DOI: 10.1002/ejic.200701070

Dynamic Behaviour of the [(Triphos)Rh(η^1 : η^2 -P₄RR')]ⁿ⁺ Complexes [Triphos = MeC(CH₂PPh₂)₃; R = H, Alkyl, Aryl; R' = Lone Pair, H, Me; n = 0, 1]: NMR and Computational Studies

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Keywords: Phosphorus / Rhodium / Tripodal phosphane ligands / Fluxionality / Density functional calculations / NMR spectroscopy

Solution multinuclear and multidimensional NMR analyses of the [(triphos)Rh(η^1 : η^2 -P₄RR')]ⁿ⁺ complex cations [triphos = MeC(CH₂PPh₂)₃; R = H, Me, Ph; R' = lone pair, H, Me; n = 0, 1] confirm the same primary structure determined by X-rays in the solid state. In addition, 2D ¹H NOESY and ³¹P{¹H} exchange NMR spectroscopy show that these complexes are nonrigid on the NMR time-scale over the 253–318 K temperature range. A dynamic process that involves the terminal phosphane groups of the triphos ligand is displayed by each compound. The NMR spectroscopic data indicate a slow

scrambling motion in which the P_4R unit tumbles with respect to the (triphos)Rh moiety. DFT calculations outline a possible turnstile mechanism involving the threefold and twofold rotors into which the complex is subdivided. The process goes through a transition state in which the axial and equatorial dispositions of the PRR' and P=P donating groups of the P_4RR' ligand are inverted with respect to the ground state.

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Introduction

White phosphorus is a cheap chemical largely used in the manufacturing of organophosphorus compounds.[1] These industrially important derivatives are generally produced by the reaction of PCl₃ or PCl₅ with suitable organic reagents. The industrial process is not very environmentally sustainable as the reaction involves gaseous chlorine and releases huge amounts of hydrochloric acid into the atmosphere. Therefore, the discovery of new and efficient methods for the preparation of organophosphorus derivatives from P₄ through non-chlorine routes remains a challenging task for the industry.^[2] Moreover, the underpinnings of the formation of P-C and P-H bonds need to be understood in more detail.^[2,3] Advances in this area entail the use of transition metal systems that are suitable for the activation of both the P₄ tetrahedron and the organic substrate. On the other hand, a recent breakthrough in the area has been presented by Bertrand et al. who have reported on the reactivity of alkyl amino carbenes with P4. This process does not lead to the formation of any metal intermediate and leads to very reactive P-C bonds.[4]

Another important advancement in P₄ activation/functionalization was our previous finding that some rhodium

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and iridium organometallic complexes containing the tripodal ligand, MeC(CH₂PPh₂)₃ (triphos), can activate the P₄ molecule and selectively deliver to it the hydride, [5] alkyl or aryl [6,7] ligand originally coordinated to the metal. Scheme 1 reports the highlights of this chemistry and, in particular, the reaction of [(triphos)MH₃] (M = Rh, Ir) and [(triphos)-RhR(C₂H₄)] (R = H, alkyl, aryl) with P₄ to yield [(triphos)-M(η^1 : η^2 -P₄R)] compounds featuring a P–H or a P–C bond. Direct alkylation of the rhodium derivatives affords cationic species of the [(triphos)Rh(η^1 : η^2 -P₄RR')]⁺ type, which can be isolated as solids. These were characterized in both the solid and solution states. [7]

These complexes are not only suitable organometallic platforms with which to study the transformation of P₄ into organophosphorus compounds,[8] but are also intriguing for their structural and bonding properties. In particular, because of their lack of molecular symmetry, these species exhibit a complicated ³¹P NMR splitting pattern resulting from the presence of eight different NMR-active nuclei (ABCDEFGX spin system). Polyphosphorus ligands usually exhibit interesting NMR properties and often show a fluxional behaviour, which can be related to different processes such as scrambling of coordinated ligands, [9] skeleton rearrangements^[10] and rotational interconversions.^[11] Since the triphos complexes formed by polyphosphorus units such as $cyclo-P_3^{[12]}$ or $cyclo-P_2X$ (X = S, Se, Te)^[13] exhibit an overall fluxionality due to both the triphos and the naked polyphosphorus unit, it is not surprising that the [(triphos)-





Scheme 1. Summary of the reactivity of [(triphos)MH₃] (M = Rh, Ir) and [(triphos)RhR(C_2H_4)] (R = H, Me, Et, Ph) with P₄.

