

Supporting Information

for

High-Resolution EXAFS Provides Evidence for a Longer Fe•••Fe Distance in the Q Intermediate of Methane Monooxygenase

George E. Cutsail III,[†] Rahul Banerjee,^{‡,§} Ang Zhou,^{§,#} Lawrence Que, Jr.,^{§,#} John D. Lipscomb,^{‡,§} Serena DeBeer^{†,*}

[†]Max Planck Institute for Chemical Energy Conversion, Stiftstr. 34, D-45470 Mülheim an der Ruhr, Germany

[‡]Department of Biochemistry, Molecular Biology and Biophysics, 321 Church St. SE, Minneapolis, MN 55455

[§]Center for Metals in Biocatalysis, University of Minnesota, Minneapolis, Minnesota 55455, USA.

[#]Department of Chemistry, University of Minnesota, 207 Pleasant Street SE, Minneapolis, MN 55455

*Correspondence to: serena.debeer@cec.mpg.de

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Fig. S1. Mössbauer spectrum of MMOH^{red} complexed with MMOB collected at 80 K.

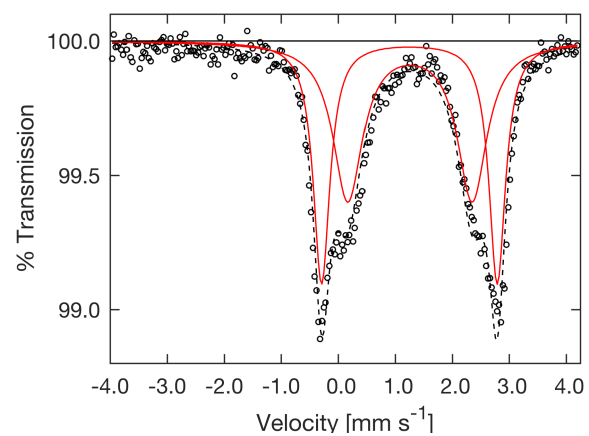


Fig. S2. Mössbauer spectrum of MMOH^{ox} complexed with MMOB collected at 80 K.

A small diiron(II) component (10%), of unoxidized MMOH^{red} is observed and fit as a single quadrupole doublet, magenta.

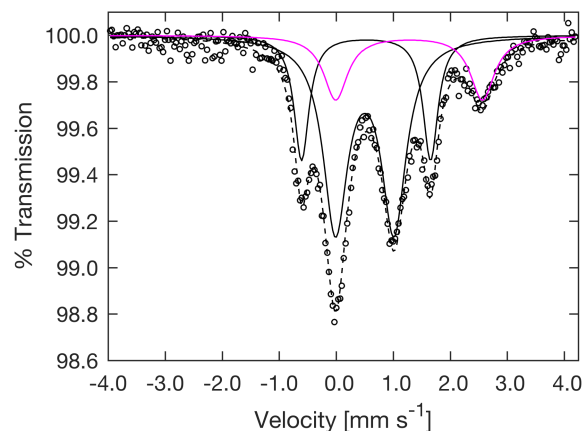
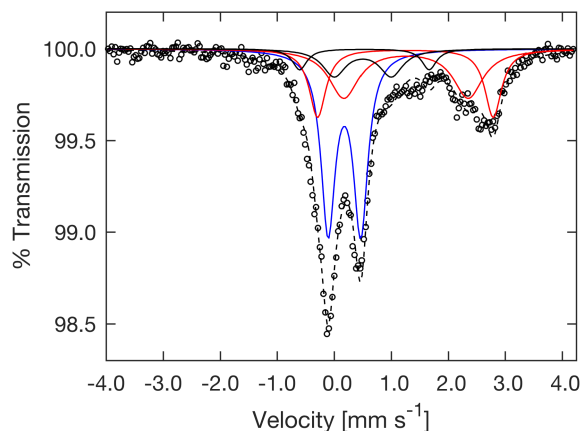


Fig. S3. Mössbauer spectra of MMOH-RFQ (powder packed) collected at 80 K.

