

Phosphabenzene–rhodium catalysts for the efficient hydroformylation of terminal and internal olefins

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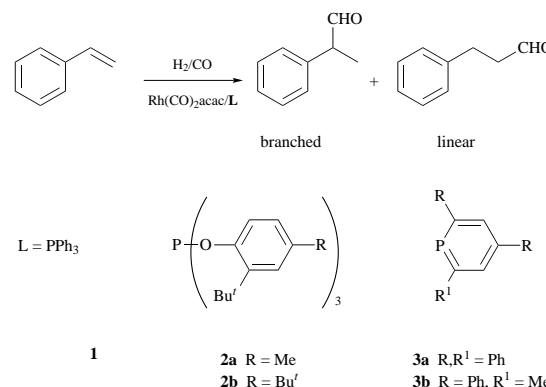
The use of rhodium-catalysts modified with bulky phosphabenzene as π -acceptor ligands as highly efficient hydroformylation catalysts for terminal and internal olefins is reported, and a catalytically active phosphabenzene complex has been structurally characterized by X-ray crystallography.

Since its discovery in 1938 hydroformylation of olefins has evolved into one of the industrially most important homogeneously catalyzed processes.¹ In 1996 worldwide about 7 million tons of oxo products were produced, resulting in a tremendous economical interest to run this reaction with a maximum of activity and selectivity.² A problem still not satisfactorily solved is that of hydroformylation of internal olefins, which is of considerable interest in both industrial and synthetic organic contexts. The main difficulty stems from a low catalyst activity of the standard triphenylphosphine (**1**)-rhodium catalysts for the hydroformylation of internal olefins. Substitution of phosphines **1** by bulky phosphites **2** as modifying ligands for rhodium catalysts was an important advance in this field.³ While the catalytic activity of such systems is extremely high, phosphites in general suffer from an inherent lability towards hydrolysis and a tendency to undergo degradation reactions. This makes the development of new catalysts for the efficient hydroformylation of internal olefins an important task.

We recently introduced new classes of strong π -acceptor ligands for homogeneous catalysis that could have beneficial effects in hydroformylation chemistry.⁴ We report here on the basis of those exploratory results on the development of rhodium catalysts modified with new bulky monophosphabenzene **3** for the efficient hydroformylation of terminal and internal olefins, as well as on the preparation and X-ray crystal structure analysis of a defined monophosphabenzene–rhodium complex which is endowed with high catalytic activity.

The hydroformylation of styrene was chosen as a first test reaction, to compare the catalytic activity of phosphabenzene **3**† catalysts with that of phosphine **1** and phosphite **2** containing systems (Scheme 1, Table 1). For this particular reaction a combination of high activity and regioselectivity favoring the branched aldehyde is desirable with regard to the preparation of the anti-inflammatory 2-arylpropionic acids.⁵ Employing extremely mild hydroformylation conditions (room temp., 20 bar CO–H₂, 1:1) the phosphabenzene catalysts showed the highest activity providing the desired 2-phenylpropanal in a regioselectivity of 20:1.

To study the catalytic activity of the monophosphabenzene rhodium catalyst for hydroformylation of internal olefins we turned to cyclohexene as a model substrate. Here the rhodium catalyst modified with the *o,o'*-diphenyl substituted monophosphabenzene **3a** showed 74 times higher activity than the standard PPh₃-rhodium catalyst for hydroformylation of cyclohexene (Table 2). Thus, this catalyst is as reactive as the most active hydroformylation catalysts known to date, the



Scheme 1 Reagents and conditions: 0.357 mol% [Rh(CO)₂acac/5 L], CO–H₂ (1:1), 20 bar, 25 °C, toluene

Table 1 Results of regioselective hydroformylation of styrene

Entry ^a	Ligand	t/min	Conv. (%) ^b	TOF/h ^{-1c}	b:l ^b
1	PPh ₃	180	8	7.5	24:1
2	2b	180	17.6	16.4	20:1
3	3a	180	30.8	28.7	20:1

^a Reactions were carried out at 25 °C in toluene ([styrene]₀ = 0.65 mol l⁻¹) in a 100 ml stainless-steel autoclave under an atmosphere of H₂ and CO (1:1) 20 bar initial total pressure, Rh:L:styrene = 1:5:280. ^b Conversions and branched to linear (b:l) ratios were determined by GC on a 0.25 mm × 30 m Supelcowax column with internal standard. For catalysts with ligands **2b** and **3b** quantitative conversion could be reached after 22 h; b:l ratio after complete conversion stays at 20:1. In all cases aldehyde selectivity was 100%. ^c Turn-over frequency (TOF) was calculated as (mol reacted substrate) × (mol catalyst)⁻¹ × (t/h)⁻¹.

bulky monophosphite **2** modified rhodium systems. Interestingly the rhodium–phosphabenzene catalysts could be reused at least two more times to catalyze this reaction without loss in catalytic activity indicating their stability.