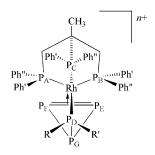
 $M(\eta^1:\eta^2-P_4R)]$ and [(triphos) $M(\eta^1:\eta^2-P_4RR')]^+$ complexes also lack stereochemical rigidity in solution. Remarkably, the occurrence of a fluxional behaviour in these species could not be evinced from the inspection of the room temperature spectra, but became evident from the observation of inverted-phase peaks in the 1H NOESY NMR spectra. This unexpected observation, which was also confirmed by variable-temperature NMR measurements, prompted us to study in detail the dynamic behaviour of these complexes. In this work, which combines experimental NMR spectroscopic data and DFT modelling, a mechanistic picture of the observed dynamic process is presented.

Results and Discussion

Dynamic Behaviour of the [(Triphos)Rh(η^1 : η^2 -P₄RR')]ⁿ⁺ Complexes in Solution: NMR Studies

The solid- and solution-state structures of the [(triphos)-Rh($\eta^1:\eta^2-P_4R$)] [R = H (1), Me (2), Et (3), Ph (4)] and [(triphos)Rh($\eta^1:\eta^2-P_4RR'$)](OTf) [R = H, R' = Me (5); R = Ph, R' = Me (6); OTf = OSO₂CF₃] Rh^I complexes were previously obtained from detailed multinuclear and multidimensional NMR analysis and, for 6, from X-ray diffraction. In both complexes, the same primary structure is retained.^[7]

As shown in Scheme 2, which also indicates the labelling scheme adopted for the NMR assignments, the coordination geometry around the metal is best described as trigonal bipyramidal (TBP) rather than pseudo-octahedral. In fact, the P_4RR' unit may be considered to be acting as a bidentate ligand in which a P=P double bond in the P_3 ring occupies an equatorial position and the PRR' σ -donor group resides in an axial position. The coordination sphere at the



R' [(triphos)Rh($\eta^1, \eta^2 - P_4H$)] Н 0 0 [(triphos)Rh(η^1 , η^2 –P₄Me)] 2 Me 3 0 [(triphos)Rh($\eta^1, \eta^2 - P_4Et$)] Et lpPh 0 [(triphos)Rh(η^1 , η^2 –P₄Ph)] lpMe H $[(triphos)Rh(\eta^1,\eta^2-P_4MeH)]^+$ [(triphos)Rh(η^1 , η^2 –P₄Ph)Me]⁺ Me 1

Scheme 2. Sketch of the [(triphos)Rh(η^1 : η^2 -P₄RR')]ⁿ⁺ complexes 1–6 showing the labelling scheme adopted for the NMR assignments; lp = lone pair.

rhodium centre is completed by the triphos P atoms occupying one axial and two equatorial positions. The bonding features of the [(triphos)M(η^1 : η^2 -P₄RR')]ⁿ⁺ (n=0,1) complexes compare well to those of the classic L₄M(η^2 -olefin) ones.^[7] In fact, the P=P double bond (localized at one side of the P₃ triangle) replaces the η^2 -coordinated olefin ligand. Four other P σ donors—three from the triphos ligand and one from the PRR' unit—confer the typical butterfly shape to the supporting metal fragment of the L₄M fragment (M = d⁸).

