where A is the major form but where B and C also contribute significantly. As predicted by this bonding description O, N, C(1), C(2), and C(1N) are coplanar (Table V (supplementary material)). A related compound has been prepared by the reduction of Ph-N=C=O with $(MeC_5H_4)_3U^{23}$ In this complex, $[(MeC_5H_4)_3U]_2(\mu-\eta^1,\eta^2-PhNCO)_2$, the U–N, 2.36 (2) Å, N–C, 1.31 (2) Å, and O–C, 1.36 (2) Å, distances are indistinguishable from those in 2a.

There are not many Cp₃UXY complexes.^{22,24} Some with monodentate, trans X and Y ligands have been reported,^{24–28} but, with the exception of those with three-membered chelate rings,^{10,11,22,23,29–31} no cis-Cp₃UXY complexes have been structurally characterized.³² Since a theoretical model that considers steric effects³⁶ predicts that Cp_3U complexes of both trans-Cp₃UXY and cis-Cp₃UXY complexes, containing a three-membered chelate ring, will be more stable than Cp₃U complexes with larger chelate rings, the characterization of 2a as the first *cis*-Cp₃UXY complex with a four membered chelate ring is particularly significant. In accord with the predictions that molecules of this type should be very sterically crowded,³⁶ the bite distance in the ligand, 2.22 (2) Å, is short, the average C(Cp)-Udistance, 2.84 (2) Å, is long, and the Cp(centroid)-U-Cp-(centroid) angles have been compressed from about 117°, typical for Cp₃U-X complexes, to 112°, 115°, and 110°.

We have earlier postulated³⁷ that



is an intermediate in the formation of Cp₃U=CHP- $(Me)(R)_2$ from Cp_3UCl and $Li(CH_2)(CH_2)PR_2$. The existance of the four-membered chelate ring in 2 supports this hypothesis.

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Supplementary Material Available: Table I, crystal, data collection, and reduction parameters, Table II, bond distances and bond angles, Table III, positional and thermal parameters for $Cp_3U[(NPh)(O)CCHPMe_2Ph] \cdot C_7H_8$, and Table V, distances from the O, N, C(1), C(2), C(1N) least-squares plane (4 pages); Table IV, observed and calculated structure factors for Cp₃U- $[(NPh)(O)CCHPMe_2Ph] \cdot C_7H_8$ (19 pages). Ordering information is given on any current masthead page.

The Slip-Fold Distortion of π -Bound Indenyi Ligands. Dynamic NMR and X-ray Crystallographic Studies of $(\eta$ -Indenyi)RhL₂ Complexes[†]

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Summary: Crystal and molecular structures of two (η - $C_{9}H_{7}$)RhL₂ complexes, $L = \eta - C_{2}H_{4}$ (2a) ($R = 0.031, R_{w}$ = 0.039) and L = PMe_3 (2b) (R = 0.026, $R_w = 0.032$), are reported. Values of the indenyl ring slip parameter, Δ , are 0.161 and 0.201 Å, respectively, indicating significantly greater slip-fold distortion for 2b than for 2a. Comparison with the known structure of $(\eta$ -C₉H₇)Rh(η duroquinone) (2c) for which $\Delta = 0.046$ Å suggests a correlation between ligand σ -donor/ π -acceptor capabilities and the degree of slip-fold distortion. The degree of distortion is also correlated with the ¹³C NMR chemical shift of the ring junction carbon atoms and with the barrier to indenyl ring rotation. The latter has been studied in solution by dynamic FTNMR spectroscopy using (η -1-CH₃C₉H₆)RhL₂ complexes. Larger barriers result from increased distortions; $\Delta G^{\dagger}_{253} = 11.20 \pm 0.03 \text{ kcal mol}^{-1}$ for $(\eta$ -1-CH₃C₉H₆)Rh(PMe₃)₂.

There has been considerable recent interest in the enhanced reactivity of η^5 -indenyl transition-metal complexes toward ligand substitution and related reactions compared with their η^5 -C₅R₅ analogues.¹⁻⁴ This rate enhancement is believed to be related to the relative ease of ring slippage from η^5 to $\eta^{3,1-4}$ Our interest in stoichiometric⁵ and cat-alytic⁶ reactions of η^5 -indenyl rhodium complexes prompted a detailed investigation of factors effecting ground-state slip-fold distortions and related ligand dynamics in a series of $(\eta$ -indenyl)RhL₂ complexes.

