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A Combined NRVS and DFT Study of Fe^{IV}=O Model Complexes: A Diagnostic Method for the Elucidation of Non-Heme Iron Enzyme Intermediates**

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Fe^{IV}=O intermediates have been shown to be key catalytic species in a growing number of mononuclear non-heme iron (NHI) enzymes. Various combinations of Mössbauer (Mb), electron-nuclear double resonance, magnetic circular dichroism (MCD), and extended X-ray absorption spectroscopy data exist on these species, providing electronic insight into and bonding descriptions of the Fe^{IV}=O unit.^[1] However, information regarding other chemically important features necessary for mechanistic insight (overall structure) is mostly lacking. Resonance Raman (rR) and IR spectroscopy can, in principle, provide such insight, but owing to selection rules, low absorption intensities (NHI species lack the intense Soret band), and other factors, experimental data are limited.[2] Herein, we present vibrational data obtained by nuclear resonance vibrational spectroscopy (NRVS) on three Fe^{IV}=O complexes interpreted by coupling with DFT calculations, with particular focus on understanding the low-energy spectral region. The data show systematic trends that provide molecular-level insight reflecting the Fe^{IV} ligand environment and reactivity. Computationally, these studies are extended to models of the Fe^{IV}=O species in the NHI enzymes HmaS and HPPD (see below) and show that Fe^{IV} species can be experimentally differentiated using NRVS.

The NRVS experiment employs synchrotron radiation tuned to the 14.4 keV ⁵⁷Fe nuclear Mb transition. ^[3–5] Inelastic

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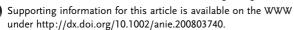
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scattering, where phonon annihilation and creation events couple with nuclear excitation, yields the Fe partial vibrational density of states (PVDOS) NRVS spectrum. This is analogous to Stokes and anti-Stokes rR scattering, but NRVS intensity is gained through Fe displacement (Δ Fe), via the mode composition factor, and provides selective enhancement of all Fe core modes with no background. [6] This method has been applied to iron–sulfur and heme enzymes, nitrogenase, and model complexes. [7-16] NRVS data for an FeV nitrido complex with a ligand similar to that in 1 (see below) have been reported. [16] However, detailed analysis in the lowenergy region was complicated by inhomogeneity in the sample. [16]

Fe^{IV}=O (S=1) functional models have been synthesized with 1,4,8,11-tetramethyl-1,4,8,11-tetraazacyclotetradecane (1); N_iN -bis(2-pyridylmethyl)-N-bis(2-pyridyl)methylamine (2); and N-benzyl- $N_iN'_iN'$ -tris(2-pyridylmethyl)-1,2-diaminoethane (3) ligands. [17-19] Each complex is six-coordinate: 1 has an equatorial (eq) ring with tertiary amines and an axial (ax) NCCH₃ ligand; 2 has eq pyridyl ligands (py) and a tethered ax tertiary amine, and 3 is quite rhombic, with two tertiary amine and three py ligands. Structures are available for 1 and 2. Note, 2 and 3 are more reactive than 1 in their ability to H-atom abstract from cyclohexane. [20]

The NRVS data for 1, 2, and 3 are shown in Figure 1. Isotope-sensitive peaks are observed at 831 (796 in ¹⁸O), 820 (788), and 824 (786) cm⁻¹ that can be assigned as v_2 , the FeO stretch (normal modes in Figure 2). [21,22] Resolution on NRVS energies is 8 cm⁻¹. These energies correlate well with the calculated energies and isotope shifts (see the Supporting Information, Figure S2). This mode has been observed only in 1 at 834 cm⁻¹ by IR spectroscopy and at 839 cm⁻¹ by rR spectroscopy, in good agreement with our results.[19,23] The comparable energies of v_2 , given the resolution of the NRVS experiment, imply that the total FeO bonding in these complexes are similar. Excited-state MCD data have shown the FeO π bond is stronger in 2 than in 1; therefore, the σ bond must be weaker. [24] This conclusion can be rationalized by considering the stronger donation of the eq ligands in 2 (see below), which donate into the donut of the d_{r^2} orbital and compete with the oxo ligand for σ bonding.

