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The X-ray Structures of $H_5Re(PMe_2Ph)_3$ and $H_7Re(PMe_2Ph)_2$

RAYMOND G. TELLER [1a], W. EAMON CARROLL [1b] and ROBERT BAU*

Department of Chemistry, University of Southern California, Los Angeles, Calif. 90089, U.S.A. Received December 16, 1983

The structures of $H_5Re(PMe_2Ph)_3$ and H_7Re - $(PMe_2Ph)_2$ have been solved by single-crystal X-ray diffraction methods. Although the hydride ligands could not be directly located in this study, the overall structure of the complexes could be deduced from a knowledge of the central rhenium/phosphorus core of the molecules. The ReP₃ skeleton of $H_5Re(PMe_2)$ - $Ph)_3$ has distorted pyramidal geometry (P-Re-P angles 149.5°, 101.9°, 99.8°) consistent with a dodecahedral structure for the H₅ReP₃ core. The ReP₂ backbone of $H_7Re(PMe_2Ph)_2$ is bent (P-Re-P angle146.8°), suggesting a tricapped trigonal prismatic geometry for the H_7ReP_2 core in which the P atoms are placed in opposing axial and equatorial positions. Crystallographic details: H₅Re(PMe₂Ph)₃: space group $P2_1/c$ (monoclinic); a = 6.876(3) Å, b =19.493(7) Å, c = 19.646(8) Å, $\beta = 103.26(2)^\circ$, V = 2563.0 Å³, Z = 4; R = 6.7% for 2067 reflections. $H_7Re(PMe_2Ph)_2$: space group $P2_1/n$ (monoclinic); a = 19.083(17) Å, b = 6.337(4) Å, c = 15.234(13)Å, $\beta = 93.72(4)^{\circ}$, V = 1834.0 Å³, Z = 4; R = 5.0%for 1672 reflections.

Introduction

During the past few years, we have been involved in the structural characterization of polyhydrido complexes of the form H_xML_y , where L is a tertiary phosphine. Most of these compounds were first prepared in the 1960's by Chatt, Shaw and coworkers [2-5] and are interesting structurally because of (a) their high co-ordination number and (b) the unusual feature of a large number of hydride ligands attached to the central metal atom. In earlier publications we have reported the results of neutron diffraction studies of H₄Os(PMe₂Ph)₃ [6] and H₈Re₂-(PEt₂Ph)₄ [7] and the X-ray diffraction structure of $H_3Ir(PMe_2Ph)_3$ [8]. In this paper we describe the results of X-ray work on the eight-coordinate complex $H_5 \operatorname{Re}(\operatorname{PMe}_2\operatorname{Ph})_3$ (I) and the nine-coordinate complex H₇Re(PMe₂Ph)₂ (II).

Experimental

All preparations were based on the original synthetic procedure of Chatt and co-workers [3] and were carried out under a nitrogen atmosphere with freshly distilled, dried and degassed solvents. Ethanol and chlorinated solvents used in the preparation of the chloride complexes were degassed by bubbling nitrogen through the solvents for half an hour prior to use. Drying agents for other solvents used in the preparation are Na/benzophenone for tetrahydrofuran (THF) and calcium hydride for hexane, toluene and benzene.

$ReCl_3(PMe_2Ph)_3$

First, 2.0 g of KReO₄ and 6 ml of 12 N HCl were added to 100 ml of ethanol in a Schlenk tube fitted with a condenser. Then 7.5 ml of PMe₂Ph was added and the mixture was heated at reflux for 18 hours. Upon cooling, an orange crystalline material precipitated. This solid was dissolved in 35 ml of CH_2Cl_2 and a pale yellow impurity removed by filtration. Recrystallization from CH_2Cl_2 /ethanol yielded 4.1 g (84% yield) of the product as orange crystals.

$ReCl_4(PMe_2Ph)_2$

The tetrachloro species was prepared by addition of 0.05 g of ReCl₃(PMe₂Ph)₃ to 25 ml of CCl₄ and heating at reflux for 24 hours. Reducing the volume of the purple solution gave a purple solid, which was recrystallized from CH₂Cl₂/hexane yielding 0.30 g (70% yield) of ReCl₄(PMe₂Ph)₂.

