

Structure of Bromotris(triphenylphosphine)rhodium(I) by Extended X-Ray Absorption Fine Structure (EXAFS)

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Summary The previously unknown interatomic bond distances and co-ordination geometry of bromotris(triphenylphosphine)rhodium(I), $[\text{RhBr}(\text{PPh}_3)_3]$, have been determined by a new structural technique, extended X-ray absorption fine structure.

THE new analysis technique, extended X-ray absorption fine structure (EXAFS), has been used to determine interatomic distances in iron-sulphur proteins,¹ copper salts in aqueous solutions,² and a polymer-bound rhodium(I) catalyst.³ We report here determination of the interatomic bond distances and co-ordination of bromotris(triphenylphosphine)rhodium(I) by EXAFS. Structural data for this complex have not been reported, although the structure of the chloro-analogue⁴ (Wilkinson's catalyst) has been determined.^{5,6} There appears to be only one report of a structure in which an $\text{Rh}^{\text{I}}\text{-Br}$ distance has been determined.⁷ The catalytic activities of $[\text{RhX}(\text{PPh}_3)_3]$, where $\text{X} = \text{Cl}$ or Br , are expected to be similar, suggesting the need for structural data on rhodium(I) systems containing bromine and phosphine ligands.⁸

The EXAFS experiments were performed with synchrotron radiation at the Stanford Synchrotron Radiation Project.⁹ The current theoretical view of EXAFS¹⁰ is that the modulation of the X-ray absorption coefficient, $\chi = (\mu - \mu_0)/\mu_0$, is due to interference between the back-scattered and outgoing photoelectrons in the photo-absorption matrix element, where μ_0 is the absorption coefficient of an isolated atom. Thus, the modulation

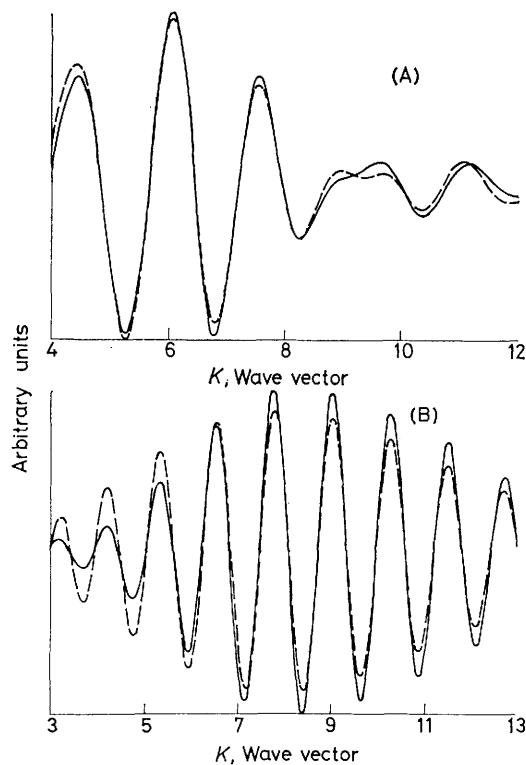


FIGURE 1. Fitting results: (A) the Rh-EXAFS in $[\text{RhBr}(\text{PPh}_3)_3]$; (B) the Br-EXAFS in $[\text{RhBr}(\text{PPh}_3)_3]$; solid lines, experimental data; broken lines, fit.

$\Delta\mu/\mu$, of the X-ray absorption coefficient of an atom is given by equation (1), where N_j is the number of scattering

$$\Delta\mu/\mu = \sum_j N_j |f_j(k, \pi) \exp(-2\sigma_j^2 k^2) \sin[2kR_j + \phi_j(k)] / R_j^2 k \quad (1)$$

atoms j at a distance R_j from the absorbing atom with a Debye-Waller like factor of $\exp(-2\sigma_j^2 k^2)$.

The $\phi_j(k)$ and $f_j(k, \pi)$ terms are energy-dependent phase shifts and amplitude functions for the scatterers respectively and k is the wave vector of the emitted photoelectron.

In the data analysis,¹¹ theoretical phase shifts and amplitudes were used to fit the data to equation (1) and the measured EXAFS for Rh and Br in $[\text{RhBr}(\text{PPh}_3)_3]$ (Figure 1). The Fourier transforms of the data in Figure 2 show that for the Rh-EXAFS in $[\text{RhBr}(\text{PPh}_3)_3]$, there are two peaks, Rh-P and Rh-Br, indicative of multiple distances. In the Br-EXAFS the Br sees only the rhodium atom as a nearest neighbour, thus giving rise to a single frequency, and hence a single peak and distance in the analysis.

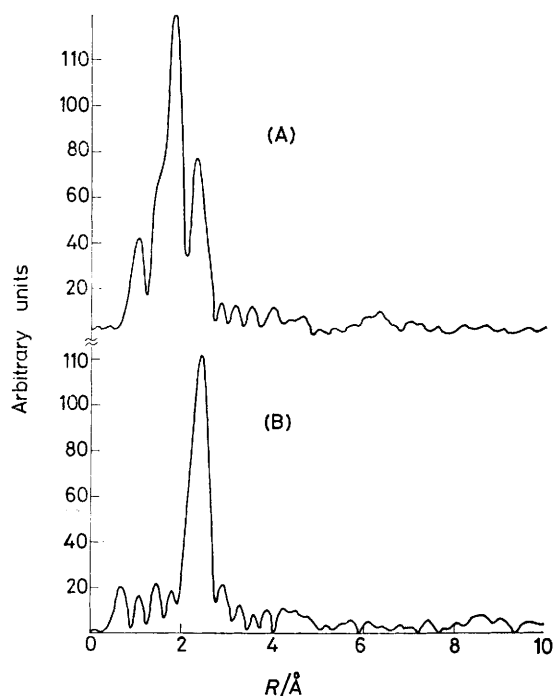


FIGURE 2. (A) Fourier transform of the Rh-EXAFS in $[\text{RhBr}(\text{PPh}_3)_3]$; (B) Fourier transform of the Br-EXAFS in $[\text{RhBr}(\text{PPh}_3)_3]$.

