Supporting Information

Coordination of the copper centers in particulate methane monooxygenase: comparison between methanotrophs and characterization of the Cu_C site by EPR and ENDOR spectroscopies

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Supporting Methods Preparation of [Cu(NH₃)₄]²⁺

 $Cu(NH_3)_4^{2+}$ was prepared in MilliQ water by adding 1 M NaOH to a solution of ~25 mM NH₄Cl and 2 mM Cu(II)Cl₂ until the pH was ~9 and the solution turned a deep blue color. A small aliquot was diluted to ~1 mM with a mixture of ethylene glycol and water.

DFT Calculations for $[Cu(NH_3)_4]^{2+}$ All density functional theory (DFT) calculations were carried out in ORCA 4.2.1.¹ Starting geometry for the copper complex was created in situ using Avogadro.² Geometry and single point energy calculations utilized spin unrestricted B3LYP/G hybrid functional set³⁻⁴ and Ahlrichs' valence triple- ξ with a polarization function basis set.⁵ For the proton hyperfine calculations Barone's EPR-III basis set⁶⁻⁷ was used, while the Cu(II) atom utilized Kutzelnigg's basis set (IGLO-III);⁸ B3LYP/G was again used as the functional set in combination with the accurate spin–orbit coupling operator [RI-SOMF(1X)].⁹

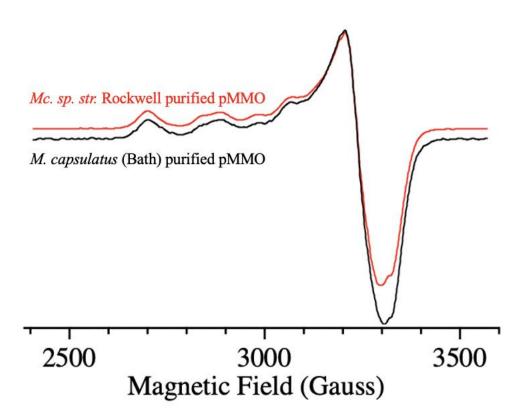


Figure S1. Overlay of X-band EPR spectra of purified *Mc*. sp. str. Rockwell (red) and *M. capsulatus* (Bath) pMMOs. The spectra are identical in terms of g and ⁶³Cu A_{II} values, the most definitive features for Cu(II) EPR. *Conditions:* 9.37 GHz, 20 K, 12.5 G modulation, 320 ms time constant, 200 μ W MW power. The spectra were collected on the same day for accurate comparison.

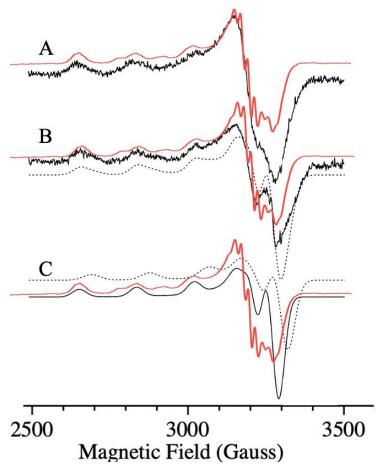


Figure S2. Overlay of X-band EPR spectrum of purified Mc. sp. str. Rockwell pMMO presented in this paper (red, from **Figure 2C** main text) with the three previously reported EPR spectra of type II Mc. sp. str. M pMMO¹⁰ (black), including (A) membrane-bound pMMO (B) solubilized pMMO, and (C) simulations of the two Cu(II) signals observed in (C) from ref. 10. The overlay of the Mc. sp. str. Rockwell spectrum (red) shows that the Cu_C site was not observed in the type II pMMOs in previous samples. Its absence may be due to not adding copper to the lysis buffer. As shown in Figure S4, this leads to only Cu_B being observed by EPR for Mc. sp. str. Rockwell pMMO. It is possible that there is some underlying Cu_C signal in (B), as evident by g_{\perp} , but due to the resolution, it is not obvious. *Conditions* are reported in ref. 10, with those for the red trace in the main text.