In agreement with our initial concept⁴ that catalyst activity in hydroformylation increases by coordination of a strong π -acceptor ligand to rhodium, it seemed reasonable to attribute this catalyst's activity to an η^1 -coordination of the phosphabenzene nucleus to the rhodium(i) center. Since no structural data on that kind of coordination of a phosphabenzene to rhodium was available⁶ we prepared a defined rhodium(i) phosphabenzene complex by reacting Rh(1,5-cod)₂BF₄ (cod = cycloocta-1,5-diene) with 4 equiv. of the phosphabenzene **3b**. This afforded the phosphabenzene rhodium complex **4** as orange–red crystals in almost quantitative yield.‡ X-Ray

‡ Preparation and selected physical data of Rh(**3b**)₄BF₄ (**4**): to a magnetically stirred solution of 23 mg (0.057 mmol) of Rh(1,5-cod)₂BF₄ in 2 ml CDCl₃ were added at room temp. 60 mg (0.228 mmol) of phosphabenzene **3b**. The reaction mixture was heated to reflux for 15 min, cooled to room temp. and filtered over Celite with 20 ml of CH₂Cl₂. The solvent was removed *in vacuo* to give 67 mg (95%) of **4** as an orange solid. **4**: δ_p (161.978 MHz, C₆D₆) +170.4 (d, ¹J_{P, Rh} 154.4 Hz); *m/z* (FD) 1151 [Rh(C₁₈H₁₅P)₄], 262 (C₁₈H₁₅P).

† The phosphabenzene **3a, b** have been prepared from the corresponding pyrylium salts and P(SiMe₃)₃ according to Märkl *et al.*⁸

Table 2 Results of hydroformylation of cyclohexene

Entry	Ligand	L/Rh	t/h	Conv. (%) ^c	TOF/h ^{-1d}
1 ^a	1	20	73	28.1	2.9
2 ^a	2b	10	1	28.8	216
3 ^a	3a	10	1	28.6	214
4 ^b	4	4	1	18	127
5 ^b	3b	4	1	12.2	92
6 ^b	3b	10	1	7	53

^a Reactions were carried out at 90 °C in toluene ([cyclohexene]₀ = 0.68 mol l⁻¹) in a 100 ml stainless-steel autoclave under an atmosphere of H₂ and CO (1:1) at 20 bar initial total pressure. Rh: cyclohexene = 1:750. ^b As described under (^a) except [cyclohexene]₀ = 0.798 mol l⁻¹ in toluene. ^c Conversions were determined by GC [see footnote (^b) Table 1]. Quantitative conversion could be reached for all rhodium catalysts with ligands **2** and **3** at longer reaction times. In all cases aldehyde selectivity was 100%. ^d TOF is calculated as (mol reacted substrate) × (mol catalyst)⁻¹ × (t/h)⁻¹.

structure analysis revealed that four phosphabenzene nuclei are η¹-coordinated through the lone pair at phosphorus to the cationic rhodium(i) center in a square planar coordination geometry (see Fig. 1). The planes of the phosphabenzene nuclei are twisted between 48 and 51.1° from the plane defined by P(1)P(2)P(3)P(4). Interestingly the phosphabenzene nucleus of each ligand shows π-stacking with the *o*-phenyl substituent of one neighbouring phosphabenzene ligand and *vice versa*. This twofold attractive π-stacking interaction causes a narrowing of the P(1)–Rh–P(4) and P(2)–Rh–P(3) bond angles to 87° each between the interacting phosphabenzene ligands. Consequently, the remaining P–Rh–P angles between the non-interacting phosphabenzene nuclei are widened to 93°.