$H_5Re(PMe_2Ph)_3(I)$

The rhenium pentahydride was prepared by dissolving 0.35 g of ReCl₃(PMe₂Ph)₃ in 15 ml of THF and adding 0.15 g of LiAlH₄ in portions. The mixture was heated at reflux for 90 minutes and hydrolyzed with wet THF (10% water) after cooling to -20 °C. After removal of the THF *in vacuo*, the solid was extracted with hexane and isolated by removal of the solvent. Recrystallization from hexane or ethanol yielded clear crystals of H₅Re(PMe₂Ph)₃ (0.18 g, 61% yield). The product was identified via its ¹H NMR spectrum in d⁶-benzene: δ 7.0–7.9 ppm

^{*}Author to whom correspondence should be addressed.

[m, 15 H], 1.89 [d, 18 H, J(P-H) = 7 Hz], -5.56 [q, 5 H, J(P-H) = 18 Hz].

$H_7 Re(PMe_2Ph)_2$ (II)

The rhenium heptahydride was prepared by dissolving 0.50 g of ReCl₄(PMe₂Ph)₂ in 15 ml of THF and adding 0.14 g of LiAlH₄ in portions. The mixture was heated at reflux for 30 minutes and hydrolyzed with wet THF. The solvent was removed *in vacuo* and the residue was extracted with methanol. Recrystallization from hexane yielded 0.11 g of H₇Re(PMe₂Ph)₂ (29% yield). The product was identified via its ¹H NMR spectrum in d⁶-benzene: δ 7.0–7.9 ppm [m, 10 H], 2.05 [d, 12 H, J(P–H) = 7 Hz], -4.75 [t, 7 H, J(P–H) = 19 Hz].

X-ray Analyses of $H_5Re(PMe_2Ph)_3$ (I) and $H_7Re-(PMe_2Ph)_2$ (II)

Precession photographs of I and II showed systematic extinctions [(0k0, k = 2n + 1) and (h0l, l = 2n + 1) for I; (0k0, k = 2n + 1) and (h0l, h + l = 2n + 1) for II] indicating space groups $P2_1/c$ and $P2_1/n$, respectively. Because I slowly oxidizes in air, the crystal used for data collection was mounted in a glass capillary. Crystal and refinement details for each complex are reported in Table I.

For I, a diamond-shaped crystal of dimensions 0.31 mm \times 0.34 mm \times 0.56 mm was mounted approximately along its a* axis. Accurate centering of 45 reflections on a Nonius CAD-3 diffractometer allowed the calculation of an orientation matrix and unit cell parameters.

For II, an approximately tetragonally-shaped crystal (0.35 mm \times 0.35 mm \times 0.65 mm) was mounted on a glass fiber approximately along its b* axis. Least-squares refinement of 36 carefully-centered reflections yielded cell constants and an orientation matrix used in subsequent data collection.

Data were collected on a Nonius CAD-3 diffractometer with Zr-filtered MoK α radiation to a (sin θ/λ) limit of 0.54 Å⁻¹ in the $\theta/2\theta$ scan mode (10°/min scan rate). Each reflection was scanned a maximum of six times, depending on intensity. Background radiation for each reflection was counted at the beginning and at the end of each scan, for a duration $(t_{\rm h})$ of one-half of the scan time $(t_{\rm s})$. The intensities of three standard reflections [(410), (190), and (008) for I; (040), (910), and (01-9) for II] were collected at intervals of 50 reflections. No systematic fluctuations were observed in these monitor reflections for either compound. Two quadrants of data were collected for each compound: $(+h, \pm k,$ ± 1 , 5442 reflections for I; $\pm h$, $\pm k$, ± 1 , 4200 reflections for II).

The intensity (I) and the standard deviation of the intensity $[\sigma(I)]$ for each reflection were calculated as:

$$I = [C - (t_s/t_b)B]$$

$$\sigma(I) = [C + (t_s/t_b)^2 \ b + (pI)^2]^{1/2}$$

where C is the integrated intensity of the reflection, B the background count, and p = 0.04 for I and 0.03 for II. Values for F_0^2 were obtained after correcting for Lorentz and polarization effects. An empirical absorption correction was also applied to each data set, with transition coefficients (normalized to unity) ranging from 1.100 to 0.846 for I and 1.540 to 0.521 for II. Equivalent reflections were merged and those with F_0^2 values less than $3\sigma(F_0^2)$ were rejected, leaving 2067 and 1672 reflections for the subsequent structure analyses of I and II, respectively.