Red: MMOH^{red}; Black: MMOH^{ox}; Blue: MMOH-Q

Sample 1:



Sample 2:

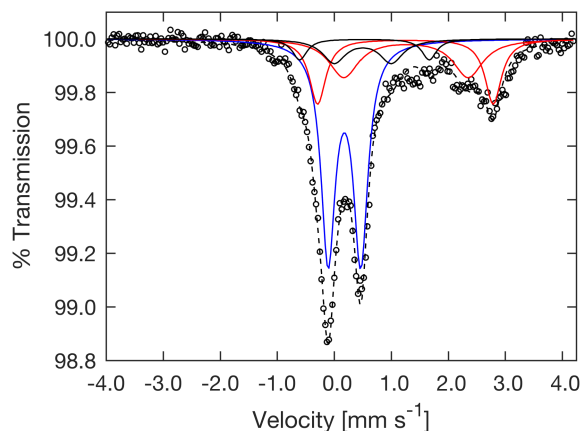
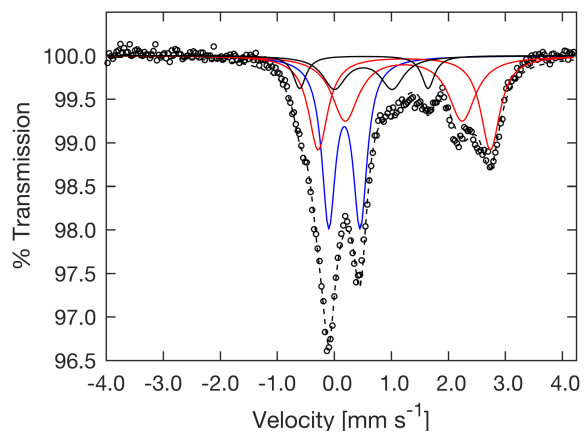


Fig. S4. Mössbauer spectra of MMOH-RFQ (frozen) collected at 80 K.

Red: MMOH^{red}; Black: MMOH^{ox}; Blue: MMOH-Q

Sample 1:



Sample 2:

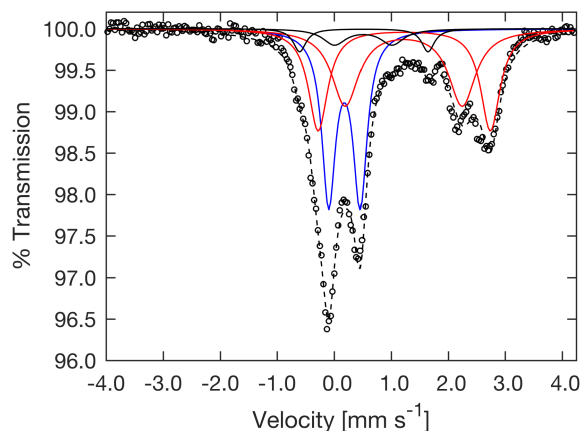


Table S1. Isomer shift (δ) and quadrupole splitting (ΔE_Q) values of the three diiron components within MMOH-RFQ samples.

Isomer shifts and quadrupole splitting experimental values were determined by fitting spectra with Lorentzian lines through the use of the software mf developed by Dr. Eckhard Bill.

	Reduced		Oxidized		Q	
δ (mm s ⁻¹)	1.25	1.26	0.53	0.50	0.18	
ΔE_Q (mm s ⁻¹)	2.18	3.08	2.27	1.00	0.56	
Component %	39.5 (36, 43)		14.5 (14, 15)		46 (50, 42)	Powder (Fig. S4) (MMOH-RFQ-1)
	54 (51, 57)		13 (17, 9)		33.5 (33, 34)	Frozen (Fig. S3) (MMOH-RFQ-2)

Fig. S5. $K\alpha$ -HERFD XAS of MMOH^{red} , MMOH^{ox} , MMOH-RFQ , Q and previously published Q.