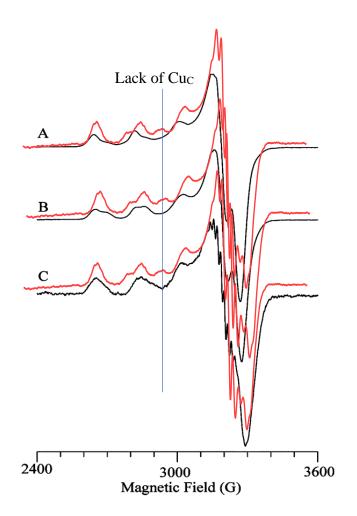


Figure S3. Overlay of purified *Mc*. sp. str. Rockwell pMMO X-band EPR spectrum presented in this paper (red, **Figure 2C** main text) with the three previously reported EPR spectra of type II *Methylosinus trichosporium* OB3b pMMO¹¹ (black), including (A) membrane-bound pMMO, (B) solubilized pMMO, and (C) purified pMMO The Cu(II) spectra in (A) and (B) are not related to pMMO since the g|| and the ^{63/65}Cu A|| do not match those of the Cu_B or Cu_C site. The purified pMMO spectrum appears to only have the Cu_B site. The lack of the Cu_C site in the purified pMMO spectrum is evident by the lack of the +1/2 hyperfine line, as shown with the blue vertical line. *Conditions* of black are described in the referece, red is described in text.

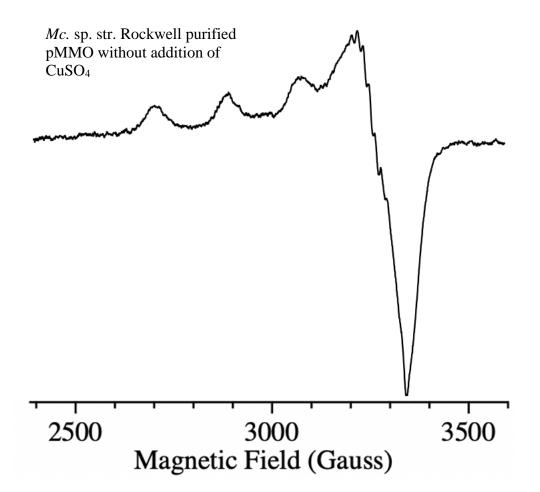
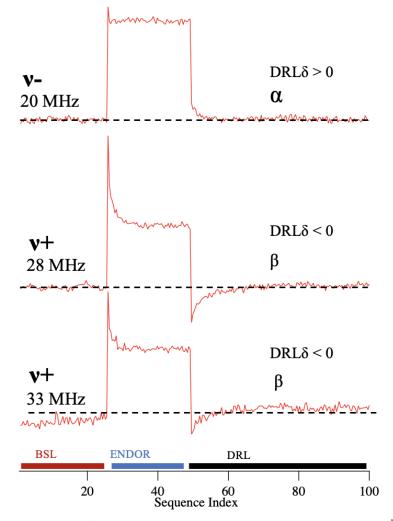
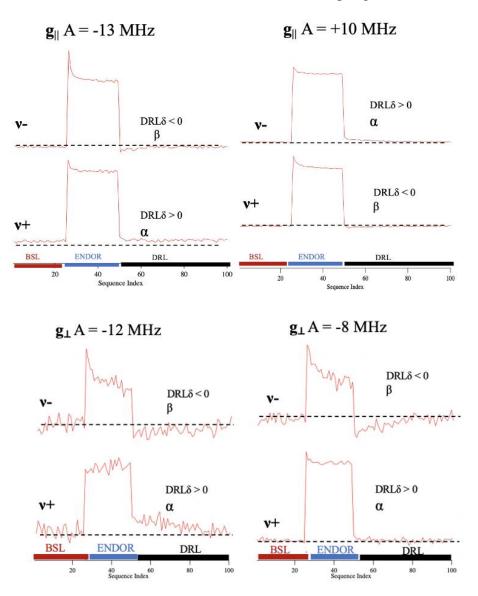


Figure S4. X -Band EPR spectrum of Mc. sp. str. Rockwell purified pMMO without the addition of CuSO₄ at lysis. The spectrum exhibits only the Cu_B signal, and is identical to that of Mc. sp. str. Rockwell whole cells (**Figure 2A**) and purified/reduced Mc. sp. str. Rockwell pMMO (**Figure 2D**), with no Cu_C signal. These samples do not have methane oxidation activity. *Conditions:* 77K, 12.5 G modulation, 10G/s sweep rate, 320 ms time constant, 200µW MW power, 9.34 GHz MW frequency.