NMR experiments to determine the solution-state structure of this class of compounds were carried out by observing the ¹H, ³¹P and ¹³C signals over the 253-318 K temperature range in CD₂Cl₂ or [D₈]THF solutions. In each complex, all the ³¹P, ¹H and ¹³C nuclei are chemically nonequivalent because of the lack of any symmetry elements in the molecules. Only the protons of the terminal methyl group in triphos, as well as the *ortho* and *meta* proton pairs on the same phenyl ring, appear equivalent on the NMR time-scale; this is because of the fast rotation of the triphos methyl group and the phenyl groups around the C-C and C–P bonds, respectively. The phosphorus atoms of triphos in a pseudo-trans disposition to the P_E, P_F and P_D phosphorus atoms of the P₄RR' unit were labelled as P_A, P_B and P_C, respectively. A higher chemical shift was arbitrarily assigned to P_A than to P_B. All the ¹H, ³¹P and ¹³C NMR resonances in 1-6 (Table 1 and Table 2) were unambiguously assigned by a combination of 1D and 2D NMR techniques (see Exp. Sect.).^[7]

Table 1. $^{31}P\{^{1}H\}$ NMR chemical shifts for the [(triphos)Rh(η^{1} : η^{2} - $P_{4}RR'$)] $^{n+}$ complexes.[a].

Nucleus	Complex					
	1 ^[b]	2 ^[b]	3 ^[c]	4 ^[d]	5 ^[c]	6 ^[e,f]
$\overline{P_A}$	15.96	15.61	12.49	13.86	15.88	15.91
P_{B}	2.81	-1.94	-2.44	-12.03	9.75	2.56
$P_{\rm C}$	-10.78	-11.77	-11.42	-4.63	5.21	9.46
$P_{\rm D}$	-280.22	-214.08	-192.91	-213.44	-172.02	-156.75
$P_{\rm E}$	-180.12	-174.52	-178.24	-181.09	-190.11	-200.77
$P_{\rm F}$	-184.98	-198.79	-199.92	-201.39	-209.25	-174.04
P_{G}	-3.33	26.79	24.28	18.14	5.60	17.39

[a] 202.46 MHz, [D₈]THF, in ppm. [b] 293 K. [c] 253 K. [d] 266 K. [e] CD_2Cl_2 . [f] 275 K; data from ref. [7].

¹H NOESY experiments were carried out on all complexes in order to obtain their structural characterization in solution. The ¹H NOESY spectra showed that, under the adopted experimental conditions, the *extreme narrowing limit* ($\tau_c \omega_L << 1$) is attained by each compound. Indeed, with the exception of complex **5**, only positive (negatively phased) NOEs were observed at 253 K.^[14,15] *Remarkably, apart from providing relevant structural information, the ¹H NOESY experiments showed that all of the examined complexes are not rigid on the NMR time-scale*. In fact, strong positively phased ¹H exchange cross-peaks were observed, which is consistent with a slow exchange motional regime. In particular, the following selective exchange pathway for the *ortho, meta* and *para* protons of the triphos ligand was

Table 2. Selected ^{1}H NMR chemical shifts for the [(triphos)-Rh(η^{1} : η^{2} -P₄RR')]ⁿ⁺ complexes.^[a]

Complex	Proton	δ [b]
1 ^[c]	P_DH	0.01 (br. dd)
2 ^[c]	P_DCH_3 (3 H)	0.65 (br. s)
3 ^[d]	$P_DCH_2CH_3$ (3 H)	0.70 (q)
	P _D CH'	1.42
	P _D CH"	0.30
4 ^[e]	$o\text{-PhP}_D$ (2 H)	7.66 (br. d)
	o-Ph'P _A (2 H)	5.71 (t)
	o-Ph"P _A (2 H)	8.12 (t)
	o-Ph'P _B (2 H)	8.48 (t)
	o-Ph"P _B (2 H)	8.09 (t)
	o-Ph'P _C (2 H)	6.47 (t)
	o-Ph"P _C (2 H)	7.33 (t)
5 ^[e]	P_DCH_3 (3 H)	0.82 (br. s)
	P_DH	3.50 (br. d)
6 ^{[f][g]}	P_DCH_3 (3 H)	0.93 (br. d)
	o-PhP _D (2 H)	8.01 (t)
	o-Ph'P _A (2 H)	6.23 (dd)
	o-Ph"P _A (2 H)	8.50 (t)
	o-Ph'P _B (2 H)	8.21 (br.)
	o-Ph"P _B (2 H)	6.66 (t)
	o-Ph'P _C (2 H)	6.06 (t)
	o-Ph"P _C (2 H)	7.49 (t)