We recently reported⁷ the first example of hindered indenyl-ring rotation in an unusual $(\eta$ -C₉H₇)RhLL' complex (1) wherein L = CO, L' = $(\eta^6 - C_6 H_3 M e_3 - 1, 3, 5) Cr(\mu - 1)$ CO)₂(CO), and the observed ΔG^* was ca. 10.8 kcal mol⁻¹.

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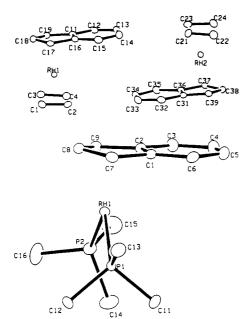


Figure 1. ORTEP views of 2a (top) and 2b (bottom). Important distances (Å) and angles (deg) for 2a, molecule (1): Rh(1)-C(11), 2.359 (5); Rh(1)-C(16), 2.354 (5); Rh(1)-C(17), 2.178 (5); Rh-(1)-C(19), 2.215 (5); Rh(1)-C(18), 2.203 (5); Rh(1)-C(ethylene), 2.135 (5) (average of four values). For 2a, molecule 2: Rh(2)-C(31), 2.361 (5); Rh(2)-C(36), 2.349 (5); Rh(2)-C(37), 2.187 (5); Rh-(2)-C(39), 2.201 (5); Rh(2)-C(38), 2.212 (5); Rh(2)-C(ethylene),2.135 (5) (average of four values). For **2b**: Rh(1)-C(1), 2.449 (4); Rh(1)-C(2), 2.431 (4); Rh(1)-C(7), 2.244 (4); Rh(1)-C(9), 2.234 (4); Rh(1)-C(8), 2.229 (4); Rh(1)-P(1), 2.213 (1); Rh-P(2), 2.210 (1); P(1)-Rh(1)-P(2), 96.69 (4).

Molecular orbital arguments⁸ suggest that an $(\eta$ -C₉H₇)RhL₂ complex will possess a ground-state conformation in which the two ligands straddle a mirror plane incorporating Rh and C₂ of the indenyl ligand and bisecting the remainder of the indenyl ligand. To test the hypothesis that the barrier to hindered indenyl rotation is correlated with the degree of slip-fold distortion,⁷ we have prepared a series of $(\eta$ -1-CH₃C₉H₆)RhL₂ complexes⁹ where L = alkene, CO, PR_3 , or RNC or $L_2 = R_2 PCH_2 CH_2 PR_2$. Incorporation of a methyl group at the 1-indenyl position breaks the molecule's ground-state C_s symmetry allowing the observation of two sets of ligand resonances at low temperature in the ³¹P or ¹³C NMR spectra. We report herein the preliminary results of this study, namely, the crystal and molecular structures of the parent $(\eta$ -C₉H₇)RhL₂ complexes, $L = C_2 H_4$ (2a) and PMe₃ (2b), and the dynamic ³¹P{¹H} NMR spectra of $(\eta$ -1-CH₃C₉H₆)Rh(PMe₃)₂ (**3b**).

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Table I. Slip-Fold Parameters for $(\eta - C_{s}H_{\eta})RhL_{\eta}$ Complexes^a

complex	Δ,Å	HA, deg	FA, deg
2a ^b	0.160	8.34	7.30
2a ^c	0.161	7.81	7.50
2a ^d	0.161	8.08	7.40
$2\mathbf{b}^{e}$	0.201	8.42	7.89
$2c^{f}$	0.046	2.35	2.55
2d ^g	0.23	10.6	8.5

^a $\Delta = d \{ (Rh-C_{3a,7a}) - (Rh-C_{1,3}) \}$; HA (hingle angle) = angle between normals to the least-squares planes defined by C_1 , C_2 , C_3 and C_1 , C_{7a} , C_{3a} , C_3 ; FA (fold angle) = angle between normals to least-squares planes defined by C_1, C_2, C_3 and $C_{3a}, C_4, C_5, C_6, C_7, C_{7a}$. ^b Molecule 1, this work. ^c Molecule 2, this work. ^d Average of values for two independent molecules, 1 and 2. ^e This work. ^f Reference 15. ^g Reference 11.

