

The complex (3) is extremely stable (*e.g.* solutions prepared in air show no evidence of decomposition) and all attempts to induce reductive elimination of the C–O bond to yield (4a) have failed. This observation and the fact that lactones arise exclusively from *trans*-addition of O–H to C=C in our catalytic systems^{3a} lead us to believe that species such as (3) are not directly involved in the catalytic cycle^{3b} for reaction (1). Although examples of mononuclear σ -alkenyl complexes arising from net *trans* insertion of activated alkynes into M–H bonds exist,⁷ in general, the alkyne insertion reaction is believed⁸ to occur in a *cis*-fashion analogous to that found for alkene insertions.

The complex (3) reacts cleanly with $\text{Me}_3\text{O}^+\text{BF}_4^-$ via alkylation at the carbonyl oxygen yielding the cationic complex (5) in high yield.[†] The CO stretching band at 1664 cm^{-1} (KBr) is shifted to higher frequency than in (3), compatible with a co-ordinated ester function,⁹ whereas the C=CH₂ stretching band (1585 cm^{-1}) is unchanged. The ¹³C{¹H} n.m.r. spectrum (CD₂Cl₂) shows a broad doublet at δ 182.32 (*J* 2 Hz) for the carbonyl group, a doublet of quartets at δ 156.61 (*J*_{Rh–C} 28, *J*_{P–C} 8 Hz) for Rh–C_α, and a multiplet at δ 120.87 for the =CH₂ group. A singlet at δ 55.34 is assigned to OMe. The molecular structure[‡] of (5) is shown in Figure 2.

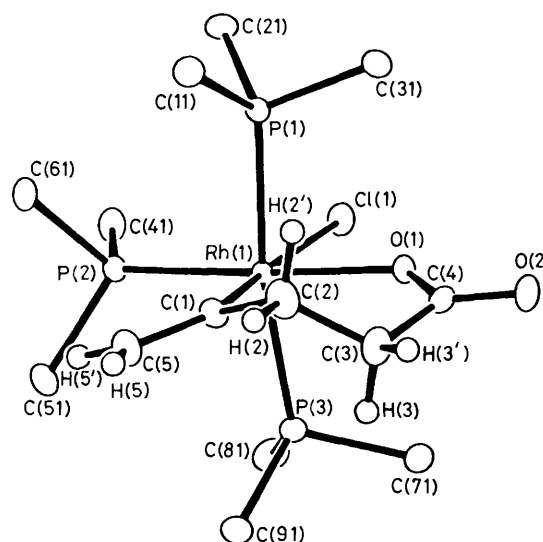


Figure 1. ORTEP drawing of a molecule of (3). Selected bond distances (Å) and angles (°): Rh(1)–P(1) 2.350(1); Rh(1)–P(2) 2.270(2); Rh(1)–P(3) 2.348(1); Rh(1)–O(1) 2.108(3); Rh(1)–C(1) 2.043(4); O(1)–C(4) 1.278(5); O(2)–C(4) 1.237(5); C(1)–C(2) 1.513(6); C(1)–C(5) 1.312(6); O(1)–Rh(1)–C(1) 91.2(1); Rh(1)–O(1)–C(4) 132.6(3); Rh(1)–C(1)–C(2) 114.8(3); Rh(1)–C(1)–C(5) 130.0(3).

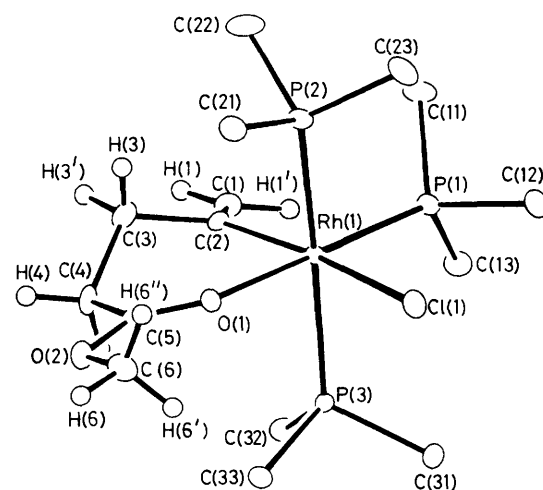


Figure 2. ORTEP drawing of a molecule of (5). Selected bond distances (Å) and angles (°): Rh(1)–P(1) 2.260(1); Rh(1)–P(2) 2.359(1); Rh(1)–P(3) 2.370(1); Rh(1)–O(1) 2.171(3); Rh(1)–C(2) 2.053(4); O(1)–C(5) 1.220(5); O(2)–C(5) 1.328(5); O(2)–C(6) 1.459(5); C(1)–C(2) 1.318(6); C(2)–C(3) 1.514(6); Rh(1)–C(2)–C(1) 128.8(3); Rh(1)–C(2)–C(3) 114.3(3); Rh(1)–O(1)–C(5) 128.2(3); O(1)–C(5)–C(4) 125.1(4).

The structures of (3) and (5) allow direct comparison of co-ordinated carboxylate and ester functions. The Rh–O(1) bond in (3) [2.108(3) Å] is significantly shorter than Rh–O(1) in (5) [2.171(3) Å], whereas C(4)–O(1) in (3) [1.278(5) Å] is longer than C(5)–O(1) in (5) [1.220(5) Å]; the C(4)–O(2) double bond in (3) [1.237(5) Å] is much shorter than the C(5)–O(2) single bond in (5) [1.328(5) Å]. These differences are consistent with the valence bond representations in Scheme 1. The Rh–C_α distances [Rh–C(1) in (3) is 2.043(4) Å and Rh–C(2) in (5) is 2.053(4) Å] are quite similar. As

expected, the longer Rh–O bond in (5) vs. (3) allows for displacement of the co-ordinated ester function by nucleophiles such as PMe_3 , as indicated by *in situ* ^1H and ^{31}P n.m.r. spectroscopy.

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