[a] 500.132 MHz, solvent [D₈]THF, signals are multiplets unless otherwise specified: s = singlet, d = doublet, t = triplet, q = quartet, br. = broad. [b] Shift values given in ppm. [c] 293 K. [d] 253 K. [e] 266 K. [f] CD₂Cl₂. [g] 275 K; data from ref. [7].

observed for 1–6 (see also Scheme 2): $Ph'P_A \leftrightarrow Ph'P_B \leftrightarrow Ph'P_C$ and $Ph''P_A \leftrightarrow Ph''P_B \leftrightarrow Ph''P_C$.

A representative example is reported in Figure 1, in which a section of the 1H NOESY spectrum of [(triphos)-Rh(η^1 : η^2 -P₄Ph)] (4) clearly displays the exchange peaks between the *ortho*-Ph'P_A and the *ortho*-Ph'P_B protons (P1) as well as between the *ortho*-Ph'P_A and the *ortho*-Ph'P_C protons (P2).

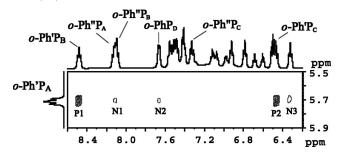


Figure 1. Section of the 1H NOESY spectrum of [(triphos)-Rh($\eta^1:\eta^2-P_4Ph)]$ (4) (500.13 MHz, [D_8]THF, 266 K, $\tau_m=1.0$ s): positively phased (exchange) cross-peaks are represented by filled circles and negatively phased (NOE) cross-peaks are represented by empty circles. Labels: P1 (o-Ph'P_A – o-Ph'P_B), P2 (o-Ph'P_A – o-Ph'P_C), N1 (o-Ph'P_A – o-Ph'P_A), N2 (o-Ph'P_A – o-PhP_D), N3 (o-Ph'P_A – m-Ph'P_A).

An analogous dynamic behaviour is displayed by the methylenic protons of the triphos chains. In some instances, the rate of the exchange process is faster than the rate of the NOE build-up. This is shown, for example, by the [(triphos)Rh(η^1 : η^2 -P₄PhMe)](OTf) (6) complex in CD₂Cl₂ solution at 275 K. In fact, the *ortho*-PhP_D protons show NOE contacts not only with the close *ortho*-Ph'P_A protons but

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also with all the other *ortho* protons exchanging with those of Ph'P_A, namely, the *ortho*-Ph'P_B and *ortho*-Ph'P_C protons (Figure 2).

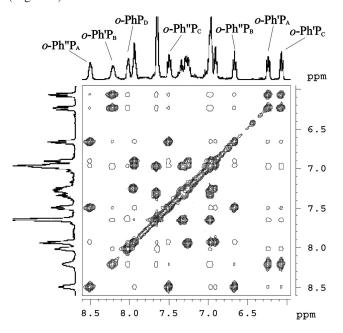


Figure 2. Section of the 1H NOESY spectrum of [(triphos)-Rh(η^1 : η^2 -P₄PhMe)] (OTf) (6) in the aromatic region (500.13 MHz, CD₂Cl₂, 275 K, $\tau_m = 0.8$ s). Positively phased (exchange) crosspeaks are represented by filled circles and negatively phased (NOE) cross-peaks are represented by empty circles.

