space for the allyl moiety to coordinate but addition of two extra methyl substituents on one of its terminal positions will result in significant steric interaction (see figure 5). This leads to a lengthening of the Pd-C3
P-P ligands, dimethyl allyl complexes.

bond and changes in bond lengths as shown in scheme 5. Therefore it is argued that the steric effect of the bite angle on the dissymmetry of the Pd-allyl bond induces electronic effects on the allyl moiety.

Scheme 5: Schematic representation of the distortion of the Pd-allyl bond as observed in the X-ray structures.

Figure 5: Space filling model of the X-ray structure of complex 7, the allyl moiety at the bottom of the figure.

The distortion of an $\eta^3$ towards an $\eta^1-\eta^2$ bonding mode as observed in the crystal structures, will result in a lower value of the overlap integral on the $\eta^2$-site and a higher value of this integral on the $\eta^1$-site. On the $\eta^2$-site, both the donating interaction (ligand to metal) and the back-donating interaction (metal to ligand) will be decreased with respect to the symmetrically $\pi$-bonded allyl moiety\[^8c, 9^c]\). Thus, as a result of the lengthening of the Pd-C3 distance the back-donation to the LUMO of this site of the allyl moiety is lower than in a symmetrically bonded $\pi$-allyl moiety. The Pd-C1 distance is decreased and the C1-C2 distance is increased, which both indicate an increase in the L→M donation as well as the M→L back-donation on the $\eta^1$-site of the allyl moiety.

The difference in the two allylic binding sites is reflected in the bond length between palladium and phosphorus. The Pd-P cis to the non-substituted C1 is shorter than the bond trans to C1. So, the two ligands that are most strongly bound to the metal center are cis to one another.

The effects of changes in donation and back-donation are also reflected in the value for the angle between the allylic CCC plane and the P-Pd-P plane. In the absence of back-donation, a value close to 90° would be expected. When the back-donation to C1 and C3 is strong, the value for this angle ($\angle((\text{CCC})-(\text{PPdP}))$) will
increase. In the crystal structures presented in this chapter, the value of \( \angle((\text{CCC})-(\text{PPdP})) \) is dependent on the ligand. When the bite angle of the ligand is larger, this angle decreases. It is concluded that an increase in the bite angle results in a decrease of the back-donation. As a consequence, the allylic LUMO orbitals, which are the sites of reaction in the nucleophilic attack, remain less occupied. Therefore, a larger bite angle of the ligand results in an increase of the reactivity of the allyl moiety towards nucleophilic attack. This has indeed been observed in the catalytic alkylation of trans substrates (chapters 4 and 5). The catalytic experiments, however, show a lower reaction rate for the Xantphos ligand. This can be explained in terms of increased steric hindrance (see below).

\( \text{7.3.2 Structure of the complexes in solution} \)

The NOE experiments carried out on the cationic \((3,3-(\text{CH}_3)_2-\text{C}_6\text{H}_3)\text{Pd(Xantphos)}^+\) complex (6) revealed steric interactions between the methyl groups and the phenyl rings of the ligand. The phenyl rings show a stronger interaction with the syn methyl, compared to the anti methyl group, whereas in the X-ray structure of complex 7, the anti hydrogen atom is closest to the phenyl rings. A slight rotation about the P-C bond in the X-ray structure would bring the phenyl rings at a closer distance to the syn substituent than to the anti substituent. In solution, some rotation about the P-C bond may take place and our results indicate that on average the syn substituent is at a closer distance to the phenyl rings. The crystal structures show that the steric interaction is stronger when the bite angle of the ligand is larger.

