argument that in TBP complexes the π -back-bonding with e' d orbitals is stronger than that with e'' d orbitals, due to the favorable mixing of equatorial 4p orbitals of the metal.¹² We believe that this discrepancy is a consequence of incomplete geometry optimizations: since the geometry of the free olefin corresponds more closely to that of the weakly complexated one, it precludes an adequate description of the forward- and back-bonding of the π orbitals. This omission has a potential for reversing the already close energy levels of the two conformations.

By the same token, comparison with Zeise's anion, $[Pt(C_2-H_4)Cl_3]^-$, made by Grima et al. 16,26 to the effect that the geometry of the olefin should not change much because of complexation is not entirely justified. In this anion, the platinum atom has an approximate D_{4h} coordination, in which the p orbitals belong to symmetry types a_{2u} and e_u , different from the symmetry types of d orbitals $(a_{1g}, b_{1g}, b_{2g}, and e_g)$. This hinders the 4p-orbital

enhancement of the π -back-bonding, ¹² which is symmetry-allowed in the D_{3h} coordination and which our calculations confirm.

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Supplementary Material Available: Bond lengths and bond angles (Table I) and frontier orbitals (Table III) of the closed-shell configurations of CHO-Co(CO)₃, bond lengths and bond angles (Table V) and frontier orbitals (Table VII) of the closed-shell configurations of HCo(CO)₃-C₂H₄, bond lengths, bond angles, and SCF energies of the conformations of CH₃CH₂Co(CO)₃ (Table VIII), CI energies of the closed-shell configurations of HCo(CO)₃-C₂H₄ and CH₃CH₂Co(CO)₃ (Table IX), summary of the results on ethylene ¹A_g (3-21G basis) (Table 10), and density difference of the CC (Figure 30) and EtC (Figure 31) conformations of HCo(CO)₃-C₂H₄ in the symmetry plane (25 pages). Ordering information is given on any current masthead page.

Communications to the Editor

Molybdenum and Tungsten Derivatives of 2,6-Diphenylphenoxide: Chelation via Metal-Carbon σ -Bonds or Metal- π -Arene Interactions

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The early transition-metal chemistry associated with 2,6-dialkylphenoxide and 2,6-dialkylthiophenoxide ligation has recently received considerable research interest.²⁻⁶ Following our studies

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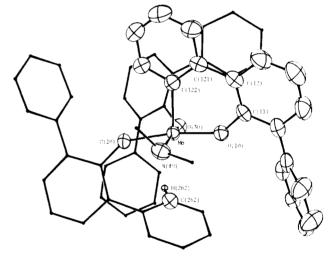


Figure 1. ORTEP view of 1 emphasizing the central coordination sphere and metallacycle ring. Selected bond distances (Å) and angles (°) are are follows: Mo-O(10), 1.919 (2); Mo-O(20), 1.976 (2); Mo-O(30), 1.962 (3); Mo-N(40), 2.288 (4); Mo-C(122), 2.075 (4); Mo-C(262), 2.814 (5); Mo-H(262), 2.67 (5); O(10)-Mo-O(20), 165.8 (1); -O(30), 97.9 (1); -N(40), 87.3 (1); -C(122), 86.4 (1); C(122)-Mo-C(262), 174.5 (1), -H(262), 164 (1).

of the sometimes mild intramolecular activation of the aliphatic carbon-hydrogen bonds in 2,6-di-*tert*-butylphenoxide ligands,⁷ we have turned our attention to the organometallic chemistry associated with 2,6-diphenylphenoxide groups.⁸ This ligand also

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Figure 2. ORTEP view of 3a emphasizing the central coordination sphere. Selected bond distances (Å) and angles (°) are as follows: Mo-P(1), 2.45 (2); -P(2), 2.453 (2); -O(1), 2.164 (5); -C(61), 2.262 (7); -C(62), 2.329 (7); -C(63), 2.255 (6); -C(64), 2.313 (7); -C(65), 2.301 (8); -C(66), 2.238 (7); Mo-H, 1.164 (8); Mo-O(1)-C(1), 118.0 (4).

offers the potential for metallacycle formation albeit via activation of an aromatic CH bond. We communicate here our initial findings concerning the coordination of this ligand to the heavier group 6 metals that indicate not only that formation of new metal-carbon σ -bonds is possible but that ground-state metalarene interactions can also be present.