The ‘pure’ Q spectrum was obtained by deconvolution of the MMOH-RFQ by subtraction of MMOH^{red} and MMOH^{ox} (complexed with MMOB) based on the Mössbauer quantification (Table S1) compared with the published $K\alpha$ -HERFD XAS of MMOH-Q from R. G. Castillo, et. al. (Q-RCG).¹

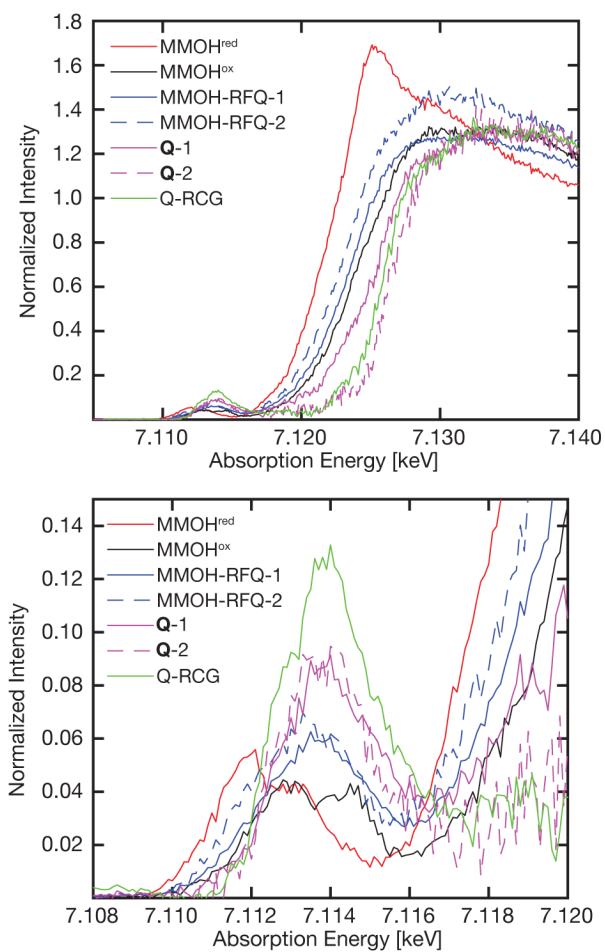


Fig. S6. Fe-K α emission spectra of MMOH^{red}, MMOH^{ox}, MMOH-RFQ and MMOH-Q.

The ‘pure’ MMOH-Q spectrum is the renormalized spectrum of MMOH-RFQ after quantitative subtraction of the MMOH^{red} and MMOH^{ox} components (complexed with MMOB) based on the Mössbauer quantification. All K α X-ray emission spectra were collected non-resonantly with an incident energy of 7800 eV.

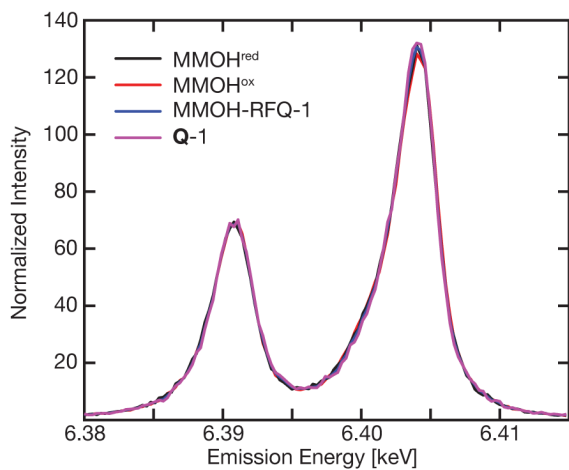
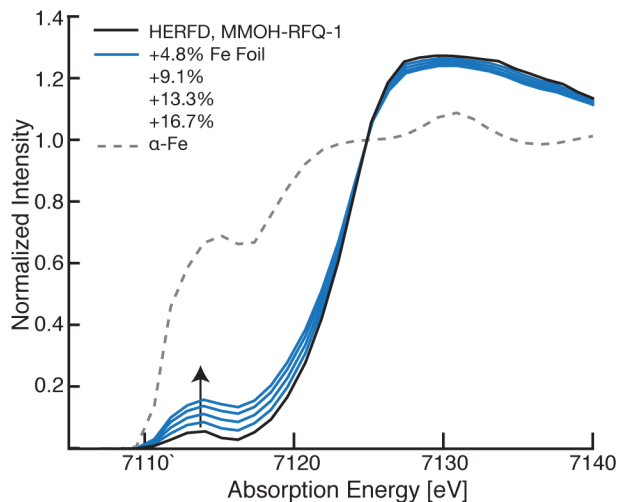