We have, therefore, two independent results with which to confirm the Rh-Br distance. The results of fitting the Rh-EXAFS and Br-EXAFS data for $[\text{RhBr}(\text{PPh}_3)_3]$ are shown in Table 1. The results gave three distinct deter-

TABLE 1. Interatomic distance in $[\text{RhBr}(\text{PPh}_3)_3]$ by analysis of the Rh-EXAFS and Br-EXAFS

	Rh-EXAFS		Br-EXAFS	
	Method 1 ^a	Method 2	Method 1	Method 2
Rh-Br	[1] ^b 2.540 (8)	—	2.536 (8)	2.535 (8)
Ph-P(1)	[1] 2.18 (1)	—	—	—
Rh-P(2)	[2] 2.31 (1)	—	—	—

^a Method 1, fitted function, equation (1); Method 2, empirical method. ^b Number of bonds of this type.

minations of the Rh-Br distances, two of which were obtained from independent experiments and a third from an empirical data analysis program. The Rh-Br distances were 2.540, 2.535, and 2.536 Å.

The Rh-P distances were 2.18 and 2.31 Å, which were determined by fitting the Rh-EXAFS data with theoretical Rh-P and Rh-Br phase shifts and amplitudes.

The fitting technique also gave information about the co-ordination number of atoms attached to the rhodium (absorber) atom. In Figure 3 the sum of the squares of the

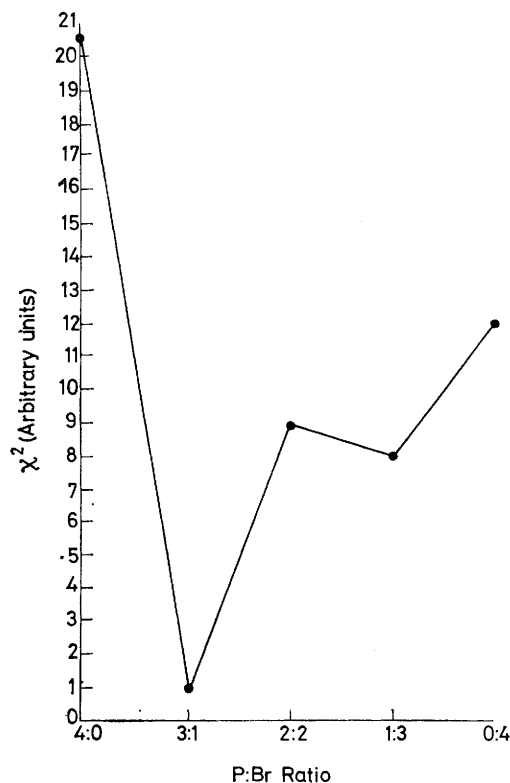


FIGURE 3. Plot of χ^2 (sum of squares of residuals) vs. P:Br ratio. χ^2 minimum occurs at a P:Br ratio of 3:1.

fit residuals (χ^2) is plotted for several values of $N_P = N_{P(1)} + N_{P(2)}$ and N_{Br} where N_j is the number of scattering atoms j at a distance R_j to the absorbing atom. In this fit, which minimized at $N_P = 3$ and $N_{Br} = 1$, only integral $N_{P(1)}$, $N_{P(2)}$, and N_{Br} values were used.

TABLE 2. Comparison of distances in $[\text{RhBr}(\text{PPh}_3)_3]$ and $\text{RhCl}(\text{PPh}_3)_3$

	$[\text{RhCl}(\text{PPh}_3)_3]$		$[\text{RhBr}(\text{PPh}_3)_3]$	$[\text{RhBrP}(\text{C}_6\text{H}_4\text{-CH}(\text{CH}_2\text{-o})_3)]$
	X-Ray ^a	EXAFS	EXAFS	X-Ray ^b
Rh-X	2.37	[1] ^c 2.35(1)	[1] ^c 2.54(1)	[1] ^c 2.587(3)
Rh-P(1)	2.214	[1] 2.23(1)	[1] 2.18(1)	[1] 2.176(10)
Rh-P(2)	[2] 2.33	[2] 2.35(1)	[2] 2.31(1)	—

^a From refs. 5 and 6. ^b From ref. 7. ^c Number of bonds of this type.

In the similar chlorotris(triphenylphosphine)rhodium(I) complex,¹² the distances which were determined by the Rh-EXAFS yielded results which were in agreement with

those determined by X-ray crystallography^{5,6} (see Table 2). In $[\text{RhBr}(\text{PPh}_3)_3]$ as in $[\text{RhCl}(\text{PPh}_3)_3]$, two of the Rh-P distances are longer than the third. This suggests that there must be an electronic effect, causing the *trans* influence of P to be greater than that of Br. The geometry of the co-ordination polyhedron about the rhodium atom in $[\text{RhBr}(\text{PPh}_3)_3]$ was not determined by the EXAFS technique. However, in view of the d^8 electronic configura-

tion, and the X-ray structure of $[\text{RhCl}(\text{PPh}_3)_3]$, $[\text{RhBr}(\text{PPh}_3)_3]$ is considered to be approximately square planar.

This technique and results reported herein demonstrate the practical application of EXAFS to the solution of structure problems, as a supplement to X-ray crystallography.

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