UVRR Spectroscopy of the Metal Receptor Site in MerR

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Although cysteine coordination to d10-metal transition-metal ions is observed in an increasing number of metal detoxification¹ and metal chaperone proteins,² there are still relatively few methods for probing such coordination environments. These metal centers have a closed-shell electronic structure and thus exhibit few spectroscopic handles. The d¹⁰-metals bound to cysteinate ligands in proteins exhibit broad and intense ligand-to-metal charge transfer (LMCT) transitions in the 200-300 nm region.³ We demonstrate here that it is possible to use ultraviolet (UV) laser excitation into these LMCT transitions of d¹⁰-thiolate centers in metalloproteins to obtain resonance enhancement of metalligand Raman bands. UV resonance Raman (UVRR) spectra are presented for a trigonal mercuric complex of an aliphatic thiolate $[Et_4N][Hg(SBu^t)_3]$ (1), and for the Hg–MerR metalloregulatory protein, a heavy metal receptor component of an Hg(II)-responsive genetic switch. Our spectroscopic data indicate that Hg(II) is trigonally coordinated in Hg-MerR and give support to results of EXAFS data,⁴ site-directed mutagenesis studies,⁵ and UV absorption,^{3,6} most recently by ¹⁹⁹Hg NMR.⁷ These results indicate that UVRR is a powerful probe of metalloprotein active sites containing d¹⁰-thiolate centers.

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(8) UVRR spectra were excited with the lines of an intracavity doubled $\rm Ar^+$ laser (Coherent Innova 300 FRED). Laser power was between 2 mW (229 nm) and 10 mW (257 nm excitation). A 135° backscattering geometry was used, and a polarization scrambler was placed in front of the monochromator. The scattered light was collected using a 1.29 m single monochromator (SPEX 1269) equipped with a diode array detector (Princeton Instruments). Raman intensities, expressed as cross sections, σ , were determined from integrated intensity ratio, I_N/I_S , of the sample (N) and internal standard (S) and Lorentzian-type model bands and including linear baseline in the optimization process.

(9) ¹⁹⁹Hg NMR spectroscopic data for a 5 mM solution of Hg(SBu^t)₃⁻ in acetonitrile indicate that the predominant species in solution is three-coordinate, and contributions to Raman band intensifies arising from dissociated ligands are considered to be negligible for solutions of equal or higher concentrations. Natan, M. J.; Millikan, C. F.; Wright, J. G.; O'Halloran, T. V. J. Am. Chem. Soc. 1990, 112, 3255–3257.

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Figure 1. UVRR spectra of 1 (26 mM, acetonitrile, 229 nm excitation, top) and of Hg-MerR (238 nm excitation, bottom). Solvent and quartz bands are labeled as (s) and (q), respectively.

Table 1. Absolute Intensities^{*a*} for Bands of [Et₄N][Hg(SBu^{*t*})₃] Solution^b and for the 288 cm⁻¹ Band of Aqueous Hg-MerR Solution^c

exc. wavelength [nm]	σ_{207}	σ_{341}	σ_{588}	σ_{820}	σ_{288} Hg-MerR
229	76	32	42	53	83 (89)
238	42	28	5	14	81 (83)
244	45	22	0	0	117
257	37	14	0	0	115
676^{d}	0.002	0	0.002	0.008	

^a Expressed in units of cross section, [millibarns/(molecule steradian)], 1 barn = 10^{-24} cm². ^b Acetonitrile bands at 379 and 918 cm⁻¹ as internal standard (ref 2). ^c 932 cm⁻¹ band of 0.2 M NaClO₄ added as internal standard.²⁰ Values in brackets based on tyrosine modes located at 1176 (ν_{9a}), 1205 (ring-C) and 1613 cm⁻¹ (ν_{8a}) with previously determined absolute intensities.²¹ *d* Kr⁺ (Coherent), Spex 1877 triple spectrometer equipped with an intensified diode array detector (Princeton Instruments).

The 229-nm excited UVRR spectrum of **1** (Figure 1, top),^{8,9} prepared by a literature method,¹⁰ contains bands at 820, 588, 341, and 207 cm⁻¹, in addition to background features due to the quartz sample tube [marked q], and an acetonitrile solvent band at 380 cm⁻¹. The bands at 588 and 820 cm⁻¹ are assigned to the C-S and the C-C stretching mode, respectively, by comparison with spectra and assignments of pure liquid tert-butanethiol.^{11,12} The band at 207 cm⁻¹ is assigned to the symmetric HgS stretching mode.⁹ The band at 341 cm⁻¹ is assigned to an internal ligand mode, most likely CCS bending, since bands at similar positions are seen in solid-state Raman spectra of Hg(SBut)2 and the