Patterson maps [9] revealed the locations of the Re and P atoms for each molecule. Successive cycles of structure factor calculations and difference-Fourier syntheses allowed the location of all non-hydrogen

TABLE I. Crystal Data and Details of Refinement for H₇Re(PMe₂Ph)₂ and H₅Re(PMe₂Ph)₃.

		$H_7 Re(PMe_2Ph)_2$	$H_5Re(PMe_2Ph)_3$		
Crystal type		monoclinic	monoclinic		
Space group		$P2_1/n$ (#14n)	$P2_{1}/c$ (#14)		
Cell constants	a	19.083(17) Å	6.876(3) A		
	b	6.337(4) Å	19.493(7) Å 19.646(8) Å 103.26(2)°		
	с	15.234(13) Å			
	β	93.72(4)°			
	V	1834.0 A ³	2563.0 Å ³		
Number of molecules in the unit cell		4	4		
Calculated density		1.70 g-cm^{-3}	1.57 g-cm^{-3}		
Observed density		1.70 g-cm^{-3}	1.58 g-cm^{-3}		
$\sin \theta / \lambda$ data collection limit		0.54 A^{-1}	0.54 A^{-1}		
Reflections used in analysis $(F > 3\alpha F)$		1672	2067		
Final agreement factors		R(F) = 0.050	R(F) = 0.067		
5		R(wF) = 0.055	R(wF) = 0.075		

atoms. Three cycles of full-matrix least-squares refinement resulted in agreement factors of R = 0.072for I and R = 0.053 for II [10]. At this point, calculated phenyl hydrogen positions were added to the coordinate list. Several more cylces of full-matrix least-squares refinement (in which the temperature factors of the Re and P atoms were allowed to vary anisotropically), with fixed hydrogen contributions, resulted in final agreement factors of R = 0.067 and R = 0.052 for I and II, respectively. In the final cycle of refinement, the largest parameter shifts were 1/4 of their estimated standard deviations. Attempts to locate the hydridic hydrogens with difference Fourier maps of decreasing $(\sin \theta / \lambda)$ maxima [11] revealed several large peaks near the Re atoms. However, these peaks gave unreasonable bond lengths and angles and are probably due to ripple peaks from the metal atoms. Final atomic positions and thermal parameters are reported in Table II for H₅Re(PMe₂Ph)₃ and Table III for $H_7(PMe_2Ph)_2$.

Discussion

Selected bond distances and angles for H_5 Re-(PMe₂Ph)₃ and H_7 Re(PMe₂Ph)₂ are reported in

TABLE II. Final Atomic Parameters for H₅Re(PMe₂Ph)₃.^a

Tables IV and V. Stereoscopic unit cell plots are illustrated in Figs. 1 and 2.

Although the hydride ligands were not located in either X-ray study, an analysis of the rhenium-phosphorus geometries can provide clues about the overall coordination geometries about the metal atoms in the complexes [12]. For $H_5 \text{Re}(\text{PMe}_2\text{Ph})_3$ (Fig. 3), the relevant parameters to consider are the P-Re-P bond angles $(149.5(2))^\circ$, $101.9(2)^\circ$, and $99.8(2)^\circ$] and the distance of the Re atom from the plane formed by the three P atoms (0.36 Å). These features are consistent with a dodecahedral geometry for $H_5 \text{Re}(\text{PMe}_2\text{Ph})_3$, as shown in *III*.