The addition of 2,6-diphenylphenol (HOAr) to the deep purple $Mo(NMe_2)_4$ in benzene results in the formation of a deep red, crystalline precipitate 1 after several hours. Microanalytical data on the sparingly soluble 1 indicated a stoichiometry "Mo-(OAr)₃(NMe₂)".¹⁰ However, infrared data on 1 is inconsistent

with this simple formulation showing a sharp band at 3250 cm⁻¹ assignable to $\bar{\nu}(NH)$ of a dimethylamine ligand. ¹⁰ A solid state crystal structure of 111 confirmed the presence of an HNMe2 group formed by the cyclometalation of one of the 2,6-diphenylphenoxide ligands (Figure 1). The geometry about the Mo atom can be seen to be square pyramidal with the metalated carbon atom occupying the axial site. The almost planar six-membered metallacycle comfortably maintains an 86° bite at the metal center due to the large Mo-O-C angle of 145°. Substitution of the HNMe₂ ligand by pyridine (py) is easily achieved to yield the monoadduct 2.¹² No evidence to support the formation of a six-coordinate, bis adduct has been obtained. A study of the coordination environment about the molybdenum atom in 1 shows a close approach of an aromatic CH bond of a nonmetalated OAr group in the vacant, axial position almost exactly trans to the Mo-C bond (Figure 1). The distances to this bond are refined

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as Mo-H = 2.67 (5) Å and Mo-C = 2.814 (5) Å, with the CH approaching the metal in a side-on fashion with Mo-H-C = 91 (3)°. However, the distances to this group are certainly on the outer fringes of those reported for well documented agostic interactions. 13-15

Reaction of the tetrahydrides $MH_4(PMePh_2)_4$ (M = Mo, W)¹⁶ with ArOH again does not lead to simple substitution products. Extended thermolysis (phenol melt, 150 °C) results in formation of a deep red mixture from which red crystals of a product of stoichiometry "M(OAr)(PMePh₂)(H)" (M = Mo, 3a; W, 3b) are obtained.¹⁷ A single-crystal X-ray analysis (Figure 2) shows the

HOAr = 2.6-diphenylphenol

phenoxide ligands in 3 to again be chelated to the metal, only this time via an η^6 -interaction through one of the ortho phenyl substituents. 19,20 The ¹H NMR spectra of these 18-electron monohydrides show a sharp set of three upfield resonances (ratio 2:2:1) for the π -ring indicating the ortho and meta protons are equivalent, as well as a triplet for the M-H hydride.¹⁷ There is no indication of rapid exchange of coordinated and noncoordinated orthophenyl rings.²⁰ A structural study on 3a (Figure 2)¹⁸ confirmed the presence of the phenoxide chelate and also managed to identify the hydride position. This position is entirely consistent with the spectroscopic data indicating a virtual plane of symmetry in the molecule equilibrating the phosphine ligands and both sides of the π -bound arene ring. The Mo-O distance of 2.163 (11) Å in 3a is significantly longer than the distances of 1.92–1.96 Å found in 1 and can be rationalized by recognizing the necessary absence of any oxygen-p to metal-d π -bonding in the 18-electron derivatives 3. The Mo-C distances found to the π -bound arene ring in 3a, 2.25-2.34 Å, are comparable to those reported for other molybdenum η^6 -arene compounds. 5c,20 However, these distances are much shorter than those found for the close aromatic CH bond

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^{13.431 (2)} Å, b = 18.213 (5) Å, c = 10.262 (1) Å, $\alpha = 101.68$ (2), $\beta = 106.53$ (1), $\gamma = 77.93$ (2)°, Z = 2, $d_{\rm calcd} = 1.306$ g cm⁻³. A total of 6113 unique data was collected with Mo K α radiation, $4^{\circ} \le 2\theta \le 45^{\circ}$ of which 4625 with I > $3\sigma(I)$ were used in the final refinement to yield residuals of R = 0.038, $R_{\rm W} = 0.054$. The hydrogen atoms H(262) and H(N) were refined. (12) Anal. Calcd for ${\rm MoC_{59}H_{43}NO_3}$ (2): C, 77.88; H, 4.76; N, 1.58. Found: C, 77.74; H, 4.74; N, 1.38.