Complex 6, as well as the other complexes, can in theory be present in two pairs of enantiomeric structures (scheme 6): the methyl groups can be oriented trans to P1 or trans to P2 and the axial phenyl rings of the ligand (figure 1-4) can be oriented in the same direction as C2 (endo) or in the opposite direction (exo). In the crystal structures, only the endo isomer is found for complexes 1, 4 and 7, whereas for complex 5 both isomers are found to crystallize together. In the NMR spectra only one complex is observed, also at low temperature, which is therefore assigned to be the endo isomer. It can not be ruled out that the different complexes which are possibly present at reaction temperature show different regioselectivities in the allylic alkylation.

\[\text{Scheme 6: Two pairs of enantiomers of complex 6.}\]
Variable temperature NMR experiments carried out for complex 6, showed that several types of fluxional behavior occur simultaneously. The slower process involves the Ha/Hb exchange and the faster process involves the phosphorus atoms and the two methyl groups of the ligand, which at low temperature appear as separate signals due to the folded structure of the backbone (see above). The exchange of the methyl groups of the ligand indicates that during the faster process, the backbone of the ligand folds from one orientation to the other, thereby exchanging the methyl groups from an equatorial to an axial position and vice versa.

The relatively high negative value for the entropy of activation suggests that the faster process occurs via an associative pathway. It is known, that temporary coordination of the counterion of cationic (allyl)palladium complexes may enhance the rate of dynamic exchange of the allyl moiety.\[^{16b}\] In a thus formed five coordinated Pd(allyl) complex, the so-called apparent rotation of the allyl moiety can easily take place.\[^{16b}\] In complex 6, such a process will cause the exchange of C1 and C3 and vice versa, resulting in the equivalency of the two phosphorus atoms. Apparently, during this process, the backbone of the ligand changes its folded structure from one form to the other, thereby exchanging the two methyl groups of the ligand.

The Ha/Hb exchange, observed in the slower process, is known to be caused by a $\eta^1$-$\eta^1$-$\eta^1$ rearrangement\[^{171}\]. The Pd-C3 and Pd-C2 bonds are broken selectively and a Pd-C1 $\sigma$-bond is formed. Rotation about the Pd-C1 bond and the C1-C2 bond results in an exchange of the signals of Ha and Hb. As at -40°C only one complex is observed, it is concluded that, contrary to our previous results,\[^{8b}\] the $\pi-\sigma$ rearrangement of the allyl moiety observed for complex 6 does not lead to the exchange of the *endo* and *exo* isomeric forms of the complex.

If the $\eta^1$-$\eta^1$-$\eta^1$ rearrangement process occurs via a T-shaped palladium intermediate, in which no exchange of the coordination sites of C1 and C3 takes place, the exchange of *endo* to *exo* and vice versa will take place.\[^{88}\] On the other hand, if it occurs via a C, symmetric Y-shaped intermediate, in which a C1-C3 exchange will take place, the *endo-*exo exchange may not occur (scheme 7). Recently, we have found evidence for coordination of the oxygen atom of the ligand backbone to palladium in cationic (Xantphos)Pd(Ph) complexes.\[^{15}\] This may also occur in the Y-shaped intermediate. The thus formed flat structure of the backbone can fold back to form the *endo* isomer of the $\eta^1$-$\eta^1$-$\eta^1$ rearranged complex. Because the $\eta^1$-$\eta^1$-$\eta^1$ rearrangement is the slower process, it can not be distinguished whether it involves a T- or a Y-shaped intermediate.

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7.3.3 Allylic alkylation

Several theoretical studies concerning the mechanism of the allylic alkylation reaction have been reported in literature.\(^{96,17}\) Recently, a theoretical rationale has been presented\(^{94}\) for the correlation between the regioselectivity and the non-symmetry of the allyl moiety\(^{91}\). By means of DFT calculations (ADF) it was shown, that an electronic preference for initial nucleophilic attack on the carbon atom of the allyl moiety with the largest Pd-C distance, leads to a lowering of the energy barrier encountered in a later stage of the reaction. Our results may be explained following the same rationale.

First, the malonate nucleophile selects the site of attack, based on electronic properties (charge, LUMO coefficients) and steric (accessibility) properties. Therefore, the regioselectivity for initial attack at C3 will be a trade-off between the larger steric hindrance the nucleophile encounters during attack at this position and the electronic preference.