PESTRE measurements for Cu_B ¹⁵N couplings

Figure S5. PESTRE measurements for sign determination of the strongly coupled Cu_B ¹⁵N in *Mc*. sp. str. Rockwell in vivo pMMO. The couplings were found to be negative in sign for both peaks shown in **Figure 3**. *Conditions:* 34.77 GHz, 100 ms repetition time, $\pi = 120$ ns, $\tau = 600$ ns, TRF = 15 µs, RF tail = 5 µs, 25 shots, 24 off.



PESTRE measurements for Cu_B ¹H couplings

Figure S6. PESTRE measurements on the large couplings of in vivo *Mc.* sp. str. Rockwell pMMO (Cu_B site) as resolved from the stochastic ¹H ENDOR experiment at both \mathbf{g}_{\perp} and \mathbf{g}_{\parallel} (**Figure 4**). The measurement resulted in A = -13 and +10 at \mathbf{g}_{\parallel} and A = -12 and -8 at \mathbf{g}_{\perp} . *Conditions:* 34.72 GHz, 100 ms repetition time, $\pi = 120$ ns, $\tau = 600$ ns, TRF = 15 µs, RF tail = 5 µs, 25 shots, 24 off.

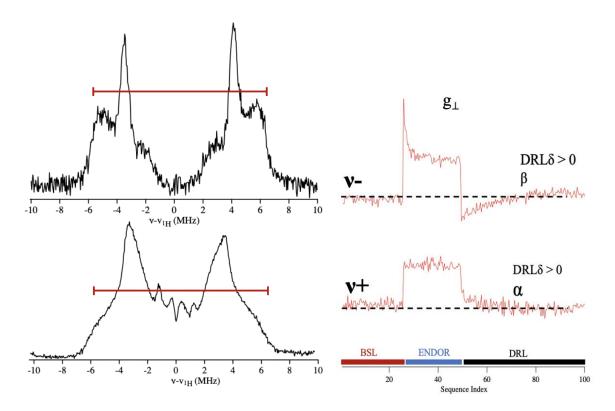
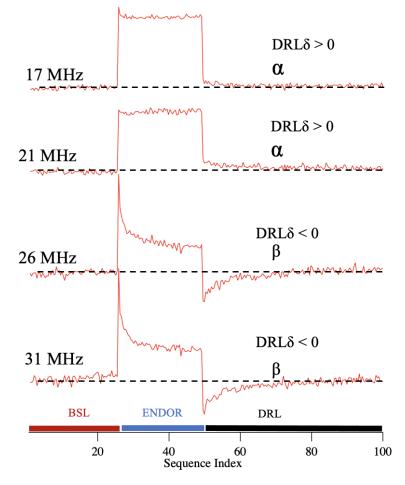


Figure S7. Q-band stochastic ¹H ENDOR and PESTRE measurements of Cu(II)(NH₃)₄ in H₂O buffer with ~20% ethylene glycol at g_{\parallel} (10800G) (top) and g_{\perp} (12020 G). Red brackets represent |A| for the largest ¹H couplings, A ~ 12 MHz. Since [Cu(II)(NH₃)₄]²⁺ does not contain axial ligands, the ~12 MHz ¹H couplings at both g_{\parallel} and g_{\perp} must come from the in-plane amine ligands. PESTRE measurements revealed the couplings at both g-values to be negative in sign. DFT model of [Cu(II)(NH₃)₄]²⁺yields a hyperfine tensor, **A** = [-15, 1.5, -16] for an in-plane amine proton. This highly rhombic tensor is achieved due to large spin delocalization from the copper to the protons of the amine when they are oriented within the x²-y² orbital of the unpaired electron. The tensor matches the signs and the identical relative magnitudes of those seen for the for the copper complex as well as for Cu_B.