Moving down in energy, a weak peak is observed at 653 cm^{-1} in **2** and **3**, while for **1** the next resolvable peak is at 526 cm^{-1} . In **2** and **3** this peak is assigned, based on DFT calculations, as v_8 , the asymmetric FeN_{eq} stretches. In the calculations, these are shifted to the 450– 470 cm^{-1} region in **1** and are not resolved in the data. In total, four modes involve



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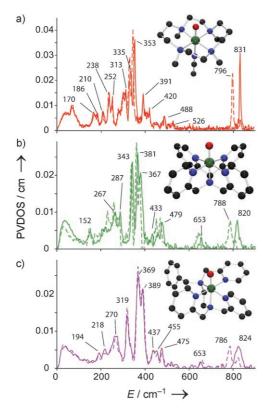


Figure 1. NRVS PVDOS for a) 1, b) 2, and c) 3 with ¹⁶O (solid lines) and ¹⁸O (dotted lines); insets show the 3D structures of complexes.

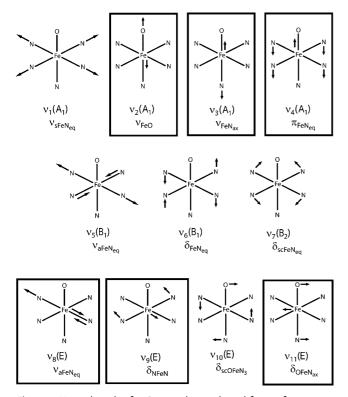


Figure 2. Normal modes for $C_{4\nu}$ complexes adapted from reference [21]. Modes with ΔFe are boxed. Nomenclature follows that of Nakamoto. [22]

FeN_{eq} stretching (including the NRVS-inactive v_1 and v_5) and probe eq bonding; the average energy of these sets is 355, 648, and 598 cm⁻¹, respectively. Calculations where the eq ring in 1 is cleaved (1', see the Supporting Information) show that the average FeN bond length decreases by 0.07 Å, the splitting of the four modes decreases, and the average energy increases to 430 cm⁻¹. These results show that the chelate in 1 influences the eq bonding but that the overall bond strength is determined by the greater σ donation of the py ligands relative to tertiary amines, which affects FeO bonding.

Continuing down in energy, the first intense feature is assigned as v_3 (FeN_{ax} stretch) at 391, 367, and 389 cm⁻¹, in **1**, **2**, and **3**, respectively. In the calculations, these bands show variable intensity (bars in Figure 3) and mix with v_{11} . The

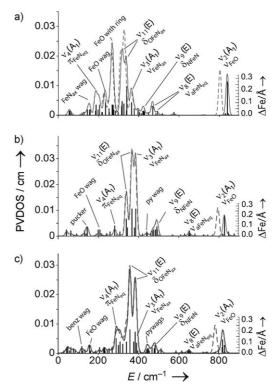


Figure 3. DFT calculated NRVS PVDOS for a) 1, b) 2, and c) 3 (16 O (solid lines) and 18 O (dotted lines)) with ΔFe included as bars.

resulting ΔFe has both in-plane (ip) and out-of-plane (op) contributions from v_{11} and v_3 . The ip motion in **1** is restricted by the chelate, thus reducing ΔFe , and consequently intensity. In **1'**, ΔFe is restored and v_3 shifts to 294 cm⁻¹ (Figure S4 in the Supporting Information). Calculations where the ax tether is removed in **2** (**2'**, see the Supporting Information) show that v_3 is shifted down in energy to 300 cm⁻¹, consistent with a 0.05 Å increase of the FeN_{ax} bond length, but the ip and op ΔFe are preserved. This mode reflects the bond *trans* to the oxo and constraints in the ligand environment.