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Figure 2. UV-absorption spectrum (top) and absolute Raman intensities for bands (lower panels) of 1.

corresponding Cd(II) and Zn(II) compounds, and they were assigned to a mode of the SBu^t group.¹³ An alternative assignment to the asymmetric Hg–S stretching mode is ruled out by the excessive separation, 134 cm⁻¹, from the symmetric stretching frequency. For trigonally coordinated HgX₃⁻ in tri-*n*-butyl phosphate solution (TBP) the separation between v_s and v_{as} Hg-X is less than 25 cm⁻¹ for X = Cl, Br, I. Moreover, the v_{as} Hg–S of **1** has been assigned to infrared bands³ at 208 (298 K) and 212 (125 K) cm⁻¹, essentially coincident with v_s Hg–S (which is IR-forbidden).

Raman intensities were measured relative to solvent bands as a function of excitation wavelength (Table 1), and the resulting excitation profiles (Figure 2) are consistent with enhancement of both Hg–S and internal ligand modes in resonance with the UV absorption bands between 230 and 280 nm (43500–35700 cm⁻¹), which have previously been assigned to $S \rightarrow$ Hg LMCT transitions.³ Thus, the LMCT state is displaced along ligand coordinates, as well as the metal–ligand stretching coordinate, as has previously been observed for Fe–S^{14a} and Cu–S^{14b} proteins.

The UVRR spectrum of Hg–MerR at 238 nm excitation (Figure 1, bottom) exhibits sample bands at 288 and 852 cm⁻¹,

which are assigned to the symmetric Hg–S[cys] stretching mode and to the ν_1 mode of tyrosine, respectively.¹⁵ MerR was purified¹⁶ and its Hg(II) derivative made as previously described.¹⁷ The absolute Raman intensity of the 288 cm⁻¹ band is essentially the same as that of the 207 cm⁻¹ band of **1**, when measured at 229 nm (Table 1), but is larger at longer wavelengths. LMCT transitions of Hg–MerR have previously been reported in the 200–300 nm region of absorption difference spectra.⁵

Why is the ν_s HgS frequency so much lower (81 cm⁻¹) for **1** than for Hg-MerR, even though the average Hg-S bond lengths are the same (244 pm)?^{3,17} The answer lies in the large effective mass of the SBut ligand. For [Hg(SMe)₃]⁻ dissolved in ethanol a Raman band at 282 cm⁻¹ has been assigned to ν Hg–S, much closer to the Hg-MerR frequency. The difference is that SMe⁻ has only H atoms attached to the Ca, while ButS- has much heavier CH₃ groups. Indeed, it has been observed that the frequencies scale as the square root of the total ligand mass for $[Hg(SMe)_3]^-$ and $[Hg(SBu^t)_3]^-$,¹¹ as well as for $[Au(SMe)_2]$ and [Au(SBu^t)₂].⁴ However, this simple behavior is deceptive because other thiolate ligands deviate from it. For example, the ν_s frequency is much higher for Hg(SBuⁿ)₂, 325 cm⁻¹, than for Hg- $(SBu^{t})_{2}$, 223 cm⁻¹, even though the ligand masses are the same. Moreover the Hg(SBuⁿ)₂ frequency is quite similar to those of the lighter homologues Hg(SPrⁿ)₂ [329 cm⁻¹] and Hg(SEt)₂ [304 cm⁻¹].¹⁷ Clearly the frequency depends not on the total ligand mass, but rather on the nature of the substituents on the C_a atom: three alkyl C atoms for SBut but one alkyl C atom and two H atoms for SEt, SPrⁿ, and Sbuⁿ.

This issue is important with respect to using the protein frequency as a gauge of coordination number. It is well-established that the ν_s frequency decreases as the coordination number increases. Thus, the frequency is 328, 296, and 265 cm⁻¹ for HgCl₂, HgCl₃⁻, and HgCl₄²⁻, in TBP solvent.¹⁷ Interestingly, the HgCl₂ and HgCl₃⁻ frequencies are close to those of the Hg(S–n-alkyl)₂ species on one hand and Hg(SMe)₃⁻, as well as Hg–MerR, on the other. Likewise, the frequency for an oligopeptide serving as a model for 2-coordinate Hg–S[cys] proteins,¹⁸ 327 cm⁻¹, is essentially the same as that of HgCl₂. Thus Cl⁻ serves as a good vibrational model for *n*-alkyl as well as methyl thiolates. However, the exact frequency obviously depends on the interaction of the Hg–S stretching coordinate with other coordinates of the thiolate ligands. This issue will be addressed in a separate vibrational analysis.¹⁹

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