PESTRE measurements for Cu_C ¹⁵N couplings

Figure S8. PESTRE measurements on Cu_C ¹⁵N couplings at $\mathbf{g} = 2.33$. The peaks examined at the frequency defined above are shown in **Figure 6**. The results of the PESTRE measurements reveal both strongly coupled ¹⁵N to the Cu_C to be negative in sign. *Conditions:* 34.71 GHz, 100 ms repetition time, $\pi = 120$ ns, $\tau = 600$ ns, TRF = 15 µs, and RF tail = 5 µs. 25 shots, 24 off.

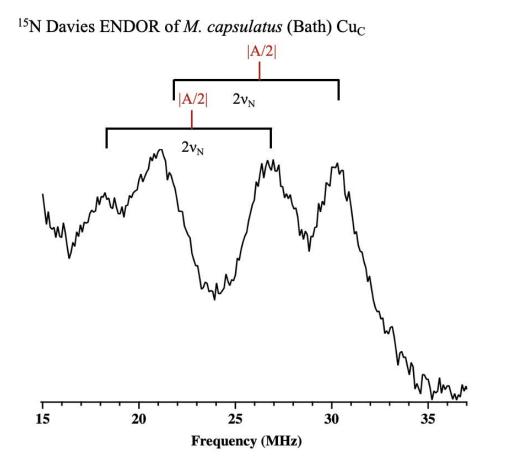


Figure S9. Medium Davies ¹⁵N ENDOR at a field of a corresponding g-value of g = 2.32 for ¹⁵N ⁶³Cu *M. capsulatus* (Bath) pMMO. The ⁵N couplings observed here are exclusive to the Cu_C site at this magnetic field and are identical in |A| to the type II *Mc.* sp. str. Rockwell pMMO, as shown in **Figure 6**, |A| ~ 50 MHz and |A| ~ 46 MHz. Brackets indicate twice the ¹⁵N Larmor frequency, and the center of the bracket (red) indicates |A/2|. *Conditions* are identical to **Figure 6** in the main text.

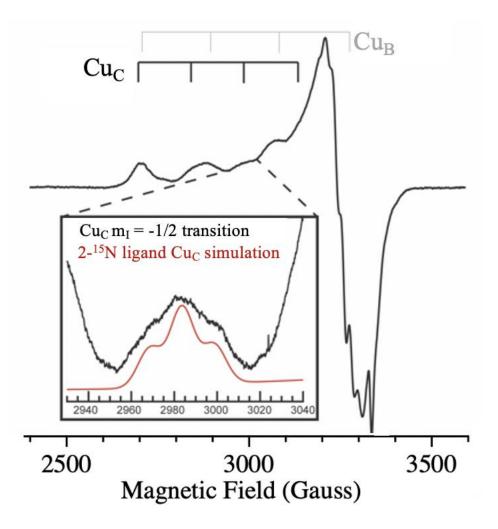


Figure S10. X-band CW EPR spectrum of the purified ¹⁵N, ⁶³Cu-labeled *M. capsulatus* (Bath) pMMO in D₂O and 10% D₃-glycerol exhibits overlapping Cu_B and Cu_C resonance. Brackets depict ⁶³Cu A_1 hyperfine splitting attributable to the two Cu(II) centers. Inset depicts baseline-corrected narrow scans collected on the g_{\parallel} Cu_C $m_I = -1/2$ transition (black) and simulation of Cu_C featuring two ¹⁵N ligands (red). *Conditions* for full spectrum: 9.365 GHz microwave frequency, 80 ms time constant, 10 G modulation amplitude, 90 s scan rate, temperature 180 K. *Conditions* for inset were the same except for a 60 s scan rate. *Simulation parameters*: $\mathbf{g} = [2.022, 2.068, 2.295]$, ⁶³Cu $A_{\parallel} = [-440]$ MHz, 2 isotropic ¹⁵N couplings $A_{iso} = -48$ MHz, 13 G full width at half maximum isotropic linewidth.