These results, in conjunction with those of McGlinchey et al.¹⁰ on $(\eta$ -1-CH₃C₉H₆)Rh $(\eta$ -C₂H₄)₂ (3a), support our original hypothesis.

> 1. R = H, $L_2 = (\eta - C_8H_3Me_3 - 1.3, 5)Cr(\mu)$ 2a. R = H, $L = \eta - C_2H_4$ b. R = H, $L_2 = \eta - durcquinone$ d. R = H, $L_2 = (\eta - C_5H_5)_2Zr(\mu - PPh_2)_2$ e. R = H, $L_2 = (\eta - C_5H_5)_2Hf(\mu - PEt_2)_2$ 3a. R = Me, $L = \eta - C_2H_4$ b. R = Me, $L = PMe_3$ c. R = Me, $L_2 = (\eta - C_5H_5)_2Hf(\mu - PEt_2)_2$ $C_{6}H_{3}Me_{3}-1,3,5)Cr(\mu-CO)_{2}(CO)$

Although several structures of indenyl rhodium complexes have been reported,¹¹ most contain complex ligands making an analysis of the relationship between slip-fold distortions and ligand electronic properties difficult. We therefore carried out X-ray diffraction studies¹² on 2a,b, the results of which are illustrated in Figure 1. We chose three parameters¹³ to define the slip-fold distortion: Δ , the slip parameter; HA, the hinge angle; and FA, the fold angle. Table I defines these parameters and lists pertinent values for 2a-d. A clear trend is observed:¹⁴ the degree of slip-fold distortion is indeed a function of the electronic parameters of the ligands; good σ -donors give rise to larger values of Δ , HA, and FA than alkene ligands, wherein each of these values diminishes as the π -acceptor capability of the alkene increases. The degree of slip-fold distortion is

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⁽¹²⁾ Crystal data collection and refinement for 2a: RhC₁₃H₁₅, mono-clinic, P2₁/n, a = 7.782 (2) Å, b = 10.838 (2) Å, c = 25.759 (4) Å, β = 98.47 (2)°, T = -102 °C, V = 2148.9 Å³, Z = 8, μ (Mo) = 15.21 cm⁻¹; Enraf-Nonius CAD4, Mo K α radiation, 4210 data collected by using the ω scan method, $1.6^{\circ} \le 2\theta \le 50.0^{\circ}$, 3219 unique reflections with $I \ge 3.0\sigma(I)$ used in solution and refinement; solution by automated Patterson analysis, refinement by full-matrix least squares, weights $\alpha [\sigma^2(I) + 0.0009I^2]^{-1/2}$ 253 parameters, all non-H atoms anisotropic, H atoms fixed; R = 0.031, $R_{\rm w} = 0.039$, largest residual density = $0.77 \text{ e}/\text{Å}^3$ near Rh. For 2b: Rh- $\begin{array}{l} \mathbf{P}_{2}\mathbf{C}_{15}\mathbf{H}_{25}, \mbox{ monoclinic, } P2_{1}, a=8.454 \ (2) \ \text{\AA}, b=12.201 \ (2) \ \text{\AA}, c=8.901 \ (3) \\ \text{\AA}, \beta=113.54 \ (2)^{\circ}, \ T=-102 \ ^{\circ}\mathbf{C}, \ V=841.7 \ \text{\AA}^{3}, \ Z=2, \ \mu(\mathrm{Mo})=11.70 \ \mathrm{cm}^{-1}; \end{array}$ Enraf-Nonius CAD4, Mo K α radiation, 3015 data collected by using the ω scan method, $3.3^{\circ} \leq 2\theta \leq 62.0^{\circ}$, 2562 unique reflections with $I \geq 3\sigma(I)$ used in solution and refinement as for 2a, 162 parameters, all non-H atoms anisotropic, H atoms fixed, R = 0.026, $R_w = 0.032$, largest residual density = $0.50 \text{ e}/\text{Å}^3$. Data for 2a,b collected at Molecular Structure Corp.