Fig. S7. $K\alpha$ HERFD-XAS of MMOH-RFQ with digital α -Fe contamination

The edge position of MMOH-RFQ is not dramatically shifted to lower energy upon the addition of $\sim 5\%$ to $\sim 17\%$ α -Fe to the spectrum. However, the pre-edge shape and intensity is strongly modulated. The pre-edge intensity increases over the entire ~ 7111 - 7118 eV range, resulting in a pre-edge that is no longer defined by a single peak, but rather a “shelf-like” plateau. This unusual pre-edge shape is the most diagnostic observable in the XAS data for metal contamination.



EXAFS fitting.

Theoretical EXAFS spectra were calculated using Artemis² utilizing the multiple scattering FEFF6 code.³ The EXAFS amplitude, $\chi(k)$, is given by

$$\chi(k) = \sum_R S_0^2 N \frac{|f_{\text{eff}}(k)|}{kR^2} \sin(2kR + \phi_k) e^{-2kR/\lambda_k} e^{-2\sigma^2 k^2}$$

where S_0^2 is the overall many-body amplitude factor, N , is the degeneracy of the paths, $|F_{\text{eff}}(k)|$ is the effective scattering amplitude, and R is the absorber-scatterer distance. A Debye-Waller like factor, $\exp(-2\sigma^2 k^2)$ is also included to account for disorder. Lastly, λ_k is the mean free path of the photoelectron and ϕ_k is the total photoelectron wave phase shift for the interaction between the absorber and the scatterer.

Models of MMOH^{red} and MMOH^{ox} were generated from the PDB deposited crystal structures 1FYZ and 1FZ1,⁴ respectively, by selection of all atoms within 8 Å of the diiron active site through the use of PyMOL.⁵ EXAFS scattering paths were calculated with FEFF6² and employed the FEFF cards: SIG2 0.001; RMAX 5; NLEGS 4.

Individual scattering paths were selected and fit within Artemis from paths calculated by FEFF6. The Fourier-transform spectrum of each were fit over a range of $R = 1.0$ to 3.4 Å (non-phase shift corrected). The FT is the product of a transform of k^3 -weighted EXAFS spectrum with a Hann window over the range of $k = 2$ to 11 Å⁻¹. By grouping similar scattering paths of a common coordination shell and increasing its degeneracy, N , the number of variables used for that coordination shell is minimal, 2 variables: σ^2 and ΔR . A single ΔE_0 variable is used for all paths in a given fit. S_0^2 was set to 0.9 for all paths. The E_0 for MMOH^{red} and MMOH^{ox} were set to 7122.17 and 7125.12 eV, respectively and MMOH-RFQ-1 and -RFQ-2 samples were 7124.10 and 7123.25 eV, respectively. Goodness of final fits were evaluated by their F -value, defined below:

$$F = \frac{\sum_{i=1}^n (d_i - f_i)^2}{\sum_{i=1}^n d_i^2}$$

where d and f represent the raw data and the fit, respectively. All reported F -values were calculated over an expanded R range of $R = 0$ to 4 Å.