¹⁵N Davies ENDOR on *Mc.* sp. str. Rockwell Cu_C

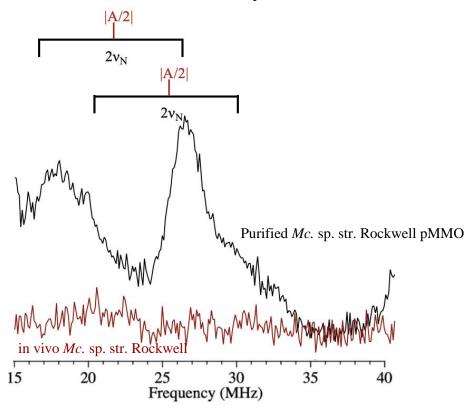


Figure S11. ¹⁵N Davies pulsed ENDOR of *Mc*. sp. str. Rockwell pMMO Cu_C (black trace) site at the low-field end of the EPR signal, at a field with a corresponding g-value of g = 2.34, (~10600 G). The red trace is in vivo *Mc*. sp. str. Rockwell pMMO, which exhibits an EPR signal from Cu_B only, at the same field position. The two spectra were echo-normalized. Brackets indicate twice the ¹⁵N Larmor frequency and the center of the bracket (red) indicates |A/2|. The larger coupling is not well resolved at this field. *Conditions* are identical to **Figure 6** in the main text.

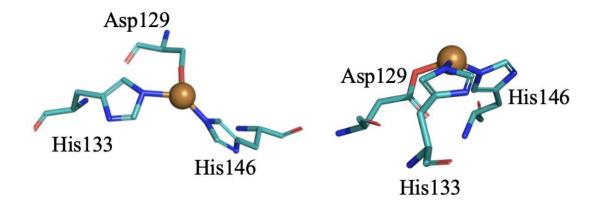


Figure S12. Alternative view of PmoC copper site from the crystal structure of *Mc*. sp. str. Rockwell (PDB accession code 4PHZ). The view shows the distortion (lack of symmetry) of the two His coordinating ligands to the copper center.

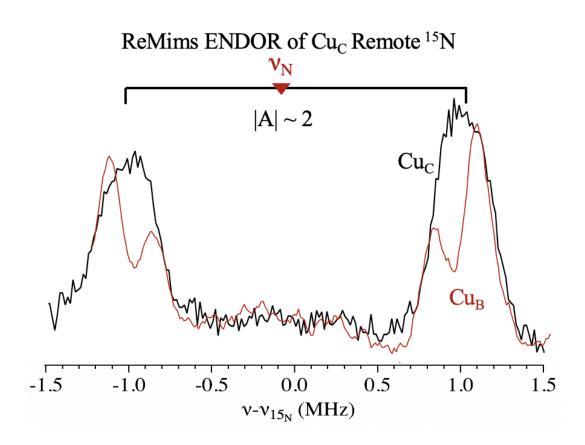
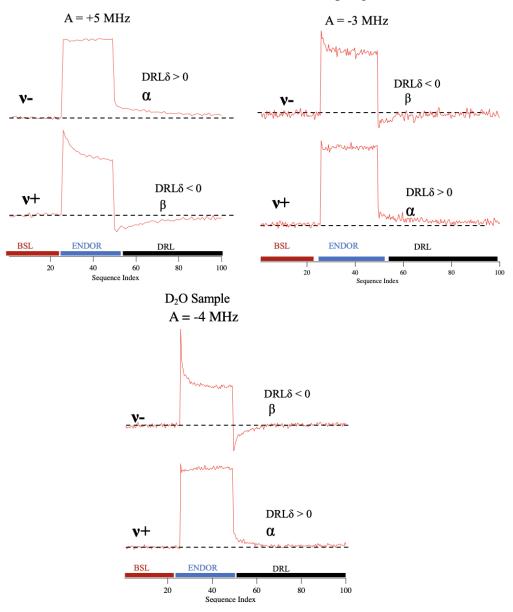


Figure S13. Overlay of ReMims ENDOR remote ¹⁵N couplings from the Cu_B (red, from **Figure 2**) and Cu_C (black, from **Figure 7**). Comparing the breadth of the unresolved doublet to the doublets in the Cu_B spectrum reveal that the couplings in the Cu_C spectrum likely arise from more than one remote ¹⁵N coupling as discussed in the text.