The presence of a dynamic process in both [(triphos)-Rh(η^1 : η^2 -P₄R)] and [(triphos)Rh(η^1 : η^2 -P₄RR')](OTf) complexes was confirmed by 2D ³¹P{¹H} EXSY spectroscopy. [16] All ³¹P{¹H} EXSY spectra are consistent with

the exchange of the phosphorus atoms of triphos. No exchange peaks related to the phosphorus atoms of the P_4 unit were detected. A section of the $^{31}P\{^1H\}$ EXSY spectrum of $\bf 6$ is reported in Figure 3 as an example showing the complete network of exchange peaks involving the three triphos P atoms.

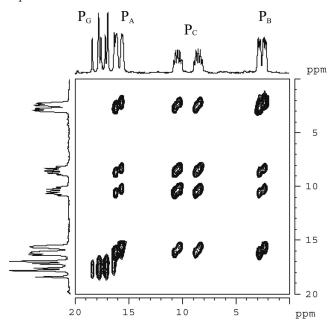


Figure 3. Section of the $^{31}P\{^{1}H\}$ EXSY spectrum of [(triphos)-Rh(η^{1} : η^{2} -P₄PhMe)](OTf) (6) (202.47 MHz, CD₂Cl₂, 275 K, τ_{m} = 0.1 s).

The only motion which may account for the observed dynamic behaviour in 1-6 is a slow scrambling of the P_4RR' and (triphos)Rh units with respect to each other. A

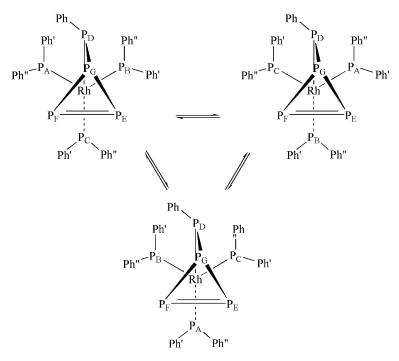


Figure 4. Schematic representation of the dynamic behaviour of [(triphos)Rh(η^1 : η^2 -P₄Ph)] (4) viewed along the Rh–C(CH₃) axis.

sketch of the proposed motion undergone by these systems is schematically represented in Figure 4 for the neutral [(triphos)Rh(η^1 : η^2 -P₄Ph)] complex (4). The process involves the interconversion of three equivalent rotamers by the switching of the phosphorus atom of triphos occupying the *trans*-axial position to the P_D atom position of the P₄Ph unit, which is always axial.

To arrive at a better understanding of the observed fluxionality, variable-temperature ¹H and ³¹P{¹H} NMR spectra were recorded for representative complexes over the 273–318 K temperature range in CD₂Cl₂ solutions. Unfortunately, both because of the decomposition of the complexes at higher temperatures and the lack of appropriate computer programs for the simulation of eight-nuclei spin systems, the activation parameters for the dynamic process could not be determined from conventional line-shape analysis. However, both proton and phosphorus NMR signals show significant temperature-dependent changes. As an example, the ¹H and ³¹P{¹H} NMR spectra in the aromatic and triphos regions for the [(triphos)Rh(η^1 : η^2 -P₄PhMe)]⁺ complex (6) are reported in Figure 5, respectively. In both spectra, the resonances due to the P₄PhMe unit (i.e., P_G and o-PhP_D) sharpen upon increasing the temperature, whereas the resonances due to the triphos nuclei (i.e., PA, PB, PC and their ortho protons) broaden and shift toward the coalescence point. These findings are

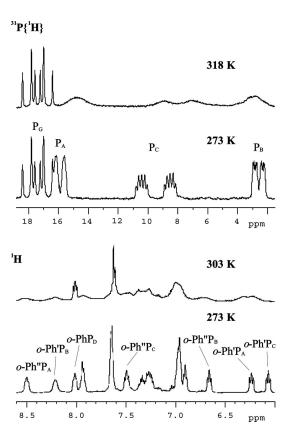


Figure 5. Sections of the variable-temperature NMR spectra of $[(triphos)Rh(\eta^1:\eta^2-P_4PhMe)](OTf)$ (6) recorded at 273 and 318 K (CD₂Cl₂). Top: $^{31}P\{^1H\}$ spectra (202.47 MHz). Bottom: 1H spectra (500.132 MHz).

in line with a rotational exchange involving the triphos nuclei only, as already indicated by the 2D NMR experiments.