Fig. S8. Additional HERFD-EXAFS fits of MMOH^{ox}

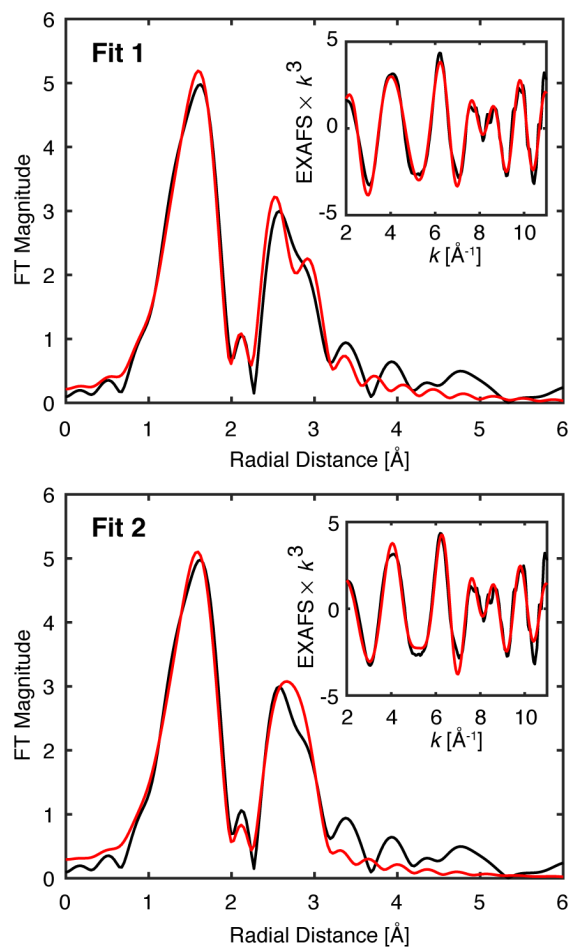


Table S2. HERFD-EXAFS Fit Parameters of MMOH^{ox} for Fig. S8

Fit 1					
N	Scattering Path	R (Å)	+/-	σ^2 ($\times 10^{-3}$ Å²)	+/-
5	Fe-O/N	2.057	0.010	12.6	1.0
1	Fe-O/N	2.480	0.036	9.9	5.7
1	Fe-Fe	3.056	0.011	1.3	0.9
3	Fe-C	3.304	0.031	4.9	4.1
$\Delta E_0 = + 2.51$ eV				$F = 0.163$	
Fit 2					
5	Fe-O/N	2.053	0.014	12.4	1.2
1	Fe-O/N	2.470	0.043	9.9	6.8
1	Fe-Fe	3.048	0.012	2.5	0.9
$\Delta E_0 = + 1.75$ eV				$F = 0.184$	

Fig. S9. Comparison of PFY MMOH^{red} with phase-shift recorection.

FT-EXAFS comparisons of HERFD- MMOH^{red} with the previously published PFY- MMOH^{red} from Shu, et al, subtracting the reported 0.35 \AA phase-shift (top panel), with a subtraction of 0.50 \AA phase-shift.

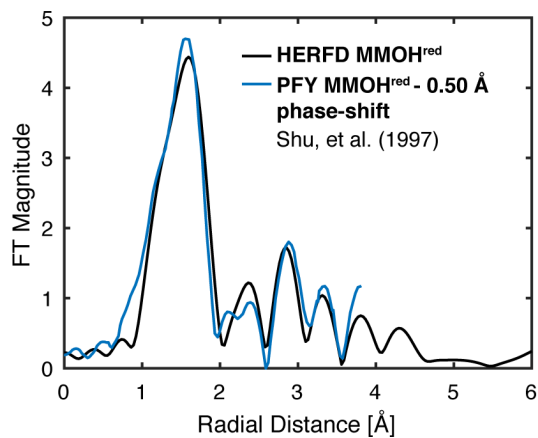
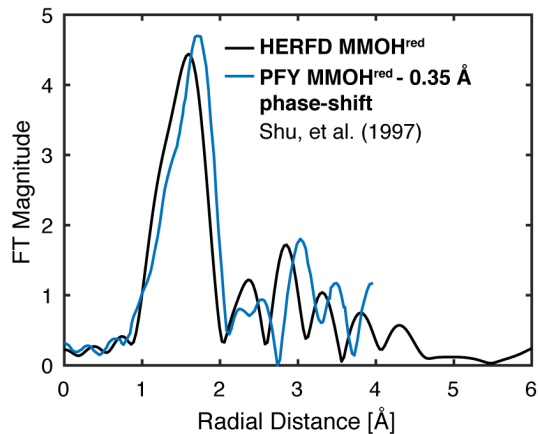