Atom	x	у	2	β_{11} or B	β22	β33	β_{12}	β ₁₃	β23
Re	2001(1)	2547(1)	4991(1)	128(2)	18(2)	15(3)	21(2)	1(1)	2(1)
P(1)	4585(8)	2622(3)	5997(3)	152(12)	29(2)	19(2)	22(7)	19(8)	3(3)
P(2)	177(8)	3041(3)	3937(3)	202(16)	26(2)	21(2)	29(9)	-2(9)	3(3)
P(3)	2518(9)	1416(3)	4635(3)	165(15)	21(2)	35(2)	16(8)	37(9)	0(3)
C(1)	5633(35)	1822(12)	6430(12)	4.37(0.47)					
C(2)	3997(39)	3087(13)	6721(14)	5.63(0.58)					
C(3)	6791(41)	3064(14)	5970(14)	5.83(0.58)					
C(4)	7431(40)	1552(13)	6338(13)	5.59(0.58)					
C(5)	8125(43)	934(15)	6687(16)	6.30(0.65)					
C(6)	7121(50)	668(16)	7052(18)	7.21(0.73)					
C(7)	5452(44)	884(15)	7228(15)	6.42(0.67)					
C(8)	4482(40)	1468(14)	6873(15)	5.72(0.59)					
C(9)	1019(31)	3872(11)	4063(11)	3.72(0.43)					
C(10)	1775(39)	3283(13)	3299(14)	5.62(0.58)					
C(11)	-1980(44)	2572(14)	3389(14)	6.81(0.63)					
C(12)	-43(40)	4482(14)	3995(14)	5.53(0.57)					
C(13)	-962(51)	5134(17)	4108(17)	7.88(0.78)					
C(14)	-2686(53)	5123(17)	4294(17)	8.16(0.82)					
C(15)	-3677(46)	4504(17)	4362(17)	7.56(0.75)					
C(16)	-2771(34)	3900(11)	4247(11)	4.21(0.46)					
C(17)	934(33)	1120(11)	3809(12)	4.10(0.45)					
C(18)	5014(42)	1162(15)	4529(15)	6.48(0.65)					
C(19)	2079(46)	733(16)	5242(17)	7.28(0.73)					
C(20)	1459(38)	1222(14)	3174(14)	5.65(0.58)					
C(21)	229(40)	1007(13)	2551(15)	5.59(0.56)					
C(22)	-1541(41)	729(14)	2568(15)	5.88(0.60)					
C(23)	-2086(38)	642(12)	3196(15)	5.21(0.54)					
C(24)	-895(33)	794(11)	3780(13)	4.01(0.45)					

^aPositional and anisotropic thermal parameters have been multiplied by 10⁴. The form of the anisotropic thermal parameter is $exp(-(\beta_{11}h^2 + \beta_{22}k^2 + \beta_{33}l^2 + \beta_{12}hk + \beta_{13}hl + \beta_{23}kl))$.

Atom	x	у	Ζ	β_{11} or B	β22	β ₃₃	β_{12}	β_{13}	β23
Re	4345(1)	5084(1)	7338(1)	171(2)	2259(18)	353(4)	59(4)	65(3)	- 90(14)
P(1)	3790(2)	6833(5)	8500(3)	223(13)	2017(92)	391(22)	91(58)	79(25)	72(71)
P(2)	5357(2)	3232(5)	6906(3)	212(13)	1817(89)	448(24)	216(55)	69(25)	- 59(68)
C(1)	3044(9)	8474(24)	8172(11)	5.36(0.36)					
C(2)	4311(9)	8619(22)	9170(10)	4.75(0.33)					
C(3)	3440(7)	5043(20)	9291(9)	3.38(0.24)					
C(4)	3851(10)	4319(24)	9987(12)	5.35(0.37)					
C(5)	3617(10)	2827(25)	10578(12)	5.65(0.38)					
C(6)	2944(10)	2119(25)	10427(12)	5.70(0.39)					
C(7)	2505(11)	2819(27)	9750(13)	6.42(0.43)					
C(8)	2777(10)	4276(25)	9154(12)	5.77(0.40)					
C(9)	5810(8)	1558(22)	7733(10)	4.84(0.34)					
C(10)	5195(9)	1478(24)	5949(11)	5.07(0.34)					
C(11)	6046(7)	4917(19)	6572(10)	3.71(0.25)					
C(12)	6558(7)	5725(18)	7200(9)	3.21(0.25)					
C(13)	7044(9)	7131(22)	6973(11)	4.97(0.34)					
C(14)	7078(9)	7758(22)	6138(11)	4.84(0.34)					
C(15)	6619(9)	6973(23)	5499(11)	4.95(0.34)					
C(16)	6091(9)	5588(22)	5726(11)	4.98(0.37)					

TABLE III. Final Atomic Parameters for H₇Re(PMe₂Ph)₂.^a

^aPositional and anisotropic thermal parameters have been multiplied by 10⁴. The form of the anisotropic thermal parameter is $\exp(-(\beta_{11}h^2 + \beta_{22}k^2 + \beta_{33}l^2 + \beta_{12}hk + \beta_{13}hl + \beta_{23}kl))$.