Scrambling Mechanism and DFT (Density Functional Theory) Modelling of the Exchange Process

Different mechanisms to account for the relative motion of the (triphos)Rh and P₄RR' units in compounds 1–6 can be proposed on the basis of the results of the NMR studies. The dynamic behaviour of the known (triphos) $M(\eta^3-P_3)$ complexes, [12,13,17] which contain the cyclo-P₃ adduct, as well as that of the related carbonyl derivatives—[(triphos)- $M\{\eta^3-P_3[M'(CO)_5]_x\}$] (M = Co, M' = Cr, x = 1-3;[17a,18,19] $M' = Mo, W, x = 1,^{[19]} 2,^{[18]} M = Rh, M' = W, x = 1,^{[19]} M$ $[(triphos)Rh\{\eta^3-P_3[W(CO)_4(PPh_3)]\}],^{[19]}$ and $[(triphos)-P_3[W(CO)_4(PPh_3)]]$ $Co\{\eta^3-P_3[Re(CO)_5]\}BF_4$, [19]—and the intrinsic threefold symmetry of the (triphos)M unit suggest an exchange process involving trihapto-P3 triangle intermediates which encompass a $\eta^1:\eta^2 \leftrightarrow \eta^3 \leftrightarrow rotation \leftrightarrow \eta^3 \leftrightarrow \eta^1:\eta^2$ sequence, as sketched in Scheme 3. Indeed, easy scrambling of the substituted (triphos) $M(\eta^3-P_3)$ species in solution has been previously reported for the metal carbonyl adducts, $[(triphos)M\{\eta^3-P_3[M'(CO)_4(L)]_x\}]Y_x^{[18,19]}$ and for the dimetallic rhodium–ruthenium species, {[(triphos)Rh](μ,η^3 :1- P_3 [CpRu(CH₃CN)(PR₃)] PF₆ (R = Me, Ph, Cy). [20]

$$\begin{array}{c} \begin{array}{c} R & R' \\ P_{B} & P_{D} \\ P_{C} & P_{F} \end{array} \end{array} \qquad \begin{array}{c} \begin{array}{c} R & R' \\ P_{B} & P_{D} \\ P_{C} & P_{F} \end{array} \end{array} \qquad \begin{array}{c} \begin{array}{c} P_{B} & P_{E} \\ P_{D} & P_{E} \end{array} \end{array} \qquad \begin{array}{c} \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \end{array} \qquad \begin{array}{c} \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \end{array} \qquad \begin{array}{c} \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \end{array} \qquad \begin{array}{c} \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \end{array} \qquad \begin{array}{c} \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \end{array} \qquad \begin{array}{c} \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \end{array} \qquad \begin{array}{c} \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{B} & P_{E} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c} P_{C} & P_{D} \\ P_{C} & P_{D} \end{array} \qquad \begin{array}{c$$

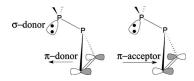
Scheme 3.

Although well-documented in the literature [18-20] and attractive for its intrinsic simplicity, the above mechanism seems unlikely in the present case, as it would require several P-P bond-forming/bond-breaking steps within the P_4RR' unit. This also implies a continuous change of the



metal coordination mode and an energetically expensive sequence of reductive elimination and oxidative addition reactions. Moreover, the trihapto coordination of the ligand would be structurally and electronically inconsistent with a P_D atom substituted by R and R' groups. [21]

If one rules out the mechanism in Scheme 3, only a turn-stile process can be invoked. [22] Indeed, the (triphos)Rh unit defines a threefold rotor with respect to the (η^1 : η^2 - P_4RR') unit. The turnstile mechanism involves the relative rotation of these two units with respect to each other while maintaining a η^1 : η^2 coordination. In this case, neither cleavage of P–P bonds within the P_4 cage nor a change in the metal oxidation state is required for the rearrangement of the pentacoordinated species. As previously reported, [7] the P_4RR' ligand is able to donate one σ and one π electron pair (Scheme 4). Because of the hybridization of the d_π orbital of the L_4M fragment, [23] the back-donation into the P=P π^* level is greatly favoured when the P=P double bond lies in the equatorial plane of the TBP.