Fig. S10. Additional HERFD-EXAFS fits of MMOH^{red}

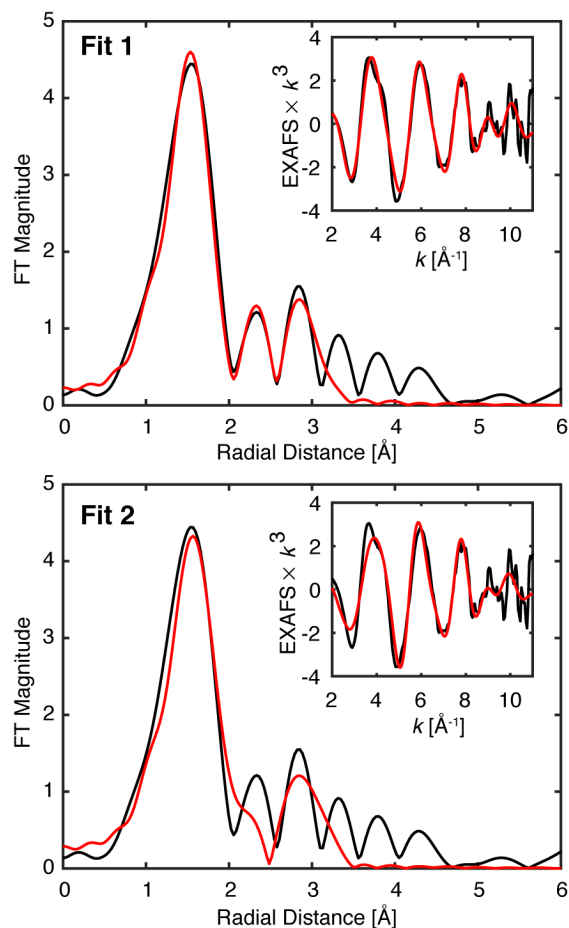


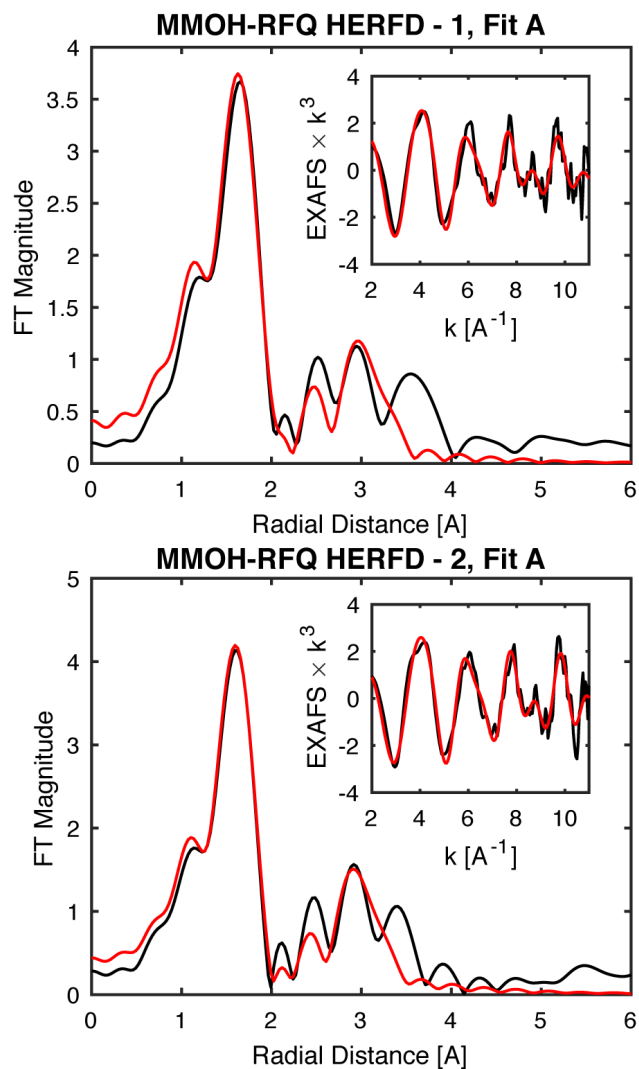
Table S3. HERFD-EXAFS Fit Parameters of MMOH^{red} for Fig. S10