TABLE IV. Selected	Distances (in A	4) and	Angles	(in degrees)
in H ₅ Re(PMe ₂ Ph) ₃ .					

TABLE V. Selected Distances (in Å) and Angles (in degrees) in $H_7Re(PMe_2Ph)_2$.^a

Bond distances				Bond distances			
Re-P(1)	2.340(5)	P(1) - C(1)	1.840(23)	Re-P(1)	2.396(4)	P(1) - C(1)	1.808(17)
Re-P(2)	2.365(6)	P(1) - C(2)	1.756(24)	Re-P(2)	<u>2.395(4)</u>	P(1) - C(2)	1.781(18)
Re-P(3)	2.365(6)	P(1) - C(3)	1.810(22)	Average Re-P	2.396(1)	P(1)C(3)	1.816(14)
Average ReP	2 357(8)	P(2)-C(9)	1.859(23)			P(2)-C(9)	1.818(15)
Avoiago no 1	2.557(0)	P(2) - C(10)	1.906(24)			P(2) - C(10)	1.841(17)
		P(2)C(11)	1.861(23)			P(2)-C(11)	<u>1.796(14)</u>
		P(3)-C(17)	1.828(22)			Average P-C	1 810(8)
		P(3) - C(18)	1.843(28)			interage i c	1.010(0)
		P(3)-C(19)	1.857(27)				
		Average P-C	1.840(14)	Bond angles			
Bond angles				P(1)-Re-P(2)	146.8(1)	C(1) - P(1) - C(2)	101.0(7)
P(1)-Re-P(2)	149.5(2)					C(1) - P(1) - C(3)	102.8(7)
P(1)-Re-P(3)	99.8(2)			Re - P(1) - C(1)	116.3(6)	C(2) - P(1) - C(3)	103.6(7)
P(2)-Re-P(3)	101.9(2)			Re-P(1)-C(2)	117.3(6)	C(9) - P(2) - C(10)	103.9(7)
Re - P(1) - C(1)	118.6(8)	C(1) = P(1) = C(2)	101.3(11)	Re-P(1)-C(3)	113.8(4)	C(9) - P(2) - C(11)	102.7(7)
Re - P(1) - C(2)	115.6(9)	C(1) - P(1) - C(3)	100.4(12)	Re-P(2)-C(9)	116.8(5)	C(10)-P(2)-C(11)	102.8(7)
Re-P(1)-C(3)	119.9(9)	C(2) - P(1) - C(3)	97.3(12)	Re-P(2)-C(10)	114.7(4)	Average C-P-C	102.8(4)
Re-P(2)-C(9)	113.6(7)	C(9) - P(2) - C(10)	102.2(10)	Re - P(2) - C(11)	114.1(4)	C	. ,
Re - P(2) - C(10)	113.9(8)	C(9) - P(2) - C(11)	100.7(11)	Average De D.C.	115 5(6)		
Re-P(2)-C(11)	120.0(8)	C(10) - P(2) - C(11)	104.1(11)	Average Re-r-C	115.5(0)		
Re-P(3)-C(17)	117.2(7)	C(17) - P(3) - C(18)	101.3(12)	P(1) = C(3) = C(4)	120 6(12)		
Re-P(3)-C(18)	119.4(9)	C(17) - P(3) - C(19)	101.1(12)	P(1) - C(3) - C(8)	120.3(11)		
Re-P(3)-C(19)	114.9(9)	C(18) - P(3) - C(19)	100.0(12)	P(2) - C(11) - C(12)	120.6(11)		
Average Re-P-C	117.0(9)	Average C-P-C	100.9(6)	P(3)-C(11)-C(16)	123.3(12)		
				Average P-C-C	121.2(7)		

complete structure determination, but an analysis of the Patterson map yielded the characteristic onelarge/two-small pattern of P-Re-P angles corresponding to structure III (134.3, 107.2, 107.1°).

^aStandard deviations of mean values (\bar{x}) are calculated as $[\Sigma(x_i - \bar{x})^2/n(n-1)]^{1/2}$. The resulting deviations are to be regarded as rough estimates of uncertainty in cases where $n \le 3$.