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in 1 (vide supra) again implying little if any agostic interaction in the amine compound.

Further studies of the possible mechanism for CH bond activation at these metal centers and the possible interconversion of the chelate types are presently underway.

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Supplementary Material Available: Tables of positional parameters, general temperature factors, bond distances, and bond angles for 1 and 3a (28 pages); listing of structure factor amplitudes for 1 and 3a (36 pages). Ordering information is given on any current masthead page.

Two-Dimensional Coherence Transfer NMR Spectroscopy by Isotropic Mixing: Application to **Protein NMR Assignments**

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Homonuclear spin-locking with matched Hartmann-Hahn radio frequency fields to obtain long-range coherence transfer within spin systems has received much attention recently.²⁻⁵ The 2D NMR experiments, often referred to as rotating frame, isotropic mixing, "TOCSY",2 or "HOHAHA"4 experiments, have considerable potential for identifying long amino acid side-chain systems in proteins (e.g., lysine, leucine, arginine, et al.). This is in contrast to the RELAY experiment,^{6,7} where the mixing time must be tuned for optimal relayed intensity for different spin systems, 8,9 and generally only single-step relays can be obtained with good signal intensity for larger proteins due to relaxation limitations. As discussed below, however, there are still compelling reasons for performing RELAY experiments in conjunction with isotropic mixing experiments.

First we demonstrate the results of applying WALTZ-16¹⁰ and MLEV-1611 pulse sequences to obtain long-range coherence

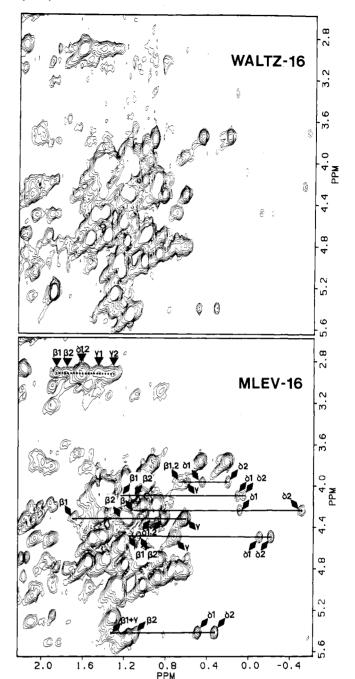


Figure 1. Selected regions of 2D isotropic mixing experiments using WALTZ-16 (above) and MLEV-16 (below) pulse cycling during the spin-locking period and performed off-resonance with 10-kHz spectral widths in both dimensions. 400 t_1 experiments were collected, zero-filled, and transformed, to generate a 1K × 1K matrix. The sample was 8 mM apoNeocarzinostatin (113 amino acids, MW 11500) in 100% D_2O ; spectra were recorded at 45 °C. The six leucine and unique lysine side-chain spin systems are indicated.

transfers (Figure 1). Each spectrum was obtained under identical conditions on our home-built 500-MHz spectrometer¹² using a 20-W pulse for both the preparation and 40-ms mixing pulses, corresponding to an applied radio frequency field of 16-20 kHz. Bax and Davis³ have modified the MLEV-16 pulse sequence by incorporating a 180° refocusing pulse and straddling "purge" pulses to correct for pulse imperfections and to eliminate antiphase and dispersive magnetization. These errors do not appear to dominate isotropic mixing spectra when the applied radio frequency field is sufficiently strong; thus the purge pulses are not required for obtaining good quality spectra. Note also that the

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