Fit 1					
N	Scattering Path	R (Å)	+/-	σ^2 ($\times 10^{-3}$ Å²)	+/-
4	Fe-O/N	2.044	0.015	10.3	1.1
1	Fe-O/N	2.538	0.022	2.6	2.6
1	Fe-Fe	3.270	0.026	8.4	2.5
3	Fe-C	2.989	0.051	20.0	8.0
$\Delta E_0 = -6.63$ eV				$F = 0.489$	
Fit 2					
4	Fe-O/N	2.062	0.026	10.3	1.5
1	Fe-O/N	2.542	0.035	3.7	4.2
1	Fe-Fe	3.306	0.041	9.2	4.1
$\Delta E_0 = -4.001$ eV				$F = 0.683$	

Table S4. HERFD-EXAFS Fit Parameters of MMOH-RFQ (Attempts to fit short diiron distance $\sim 2.8 \text{ \AA}$).

Fit A					
MMOH-RFQ HERFD – 1					
N	Scattering Path	R (Å)	+/-	$\sigma^2 (\times 10^{-3} \text{ \AA}^2)$	+/-
5	Fe-O/N	2.053	0.016	13.7	1.2
1	Fe-O/N	1.780	0.020	7.5	2.4
0.5	Fe-Fe	2.668	0.040	14.6	5.4
0.5	Fe-Fe	3.385	0.020	3.0	2.1
$\Delta E_0 = -2.002 \text{ eV}$				$F = 0.41487$	
MMOH-RFQ HERFD – 2					
5	Fe-O/N	2.035	0.015	12.0	1.0
1	Fe-O/N	1.759	0.018	7.14	2.4
0.5	Fe-Fe	2.649	0.051	16.7	7.2
0.5	Fe-Fe	3.337	0.016	1.34	1.5
$\Delta E_0 = -4.186 \text{ eV}$				$F = 0.28673$	
Best Fit (as shown in Fig. 10, Table 3)					
MMOH-RFQ HERFD – 1					
N	Scattering Path	R (Å)	+/-	$\sigma^2 (\times 10^{-3} \text{ \AA}^2)$	+/-
1	Fe-O	1.781	0.018	8.3	2.3
5	Fe-O/N	2.057	0.014	13.8	1.0
0.3	Fe-Fe	3.067	0.025	1.9	2.4
0.7	Fe-Fe	3.343	0.030	8.0	4.2
$\Delta E_0 = -1.454 \text{ eV}$				$F = 0.35461$	
MMOH-RFQ HERFD – 2					
1	Fe-O	1.758	0.011	7.6	1.5
5	Fe-O/N	2.035	0.009	12.1	0.6
0.3	Fe-Fe	3.053	0.019	1.2	1.8
0.7	Fe-Fe	3.303	0.015	3.7	1.6
$\Delta E_0 = -4.060 \text{ eV}$				$F = 0.20059$	

Fig. S11. HERFD-EXAFS fits of MMOH-RFQ – 1 and 2, as Detailed in Table S4.



1. Castillo, R. G.; Banerjee, R.; Allpress, C. J.; Rohde, G. T.; Bill, E.; Que, L.; Lipscomb, J. D.; DeBeer, S., High-energy-resolution fluorescence-detected x-ray absorption of the Q intermediate of soluble methane monooxygenase. *J. Am. Chem. Soc.* **2017**, *139* (49), 18024-18033.
2. Ravel, B.; Newville, M., ATHENA, ARTEMIS, HEPHAESTUS: data analysis for X-ray absorption spectroscopy using IFEFFIT. *J. of Synchrotron Rad.* **2005**, *12* (4), 537-541.
3. Zabinsky, S. I.; Rehr, J. J.; Ankudinov, A.; Albers, R. C.; Eller, M. J., Multiple-scattering calculations of X-ray-absorption spectra. *Phys. Rev. B: Condens. Matter Mater. Phys.* **1995**, *52* (4), 2995-3009.
4. Whittington, D. A.; Lippard, S. J., Crystal structures of the soluble methane monooxygenase hydroxylase from *Methylococcus capsulatus* (Bath) demonstrating geometrical variability at the dinuclear iron active site. *J. Am. Chem. Soc.* **2001**, *123* (5), 827-838.
5. The PyMOL Molecular Graphics System, Version 1.8. **2015**.