During the process of bond formation,\(^{17}\) the allyl moiety rotates to form a transient palladium-olefin complex\(^{18}\). When the attack takes place at C3, the branched site, a terminal C=C double bond is formed. The formation of the transient palladium-olefin complex implies that the substituted carbon atom has been rotated out of the P-Pd-P plane, thereby minimizing the steric interaction with the phenyl rings of the ligand. Alternatively, attack on C1 results in the formation of an internal C=C double bond, substituted by three groups. In the resulting palladium-olefin complex, the methyl substituents are at a closer distance to palladium than they were in the allyl complex, so the steric hindrance with the ligand is increased. In this stage of the reaction, the height of the energy barrier is also influenced by the thermodynamically favorable formation of an internal C=C double bond compared with that of a terminal alkene.

The preference for formation of the branched product when the bite angle of the ligand is larger, can therefore be explained both in steric and electronic terms. Thus, the selectivity towards the formation of the linear product for the dppe ligated complex can be explained by a less activated allyl moiety due to less steric interaction with the ligand (see above). Consequently, the steric accessibility of C3 relative to C1 is important. Also in the second stage of the reaction, when the more favored internal alkene is formed, the steric hindrance in the palladium-olefin complex will be less than that in complexes of the other ligands.

Scheme 7: Different intermediates for the $\eta^3$-$\eta^1$-$\eta^3$ rearrangement.
P-P ligands, dimethyl allyl complexes.

In contrast, the use of ligands with a larger bite angle will result in the formation of a more activated allyl moiety in the corresponding palladium complex. In the trade-off between steric access and electronic reactivity, the electronic factors may prevail. During the rotation of the allyl moiety, the steric hindrance encountered during terminal olefin formation (branched product) will be less than that for internal olefin formation (linear product). We therefore propose that the regioselectivity of the allylic alkylation is determined by a set of steric / electronic trade-off pathways (see scheme 8).

The large difference in the amount of branched product formed from the dpff (4) and the DPEphos (5) modified complexes would not be expected based on the small difference in bite angle. In general, comparing the values found for the three different ligand parameters (bite angle, cone angle, solid angle) with the regioselectivity, reveals that the cone angle shows the best correlation.

The total effect of the ligand geometry on the rate of the catalytic reaction is dependent on several effects. The reaction consists of a number of consecutive stages: (1) substrate coordination to palladium, (2) oxidative addition, (3) nucleophilic attack and (4) product dissociation. A large cone angle will slow down stage (1), as a result of large steric hindrance. The study of the crystal structures shows that a large cone angle may enhance the reaction rate of stage (3) and (4). The oxidative addition process can be regarded as the reverse process of nucleophilic attack, so it can be expected that also stage (2) may proceed slower when the cone angle is larger. Thus, the total effect of the cone angle of the ligand on the rate is determined by a trade-off between the accelerating and the decelerating effects.

Scheme 8: Trade-off pathways that may determine the regioselectivity.
Chapter 7

7.4 Conclusion

We have shown, that in crystal structures an increase of the bite angle of a bidentate phosphorus ligand leads to an increase in both the cone angle and solid angle of the ligand. As the effect of the ligand on the geometry and reactivity of the complexes is best described in terms of steric hindrance, the cone angle of the ligand is the parameter of choice.

The more pronounced embracing of the allyl moiety by the ligand distorts the symmetry of the Pd-(3,3-(CH$_3$)$_2$-C$_3$H$_5$) bond to an $\eta^1$-$\eta^2$ type coordination. Furthermore, the backbonding to the substituted site of the allyl moiety decreases when the cone angle is large. This sterically induced electronic effect has a pronounced influence on the reactivity of the allylic carbon atoms and the regioselectivity of stoichiometric alkylation.

A larger cone angle of the ligand enhances the electronic preference for nucleophilic attack on the branched position, but also the steric hindrance at this position is enhanced. Therefore it is concluded, that the regioselectivity of the reaction is a result of a trade-off between electronic and steric interactions.

7.5 Experimental Section

7.5.1 General procedures

$^1$H NMR (300 MHz, TMS, CDCl$_3$), $^{31}$P ($^1$H) (121.5 MHz external 85% H$_3$PO$_4$, CDCl$_3$) were recorded on a Bruker AMX-300 spectrometer.