The remaining two intense features are assigned as $v_{11a/b}$ (OFeN_{ax} bends), which are split due to orthogonal differences in the eq ligand environment and to ring size in **1**, ax tether in **2**, and eq chelate in **3**. The splitting is 50 cm⁻¹ in **3** but only approximately 25 cm⁻¹ in **1** and **2**, reflecting the contracted eq

chelate between a pair of cis nitrogen atoms in 3, and affords a probe of rhombic (perpendicular to the FeO bond) distortion. In $\mathbf{1'}$, the most symmetric model, the splitting is 6 cm⁻¹. In $\mathbf{2'}$ (where the x and y axes are still inequivalent), the energy positions and splitting are unchanged, but the calculated intensity is reduced because of increased N_{ax} motion. The splitting of v_{11} reflects the ligand environment perpendicular to the FeO bond and, combined with high intensity, should provide structural insight for Fe^{IV}=O intermediates.

(4-Hydroxyphenyl)pyruvate dioxygenase (HPPD) and (4hydroxy)mandelate synthase (HmaS) are NHI enzymes containing a 2-His-1-Glu facial triad and use the same substrate ((4-hydroxyphenyl)pyruvate, HPP) but exhibit different reaction mechanisms (electrophilic aromatic attack and H-atom abstraction, respectively) dependent on the orientation of HPP in the protein pocket. [25] The first step in both reactions is decarboxylation of HPP, and computational studies on the resultant Fe^{IV}=O site show that two 5C geometries are reasonable: square pyramidal (sqp, 4) with the oxo ligand in the eq plane, and trigional bipyramidal (tbp) with the oxo ligand axial (5).[25] DFT calculations show that NRVS can discriminate between these proposed structures. The calculated PVDOS are shown in Figure 4. The intense

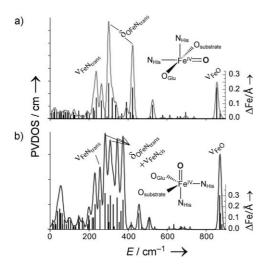


Figure 4. DFT calculated NRVS PVDOS for 4 and 5, with ΔFe included as bars.

peaks at 301 and 424 cm⁻¹ in 4 are bends of the OFeN_{trans} unit (trans to the oxo ligand) reminiscent of $v_{11a/b}$, where the large splitting arises from loss of the cis ligand. Intense peaks are also at 237 and 837 cm⁻¹ which involve FeN_{trans} (reminiscent of v_3) and FeO (reminiscent of v_2) stretches. For 1, 2, and 3, v_3 was observed at a higher energy than v_{11} ; the shift to lower energy in 4 is due to the increased effective mass of histidine. The NRVS data of 5 are distinctly different and can be understood considering the normal modes for a tbp complex (Figure S6 in the Supporting Information). The degenerate v_6 and v₇ pairs involve trans ax bends and cis metal-ligand stretches. These are highly mixed in 5 and produce four peaks with similar intensity. The trans N_{His} stretch (reminiscent of v_3) is also intense at 257 cm⁻¹. In summary, with the exception of the FeO stretch, the intense modes occur at low energies and,

because of the inverse energy dependence of NRVS intensity, these are likely to be detectable in enzyme intermediates and afford a mechanism for structural characterization.

Analysis of the ground-state NRVS vibrational data coupled with excited-state data from previous MCD studies provides detailed bonding descriptions of 1, 2, and 3. [24,26] The increased reactivity of 2 and 3 relative to 1 correlates with the strength of the FeO π bond (oxo p character in the π * LUMO, the frontier molecular orbital) and is determined by equatorial ligand σ donation. Additionally, the v_3 and v_{11} modes show systematic trends that reflect the ligand environment. Lowerenergy modes have generally not been used for geometric insight, but in NRVS spectra these are enhanced in intensity relative to higher-energy stretches (e.g. characteristic metalligand modes such as v_{FeO}) and are more likely to be detected in intermediates. This study uses these modes for electronic structural insight on Fe^{IV}=O models and computationally evaluates their utility for determination of active site structure in NHI enzyme intermediates.

Experimental Section

Samples were prepared as previously described. [17,18] NRVS was collected on beamline 3-ID at Advance Photon Source on multiple visits. [6,7,27] Spectra were collected from -25 to 125 meV in 0.25 meV steps. Five to 12 scans were averaged and normalized from which the PVDOS was generated using the PHOENIX program. [28] Spinunrestricted DFT calculations were performed using Gaussian 03 (see the Supporting Information).

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