Scheme 4. Orbital origin of the donor and acceptor capabilities of the P_4RR' ligands.

The turnstile mechanism requires that the positions of the σ and π electron pairs are switched at some point; that is, the P=P bond must also be able to occupy the TBP apical position during the interconversion process. The feasibility of such a structural rearrangement is confirmed by the DFT optimization on a model of complex 1 (R = H, R' = lone pair)—namely, [(Htriphos)Rh(η^1 : η^2 -P₄H)]—in which Htriphos is a theoretical counterpart of the real tripodal phosphane with all the phenyl substituents in the ligand replaced with H atoms.^[7] This model compound is characterized by an actual transition state (one negative frequency for structure **b** in Figure 6), which lies about 20 kcal mol⁻¹ above the ground state (structure **a** in Figure 6).

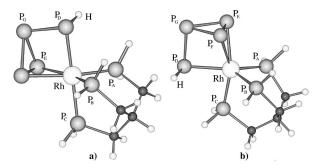


Figure 6. DFT-optimized geometries of the ground state (a) and transition state (b) of the model compound, [(Htriphos)Rh($\eta^1:\eta^2-P_4H$)].

The significantly larger stability of the latter structure is due to the metal π back-donation, which occurs when the P=P bond lies in an equatorial position of the trigonal bipy-

ramidal structure; this is reduced in the apical position. As shown by the distances reported in Table 3, the different conformation has an obvious effect on the geometrical parameters. Thus, in model b the P=P distance is shorter than in a (2.14 vs. 2.16 Å, respectively) while the Rh-P_E and Rh- $P_{\rm F}$ bonds are longer (2.53 and 2.52 vs. 2.48 and 2.49 Å, respectively). The relatively high energy barrier for the rearrangement of the whole P₄R unit with respect to the (triphos)Rh fragment justifies the slow dynamic exchange process observed under the experimental NMR conditions. In fact, the ¹H NOESY and the ³¹P{¹H} EXSY spectra are consistent with a slow scrambling of the P₄R and (triphos)-Rh units with respect to each other. In conclusion, the reciprocal rearrangement of the bulky threefold and twofold rotors—that is, (triphos)Rh and Rh(η^1 : η^2 -P₄RR')—must be significantly hindered. The situation is fairly similar to $L_4M(\eta^2$ -olefin) complexes for which comparable rotation barriers are found.[24]

Table 3. Selected bond lengths [Å] of the optimized geometries of the ground state (a) and transition state (b) of the model compound, $[(Htriphos)Rh(\eta^1:\eta^2-P_4H)]$.

	Model a	Model b
Rh-P _A	2.35	2.35
Rh-P _B	2.36	2.36
Rh-P _C	2.41	2.30
Rh-P _D	2.43	2.51
Rh-P _E	2.48	2.52
Rh-P _F	2.49	2.53
P _D -H	1.43	1.43
$P_D - P_G$	2.26	2.26
$P_E - P_E$	2.16	2.14
$P_E - P_G$	2.26	2.25
$P_F - P_G$	2.27	2.25