The product distribution of allylic alkylation was measured on an Interscience Mega2 apparatus, equipped with a DB1 column, length 30 m, inner diameter 0.32 mm, film thickness 3.0 $\mu$m, and a F.I.D detector.

All experiments were carried out using standard Schlenk techniques. All solvents were freshly distilled prior to use. All reactions have been performed at room temperature (292 K).

The variable temperature NMR experiments have been carried out by measuring spectra at different temperatures, ranging from 233K (dynamic behavior frozen out) to 328 K (coalesced signals, towards fast exchange). The simulation of the spectra was carried out using gNMR software.$^{109}$

Sodium diethyl 2-methylmalonate (0.5 M in THF) was prepared from diethyl 2-methylmalonate and NaH in THF at 273 K. All alkylation experiments were carried out in triplo.

The stoichiometric alkylation reactions were performed by adding an excess of sodium diethyl 2-methylmalonate (0.1 ml of a 0.5 M solution in THF) to a solution of 10 mg of the Pd-complex in 1 ml of THF. Reaction was instantaneous and after one minute, the mixture was worked up with water, filtered over silica and analyzed by GC.

The catalytic reactions were performed in THF (10 mL), using 0.05 mol% of catalyst (0.00050 mmole), 1.0 mmole of 3-Me-but-2-enyl acetate and 2.0 mmole of sodium diethyl 2-methylmalonate. The reaction was
P-P ligands, dimethyl allyl complexes.

monitored by taking samples from the reaction mixture, which after quenching with wet ether, were analysed by GC using decane as the internal standard.

The Pd-complexes were prepared in CH$_2$Cl$_2$ from [(C$_5$H$_5$)-Pd-μCl]$_2$ by adding 2 equiv. of ligand and abstracting the Cl-atom with AgOTf. The complexes were isolated in quantitative yield (white microcrystalline powder) as their analytically pure equilibrium mixtures and were used as such in the alkylation reaction. The syntheses of DPEphos and Xantphos have been published elsewhere. Dppe, dpp, dppb and dpff were obtained from Acros chemicals and used as received.

7.5.2 Characterization

(C$_5$H$_5$)Pd(dppe)OTf (1): $^1$H: 1.05 (t, 3H, $J_1 = J_2 = 6.3$ Hz, anti-CH$_3$); 1.90 (t, 3H, $J_1 = J_2 = 8.4$ Hz, syn-CH$_3$); 2.5-2.8 (br m, 4H, 2P-CH$_2$); 3.6-4.0 (br, 2H, allylic-CH$_2$); 5.53 (t, 1H, $J_1 = J_2 = 11.0$ Hz); 7.3-7.9 (br m, 20H, aromatic H)

$^{31}$P: 46.8 (d, 1P, $J = 34$ Hz); 51.1 (d, 1P, $J = 34$ Hz)

$^{13}$C{$^1$H}: 20.2 (d, $J = 5$ Hz); 26.6; 26.9 (d, $J = 13$ Hz); 28.5 (d, $J = 13$ Hz); 60.6; 61.0; 103.5; 104; 107.9; 115.0; 115.1; 115.2; 116.7; 116.8; 117.1; 117.3; 129.5 (d, $J = 11$ Hz); 131.8; 132.3 (d, $J = 13$ Hz)

HR-MS (FAB): C$_{31}$H$_{33}$P$_2$Pd$^+$ requires m/z = 573.1092, found m/z = 573.1102

(C$_5$H$_5$)Pd(dppp)OTf (2): $^1$H: 1.14 (dd, 3H, $J_1 = J_2 = 6.3$ Hz, anti-CH$_3$); 1.28 (dd, 3H, $J_1 = J_2 = 9.6$ Hz, syn-CH$_3$); 1.6 (br m, 1H, C(H)H-CH$_2$-P); 2.2 (br m, 1H, C(H)H-CH$_2$-P); 2.75 (m, 2H, -CH$_2$-P); 2.8 (dd, 1H, $J_1 = J_2 = 13$ Hz, allylic H); 2.95 (m, 2H, -CH$_2$-P); 3.65 (dd, 1H, $J_1 = J_2 = 7.2$ Hz, allylic H); 5.27 (dd, 1H, $J_1 = 8.1$ Hz, $J_2 = 13.8$ Hz, allylic H); 7.2-7.7 (br m, 20H, aromatic H)