PESTRE measurements for Cu_C ¹H couplings

Figure S14. 35 GHz PESTRE performed on the Cu_C ¹H couplings at g = 2.34 (10700G). The +5 MHz and -2.5 MHz couplings were determined using a purified sample in H₂O, while the -4 MHz coupling was determined using a purified *Mc*. sp. str. Rockwell pMMO sample exchanged into D₂O. *Conditions:* 34.67 – 34.69 GHz, 100 ms repetition time, $\pi = 120$ ns, $\tau = 600$ ns, TRF = 15 µs, and RF tail = 5 µs. 25 shots, 24 off.

¹⁷O Mims ENDOR on *M. capsulatus* (Bath) Cu_C

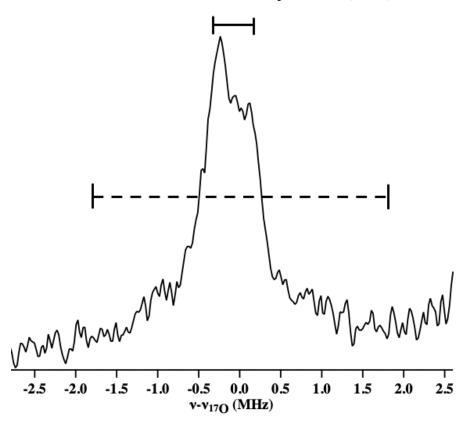


Figure S15. ¹⁷O ENDOR at g = 2.33 (Cu_C site). The spectrum specifically lacks the ~3.6 MHz doublet (indicated by the dashed 'goalposts') previously reported observing at g = 2.01 for Cu_C. The spectrum instead exhibits a response centered at the Larmor frequency, which is expected for remote H₂¹⁷O solvent in the sample, and may exhibit unresolved intensity, breadth ~ 2 MHz, from a secondary coordination sphere water, whose ¹H ENDOR response is discussed in Figure 8.

¹⁷O Mims ENDOR on purified *M. capsulatus* (Bath) pMMO

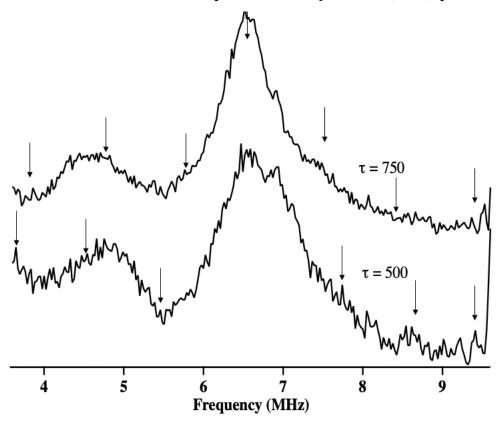


Figure S16. ¹⁷O Mims Pulsed ENDOR of purified *M*. capsulatus (Bath) pMMO in 45% enriched H₂¹⁷O buffer at g = 2.02 (~12050 G). This experiment was conducted at different τ values to observe whether there is suppression of the large coupling (3.5 MHz). The $\tau = 750$ experiment should suppress the coupling, but it looks quite resolved here. The coupling is now believed to derive from either a weakly coupled ¹⁴N or 5/2 quadrupole of the Cu_B axial H₂O ligand.