Conclusions

A detailed multinuclear, multidimensional NMR investigation of both neutral $\eta^1:\eta^2-P_4R$ and cationic $\eta^1:\eta^2-P_4RR'$ (triphos)Rh complexes of the general formula [(triphos)- $Rh(\eta^1:\eta^2-P_4RR')]Y_n$ has allowed for a satisfactory structural characterization of the species in solution. Moreover, the fluxional behaviour, which entails the exchange of the triphos phosphorus atoms, has been highlighted. The conclusions are supported by DFT calculations for the [(Htriphos)Rh(η^1 : η^2 -P₄H)] model compound. For this compound, a transition state is detected about 20 kcal mol⁻¹ above the ground state and this provides a convincing justification for a turnstile pseudorotation process. This implies that the P=P bond, which belongs to the P₃ triangular face of the P₄RR' ligand and is coordinated to the metal in a dihapto fashion, switches between the equatorial and axial positions of the bipyramidal geometry. Given the significant energy barrier, the process is hindered and slow, which is fully consistent with the NMR response.

Experimental Section

General Data: The [(triphos)Rh(η^1 : η^2 -P₄RR')]Y [triphos = MeC(CH₂PPh₂)₃; R = H, alkyl, aryl; R' = lone pair, H, Me] complexes were prepared as previously reported.^[7]

NMR Experiments: ³¹P{¹H}, ¹H and ¹³C{¹H} NMR spectra were recorded with a Bruker Avance DRX-500 spectrometer equipped with a variable-temperature control unit accurate to ± 0.1 °C and operating at 202.47, 500.13 and 125.76 MHz, respectively; deuterated solvents dried with molecular sieves (4 Å) were used. ³¹P{¹H} chemical shifts are reported relative to external 85% H₃PO₄, with downfield values reported as positive. ¹H and ¹³C{¹H} chemical shifts are reported relative to tetramethylsilane as an external reference and were calibrated against the residual solvent resonance (1H) or the deuterated solvent multiplet (13C). The assignments of the signals resulted from 1D spectra, 2D ¹H DQF-COSY, ³¹P{¹H} COSY, ¹H NOESY and proton-detected 2D ¹H-¹³C and ¹H-³¹P correlations. 2D NMR spectra were recorded with degassed nonspinning samples by using pulse sequences suitable for phase-sensitive representations and by using TPPI. Standard pulse sequences were used for the ¹H NOESY^[14] experiments: 1024 increments 2 K in size (with 8 scans each) and covering the full range in both dimensions (ca. 5000 Hz) were acquired with a relaxation delay of 2 s and a mixing time of 1.0 or 0.8 s. The 2D ${}^{31}P{}^{1}H{}^{1}$ EXSY[16] spectra were recorded using the NOESY pulse sequence^[14b,25] with ¹H decoupling during acquisition: 1024 increments 2 K in size (with 64 scans each) and covering the full range in both dimensions were collected with a relaxation delay of 0.8 s and mixing times of 0.10, 0.50, 0.80 and 1.00 s.

Computational Studies: Structural optimizations were carried out at the hybrid density functional theory (DFT) using the Gaussian98 program.^[26] The triphos ligand was modelled by replacing all of the phenyl substituents with H atoms (Htriphos). This allows for the maintenance of the typical geometric constraints at the metal. In particular, the three P-Rh-P angles are forced to be almost 90°. The method used was the Becke three-parameter hybrid exchangecorrelation functional containing the Lee, Yang and Parr nonlocal gradient correction (B3LYP).[27,28] The nature of the optimized structures was confirmed by calculation of the frequencies. The basis set for the Rh atom utilized the effective core potentials of Hay and Wadt with the associated double-ζ valence basis functions.^[29] The basis set used for the remaining atomic species was 6-31G(d,p).[30] The geometrical parameters for the minimum structure of model a (see text above) were compared with the X-ray solid-state features of [(triphos)Rh(η¹:η²-P₄PhMe')](OTf)^[7] and indicate a good agreement between the two structures.

Acknowledgments

The authors are thankful for the support of the European Community through the FP6 IDECAT Network of Excellence (contract NMP3-CT-2005-011730). AI and CM are grateful for the computing time provided by CINECA under the agreement of the National Council for Research (CNR). Thanks are also given to Thermphos Intl., Vlissingen, NL for a generous loan of white phosphorus and for supporting this research activity.

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Published Online: January 15, 2008