$^{31}$P: 6.5 (d, $J = 65.7$ Hz); 9.8 (d, $J = 65.3$ Hz)

$^{13}$C{$^1$H}: 19.0; 20.7; 25.9; 26.3; 26.6; 64.1 (d, $J = 29$ Hz); 110.1 (d, $J = 26$ Hz); 114.5; 128.4; 129.2; 129.3; 129.5; 129.6; 130.5; 131.0; 131.8; 131.9; 132.7 (d, $J = 11$ Hz); 133.8 (d, $J = 12.4$ Hz); 134.3 (d, $J = 14$ Hz)

HR-MS (FAB): C$_{32}$H$_{35}$P$_2$Pd$^+$ requires m/z = 587.1249, found m/z = 587.1255

(C$_5$H$_5$)Pd(dppb)BF$_4$ (3): $^1$H: 0.82 (dd, 3H, $J_1 = J_2 = 6.0$ Hz, anti-CH$_3$); 1.36 (dd, 3H, $J_1 = 7.4$ Hz, $J_2 = 10.1$ Hz, syn-CH$_3$); 1.7 (br m, 4H, CH$_2$-CH$_2$-P); 2.7 (br m, 5H, CH$_2$-CH$_2$-P, allylic H); 3.7 (br, 1H, allylic H); 5.45 (t, 1H, $J_1 = J_2 = 10.8$ Hz); 7.2-7.8 (br m, 20H, aromatic H)

$^{31}$P: 21.3 (d, AB, 1P, 49 Hz); 22.2 (d, AB, 1P, 48 Hz)

$^{13}$C{$^1$H}: 20.0; 22.9; 25.4; 26.3; 26.9; 63.1; 106.1; 106.9; 113.9; 116.5; 116.6; 129.2; 130.7; 131.3; 134.2; 135.0

HR-MS (FAB): C$_{33}$H$_{37}$P$_2$Pd$^+$ requires m/z = 601.1405, found m/z = 601.1400
(C₅H₅)Pd(dpff)BF₄ (4): ¹H: 1.1 (m, 6H, 2CH₃); 2.82 (t, 1H, J₁ = J₂ = 11.4 Hz, anti-allylic H); 3.68 (t, J₁ = J₂ = 7.2 Hz, syn-allylic H); 3.87 (s, 1H, FcH); 4.19 (s, 1H, FcH); 4.27 (s, 1H, FcH); 4.32 (s, 1H, FcH); 4.41 (s, 1H, FcH); 4.48 (s, 2H, FcH); 4.55 (s, 1H, FcH); 5.49 (dd, 1H, J₁ = 8.1 Hz, J₂ = 13.8 Hz, Hc); 7.3-7.6 (br m, 18 H, aromatic H); 7.85 (br m, 2H, aromatic H)

³¹P: 22.7 (d, J₁ = 46 Hz); 26.9 (d, J₁ = 46 Hz)

¹³C¹¹[H]: 15.5; 21.1; 26.5; 66.1; 114.3; 116.0; 118.4; 129.2; 129.3; 129.4; 129.5; 129.6; 130.3; 130.8; 131.3; 131.4; 131.5; 132.1; 132.4; 132.6; 133.1; 133.5; 134.0; 134.4; 134.9

HR-MS (FAB): C₃ số₉ FeP₂Pd⁺ requires m/z = 729.0755, found m/z = 729.0748

(C₅H₅)Pd(DPEphos)BF₄ (5): ¹H: 1.06 (t, 3H, J₁ = J₂ = 5.6 Hz, anti-CH₃); 1.23 (dd, 3H, J₁ = 5 Hz, J₂ = 11 Hz, syn-CH₃); 2.84 (t, 1H, J₁ = J₂ = 12.5 Hz, anti-allylic H); 3.63 (dt, 1H, J₁ = J₂ = 7.3 Hz, J₃ = 2.0 Hz, syn-allylic H); 5.51 (dd, 1H, J₁ = 7.9 Hz, J₂ = 13.4 Hz, Hc); 6.6-7.6 (m, 28H, aromatic H)