In a catalytically active rhodium hydroformylation catalyst no more than two P-donors may be coordinated to the rhodium center.² In the case of the bulky monophosphite ligands **2** only a single phosphite ligand is coordinated to rhodium within the catalytically active species.⁷ Accordingly, a complex such as **4** has to lose at least two phosphabenzene ligands to generate the catalytically active species. To find out whether a coordination of four phosphabenzene ligands to rhodium(i) is reversible or rather a thermodynamic sink, we subjected complex **4** to hydroformylation conditions. Under the conditions indicated in Table 2 (entry 4) complex **4** showed a remarkable catalytic activity in the hydroformylation of cyclohexene. A catalyst prepared *in situ* from Rh(CO)₂acac (Hacac = acetylacetonate) and the phosphabenzene **3b** performed with a comparable reactivity (entries

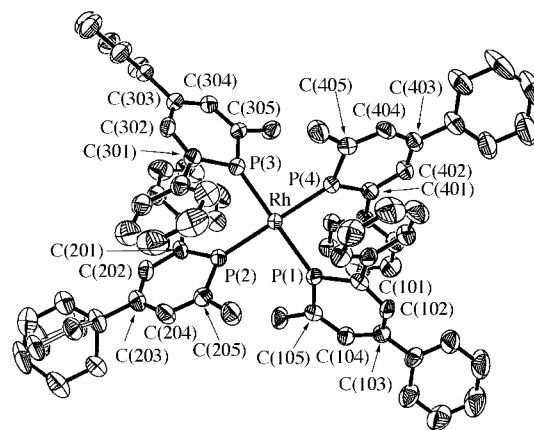


Fig. 1 Structure of the cation of **4** in the solid state; selected bond lengths (Å) and angles (°): Rh–P1 2.2918(11), Rh–P2 2.2661(11), Rh–P3 2.2628(11), Rh–P4 2.2721(11), P(3)–Rh–P(2) 86.85(4), P(2)–Rh–P(1) 93.55(4), P(3)–Rh–P(4) 92.99(4), P(4)–Rh–P(1) 86.60(4)

4 and 5). These results indicate two things: first, the coordination of at least two phosphabenzene nuclei to the rhodium(i) center of **4** is reversible although it remains unclear whether one or two phosphabenzene ligands stay coordinated within the catalytically active species. Second, η¹-coordinated phosphabenzene ligands to a rhodium(i) transition metal center can be regarded as the origin of the observed high catalyst activity.

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§ Crystal data for **4**: crystals suitable for X-ray crystal structure analysis were obtained from a CHCl₃–C₆H₆ solution at room temperature. [C₇₂H₆₀P₄Rh]⁺[BF₄]⁻·CHCl₃·2C₆H₆, *M* = 1514.38; monoclinic, space group *P*2₁/*c*, *a* = 18.666(3), *b* = 19.977(2), *c* = 21.253(3) Å, β = 107.96(1)°, *U* = 7539(2) Å³, *D*_c = 1.334 g cm⁻³ for *Z* = 4, *F*(000) = 3120, (Mo–Kα) = 0.472 mm⁻¹, λ = 0.710 73 Å, *T* = 213(2) K, crystal size 0.3 × 0.3 × 0.2 mm. A total of 13 684 reflections were collected (Enraf-Nonius CAD4) using scans in the range 4.6 < 2θ < 50°, 13 252 were unique (*R*_{int} = 0.0296). The structure was solved by direct methods. Full matrix least squares refinement was based on *F*², with all non-hydrogen atoms anisotropic and with hydrogens included in calculated positions with isotropic temperature factors 1.2 times that of the *U*_{eq} of the atom to which they were bonded.

The BF₄⁻ anion, one phenyl ring of the cation, and the solvent molecules (CHCl₃ and two molecules benzene in the asymmetric unit) were disordered. The 'calc squeeze' procedure⁹ in the program PLATON¹⁰ has been applied to the heavily disordered solvent molecules. The refinement converged at *R*1 = 0.0527 [for 8698 reflection with *I* > 2σ(*I*)] and *wR*2 = 0.1604 (all data) {*w* = [σ²(*F*_o)² + (0.0940*P*)²]⁻¹ where *P* = (*F*_o² + 2*F*_c²)/3}; final GOF = 1.034; largest peak and hole in the final difference fourier: 0.479/–0.450 e Å⁻³.

Programs used were SHELXS-96,¹¹ SHELXL-96¹² and PLATON.¹⁰ Atomic coordinates, bond lengths and angles, and thermal parameters have been deposited at the Cambridge Crystallographic Data Centre (CCDC). See Instructions for Authors, *J. Chem. Soc., Perkin Trans. 1*, 1997, Issue 1. Any request to the CCDC for this material should quote the full literature citation and the reference number 207/138.