X-ray Structures of Rhenium Polyhydrido Complexes



Fig. 1. A stereoscopic unit cell plot of H₅Re(PMe₂Ph)₃.



Fig. 2. A stereoscopic unit cell plot of H₇Re(PMe₂Ph)₂.



Fig. 3. A molecular plot of $H_5Re(PMe_2Ph)_3$. Phenyl hydrogens are omitted for clarity.

A very recent neturon-diffraction study [13b] of compound closely related to ours, $H_5 \text{Re}(\text{PMePh}_2)_3$, has confirmed structure *III*, again with the same pattern of P-Re-P angles: [146.5(2), 103.4(2), 99.0(2)°] and [148.0(2), 101.6(2), 100.9(2)°] for the two independent molecules in the unit cell. These three studies suggest that the nature of the phosphine ligand has little effect on the coordination geometry of this particular class of eight-coordinate complexes. The dodecahedral geometry *III*, incidentally, has also been confirmed by the X-ray analysis of another eight-coordinate polyhydride complex, $H_4 Mo(PMe_2Ph)_4$ [14], a study in which the hydride ligands (as in the neutron study of $H_5 \text{Re}$ -(PMePh₂)₃ [13b]) were actually located.

The ReP₂ geometry of $H_7Re(PMe_2Ph)_2$ is illustrated in Fig. 4. The central P-Re-P backbone is found to be distinctly bent, with an angle of



Fig. 4. Molecular plot of $H_7 \text{Re}(PMe_2Ph)_2$. Phenyl hydrogens have been omitted for clarity.

146.8(1)°. Structural data on nine-coordinate molecules indicate that the tricapped trigonal prism is by far the most preferred geometry [15] and the structure of $H_7Re(PMe_2Ph)_2$ can be rationalized in terms of this model. One can envisage several ways in which a bent P-Re-P backbone can be accommodated in a tricapped trigonal prism: a structure with equatorial-equatorial phosphines (*IV*), axialequatorial (*V*), or axial-axial (*VI*). The 'normal'



values expected for the P-Re-P angles in the three cases (i.e., in the absence of steric effects) are 120° for IV, 139.1° for V and 100.0° for VI. It is clear that, for structures IV and VI, a fair amount of distortion would have to take place in order for the P-Re-P angle to expand to the observed value of 146.8°. This distortion is presumably largely caused by steric effects arising from the relatively bulky phosphine ligands. The axial-equatorial model V, on the other hand, is the least crowded arrangement, and would be the one favored solely on steric grounds. At first glance, the ¹H NMR data [3], which indicate the presence of two equivalent phosphorus atoms, might argue against structure V, but it should be kept in mind that virtually all polyhydride/phosphine complexes are fluxional on the NMR time-scale [16].

Very recent neutron-diffraction work by Howard, Spencer and co-workers on two different crystalline modifications of $H_7Re(PPr_1^iPh)_2$ [17] also revealed a bent backbone (P-Re-P = 141.0°, 147.0°), but unambiguous location of the hydride ligands was prevented by complicated disorder effects. Nevertheless, sufficient detail was seen, through the partial resolution of the hydride peaks, to allow the authors to suggest a monocapped square-prismatic structure (VII) for $H_7Re(PPr_1^iPh)_2$, as an alternative to the tricapped trigonal prismatic structure IV-VI.



The same authors also examined the related ninecoordinate complex $H_6W(PPr^i_2Ph)_3$, again with neutron diffraction [18]. In that compound they found



a WP₃ skeleton which was planar, but curiously distorted from three-fold symmetry. The unexpected two-large/one-small pattern of P-W-P angles in H₆W(PPrⁱ₂Ph)₃ [126.49(5)°, 125.53(6)°, 107.89(6)°] led Howard, Spencer *et al.* to suggest structure VIII (ax-ax-eq) over the more symmetrical all-equatorial structure IX. (The W-P distances in H₆W(PPrⁱ₂-Ph)₃, incidentally, also show a two-long/one-short pattern consistent with structure VIII: W-P = 2.524-(1), 2.519(2) Å, 2.424(2)) [18].

In summary, our X-ray structural results on H_7 -Re(PMe₂Ph)₂ are consistent with any of the models IV to VI (or, indeed, even VII) but on steric arguments we prefer structure V. The final assignment must await more definitive hydride positions from a neutron diffraction analysis.

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