³¹P: 13.8 (d, J₁ = 41 Hz); 14.7 (d, J₁ = 42 Hz)

¹³C¹¹[H]: 15.1; 21.0; 26.0; 65.6; 113.1; 116.3; 116.4; 118.4; 122.2; 122.6; 122.8; 124.1 (d, J = 6.5 Hz); 125.5 (d, J = 6.5 Hz); 127.4; 128.0; 128.8; 129.0; 130.8; 131.0; 131.2; 134.3 (d, J = 13 Hz); 135.1; 157.9 (d, J = 7.9 Hz)

HR-MS (FAB): C₄₁H₃₇OP₂Pd⁺ requires m/z = 713.1354, found m/z = 713,1373

(C₅H₅)Pd(Xantphos)OTf (6): ¹H: 0.9 (t, 3H, J₁ = J₂ = 8.6 Hz, syn-CH₃); 1.04 (t, 3H, J₁ = J₂ = 5.7 Hz, anti-CH₃); 1.4-1.9 (br, 6H, ligand-CH₃’s); 2.74 (br, 1H, anti-allylic H); 3.5 (br, 1H, syn allylic H); 5.70 (t, 1H, J₁ = J₂ = 10.4 Hz); 6.67 (t, 2H, J₁ = J₂ = 7.9 Hz); 7.0-7.6 (br m, 22H, aromatic H); 7.64 (d, 2H, J = 7.6 Hz)

³¹P: 5.3 (br, 1P); 10.4 (br, 1P)

¹³C¹¹[H]: 36.0; 116.0; 116.3; 116.4; 118.8; 123.8; 123.9; 124.6; 124.7; 127.8; 128.8; 129.0; 130.6; 131.7; 132.7; 134.0; 135.4

HR-MS (FAB): C₄₄H₃₇O₂Pd⁺ requires m/z = 753.1667, found m/z = 753.1685

(C₃H₅)Pd(Xantphos)BF₄ (7): ¹H: 1.54 (s, 3H, ligand-CH₃); 1.81 (s, 3H, ligand-CH₃); 3.5 (m, 2H, anti-allylic H); 3.8 (m, 2H, syn-allylic H); 6.0 (m, 1H, central allylic H); 6.6 (br t, 2H, Ar-H); 7.0-7.5 (br m, 22H, aromatic H); 7.63 (d, J = 7.8 Hz, 2H, aromatic H)

³¹P: 4.3 (s); ¹³C¹¹[H]: 25.9; 30.1; 36.3; 118.0; 118.0; 123.1; 125.0; 128.4; 129.3; 130.9; 132.3; 132.7; 133.2; 134.4; 155.2

HR-MS (FAB): C₄₂H₃₇OP₂Pd⁺ requires m/z = 725.1354, found m/z = 725.1355
7.5.3 Calculation of cone angle and solid angle

In the crystal structures, the allyl moiety was removed and a dummy atom was placed in the center of the P-P axis at a non fixed distance to palladium. The two-dimensional cone angle in the P-Pd-P plane has been calculated viewed from the palladium atom in the direction of the dummy atom. The values thus found for the ligands are listed in table 6. In all cases, the ligand has a cone angle larger than 180°. The value found for dppe is the smallest (224.6°) and increases for ligands with a larger bite angle to 246.9° for the Xantphos ligand.

A direct comparison between the different ligands is hampered by the fact that some atoms of the phenyl rings of the ligand can be located slightly below or above the P-Pd-P plane. Since these atoms are not taken into account by the two-dimensional cone angle θ, we calculated the three-dimensional solid angle Ω of these ligands in an analogous manner (see scheme 9). The ligand is again viewed from the palladium atom in the direction of the dummy atom. The solid angle is then determined as a three-dimensional cone angle. The values listed in table 6 represent the space that the ligands occupy after projection on the central metal atom. The thus determined value for the size of the ligand appears to be dependent on the bite angle of the ligand and increases when the bite angle is larger. Again, the smallest value is found for dppe (4.47 sterrad) and the largest is found for Xantphos (5.56 sterrad). The relatively large value found for the dppf ligand is mainly due to the presence of the large ferrocene unit at a short distance to palladium.