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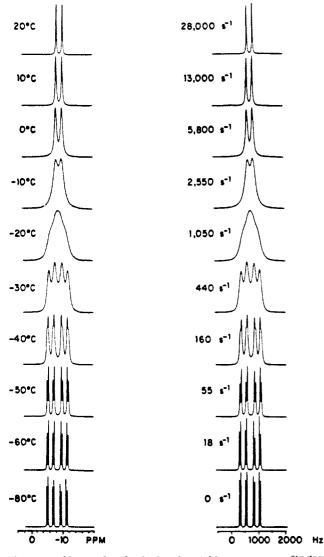


Figure 2. Observed and calculated variable-temperature ³¹P{¹H} NMR spectra (121.69 MHz) for 3b. Simulations were performed using a locally modified version of DNMR3. Activation parameters for indenyl rotation are $\ln A = 29.81 \pm 0.55$, $E_a = 11.45 \pm 0.28$ kcal mol⁻¹, $\Delta G^*_{253} = 11.20 \pm 0.03$ kcal mol⁻¹, $\Delta H^* = 10.95 \pm 0.28$ kcal mol⁻¹, and $\Delta S^* = -0.97 \pm 1.07$ eu.

also strongly correlated with the ¹³C chemical shift of the ring-junction carbons $C_{3a,7a}$: larger distortions resulting in larger downfield shifts.^{14a,16}

We also find that the degree of distortion is paralleled by the barrier to hindered rotation of the indenyl ring.⁷ In order to ascertain the effect of 1-CH₃ substitution on the observed barriers, unsubstituted and substituted complexes 2e and 3c were examined;^{11,17} observed barriers were identical within experimental error.¹⁷ We therefore conclude that the activation parameters obtained for $3b^{18}$ (Figure 2) are good estimates of those which are unavailable for 2b. The values for 3b are significantly larger than those for 3a ($\Delta G^* = 8.5 \text{ kcal mol}^{-1}$)¹⁰ in keeping with our proposal⁷ that an increase in the degree of slip-fold distortion should result in a larger barrier to rotation.

We will report further analyses of (indenyl) RhL_2 (L = PR_3 , CO, RNC; $L_2 = R_2PCH_2CH_2PR_2$) complexes in due course.19

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Registry No. 2a, 63428-46-6; 2b, 75094-84-7; 2e, 100113-59-5; 3b, 108010-58-8; 3c, 108010-59-9.

Supplementary Material Available: A drawing of 2a and tables of crystal data, fractional coordinates and isotropic thermal parameters, anisotropic thermal parameters, interatomic distances, and intramolecular angles for 2a and 2b (8 pages); listings of structure factors for 2a and 2b (16 pages). Ordering information is given on any current masthead page.

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(19) Full details will appear in a manuscript describing our joint efforts with the authors in ref 10.

Novel Pyrazolylborato- and Pyrazolyl-Ruthenium **Complexes.** The Crystal and Molecular Structures of $[{H(\mu-H)B(3,5-Me_2pz)_2}Ru(\eta^4-C_8H_{12})H]$ and $[(\eta^4 - C_8 H_{12})_2 Ru_2 H[\mu - N = C(Me)pz](\mu - pz)(\mu - H)]$ Confirmation of the First Examples of the Ru-H-B Agostic Interaction and the Bridging Amidine Ligand

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Summary: The reactions of $[(\eta^4-C_8H_{12})RuH(NH_2NMe_2)_3]$ - PF_6 and $[{(\eta^4-C_8H_{12})RuCl(Me)(NCMe)}_2]$ with $K[H_2B(3,5 Me_2pz_2$] (pz = 1-pyrazolyl) give [{H(μ -H)B(3,5-Me_2pz)_2}- $Ru(\eta^4-C_8H_{12})X$] (X = H, Me), the first examples of the Ru····H–B interaction and verified in $[{H(\mu-H)B(3,5-)}]$

^{(14) (}a) A similar trend has been observed for analogous iridium complexes: Merola, J. S., private communication. (b) The observed slip-fold distortions do not appear to arise from steric interactions of the ligands with the indenyl ring as there are no short intramolecular nonbonded interactions. In addition, they do not appear to be related to crystal packing constraints. Thus, real-time computer graphic analyses of the unit cell contents do not indicate any unusual intermolecular interactions, and the phenomenon has been shown to persist in solution (vide infra). (c) Preliminary results indicate significant slip-fold distortions are present in $(\eta$ -indenyl)RhL₂ complexes containing certain cylindrical π -acceptor ligands, e.g. L = CO or RNC. A complete analysis of these and related complexes will appear in due course.¹⁹

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