Scheme 9: Determination of the cone angle and the solid angle, D=dummy atom (necessary for determining plane of the cone angle).
### 7.5.4 Crystal structure determination

Table 7: Crystallographic data for compounds 1, 4, 5 and 7.

<table>
<thead>
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<th>Complex</th>
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<th>4</th>
<th>5</th>
<th>7</th>
</tr>
</thead>
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<td><strong>Formula</strong></td>
<td>[C₅H₁₃OP₂Pd]⁺ SO₃CF₃</td>
<td>[C₆H₁₃OP₂FePd]⁺ BF₄⁻</td>
<td>[C₄H₁₃OP₂Pd]⁺ BF₄⁻</td>
<td>[C₆H₁₃OP₂Pd]⁺ BF₄⁻ CH₂Cl₂</td>
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<tr>
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<td>13.067(2)</td>
<td>12.519(2)</td>
<td>18.0851(9)</td>
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<tr>
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<td>18.267(3)</td>
<td>14.435(2)</td>
<td>12.6484(8)</td>
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<tr>
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<td>78.01(1)</td>
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<tr>
<td>β, °</td>
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<td>83.84(1)</td>
<td>108.01(1)</td>
<td>105.409(9)</td>
</tr>
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<td>γ, °</td>
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<td>1930.7(5)</td>
<td>2018.5(5)</td>
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<td>μ, cm⁻¹</td>
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<td>R_w</td>
<td>0.067</td>
<td>0.080</td>
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* R = Σ(|F_obs| - |F_calc|) / Σ(|F_obs|), ** R_w = Σ(|F_obs| - k |F_calc|)^2 / Σ(|F_obs|^2)

Data (table 7) were collected on an Enraf-Nonius CAD-4 diffractometer with graphite-monochromated CuKα radiation and θ-2θ scan. Corrections for Lorentz and polarisation effects were applied.

Absorption correction was performed with the program PLATON following the method of North et al. using Ψ-scans of five reflections for all four compounds. The structures were solved by the PATTY option of the DIRDIF96 program system. The hydrogen atoms were calculated. Full-matrix least-squares refinement was carried out on F, anisotropic for the non-hydrogen atoms and isotropic for the hydrogen atoms, restraining the latter in such a way that the distance to their carrier remained constant at approximately 1.0Å converged. Scattering factors were taken from Cromer and Mann; International Tables for X-ray
Crystallography.[24] Anomalous scattering was taken into account.[25] All calculations were performed with XTAL,[26] unless stated otherwise. For (4) the BF$_4^-$ moiety was kept fixed at ideal geometry with isotropic temperature factors U=0.15 Å$^2$ for B and U=0.25 Å$^2$ for F; the hydrogen atoms were kept fixed at their calculated positions with U=0.10 Å$^2$. (5) showed some disorder which was dealt with in the following way: C(2) was divided over two half occupied positions and all H atoms connected to atoms C(1)-C(5) were kept fixed at their calculated positions with U=0.10 Å$^2$; the BF$_4^-$ moiety also showed disorder and all F-atoms were divided over three 1/3 occupied position and were kept restrained geometrically to values from literature and were refined isotropic. After refinement some residual electron density was found in a ΔF synthesis. It was impossible to interpret this density so it was decided to correct for it by using the option SQUEEZE in the program package PLATON.[20, 21] For compound (7) BF$_4^-$ was refined isotropic, the solvent molecule consists of two half occupied molecules which share their Cl-atoms through the centre of symmetry; the solvent molecule was refined isotropically and the H-atoms of the solvent were kept fixed with U=0.10 Å$^2$ the remainder of the H-atoms were restrained with fixed temperature factors U=0.10 Å$^2$.

Supporting information available: X-ray crystallographic files in CIF format for the structure determinations of complexes 1, 4, 5 and 7. This material is available free of charge via the Internet at http://pubs.acs.org
7.6 References


P-P ligands, dimethyl allyl complexes.