¹H CW ENDOR on *Mc.* sp. str. Rockwell Cu_C

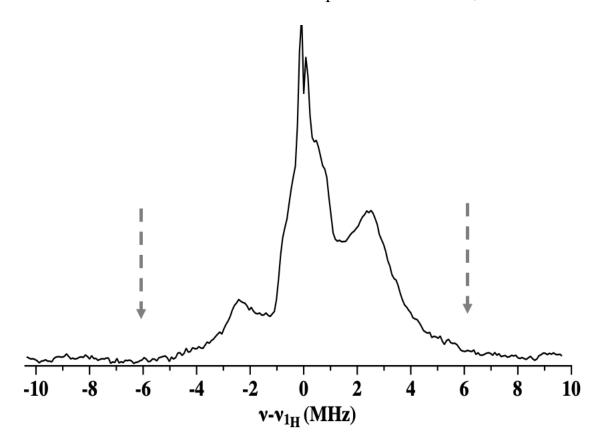


Figure S17. ¹H CW ENDOR spectrum at g = 2.35 of *Mc*. sp. str. Rockwell purified pMMO Cu_C site. The gray dashed arrows highlight the absence of any couplings with A >8 MHz, which signifies the absence of a bound axial solvent molecule. This also serves as an additional control that only the Cu_C site is observed at this field position since the Cu_B site exhibits couplings of ~12 MHz at g_{||}.

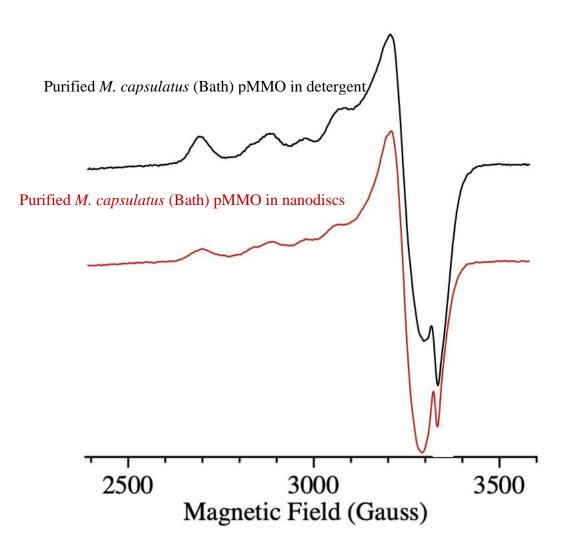


Figure S18. X-band EPR spectra of detergent-purified natural abundance *M*. capsulatus (Bath) pMMO (black) and purified *M*. capsulatus (Bath) pMMO in nanodiscs (red). The spectra are scaled by EPR intensity to facilitate comparison. The two sites exhibit identical g values and 63 Cu A values. *Conditions:* 9.37 GHz MW Frequency, 30 K, 12.5 G modulation, 10 G/s sweep rate, 320 ms time constant, 200µW power.

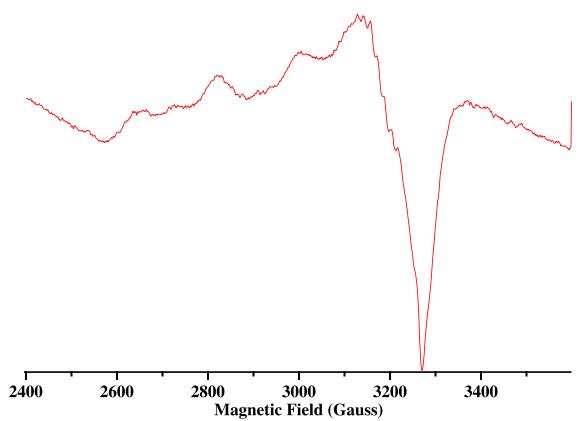


Figure S19. X-band EPR spectrum of reduced *M. capsulatus* (Bath) pMMO in nanodiscs. 12:1 ascorbic acid per 100 kDa protomer of pMMO was added to reduce the Cu_C site and obtain a pure Cu_B site to investigate with ENDOR. *Conditions:* 77K, 12.5 G modulation, <u>200µW</u> MW power, 320 ms time constant, 9.37 GHz frequency.

¹H CW ENDOR on nanodisc pMMO Cu_B

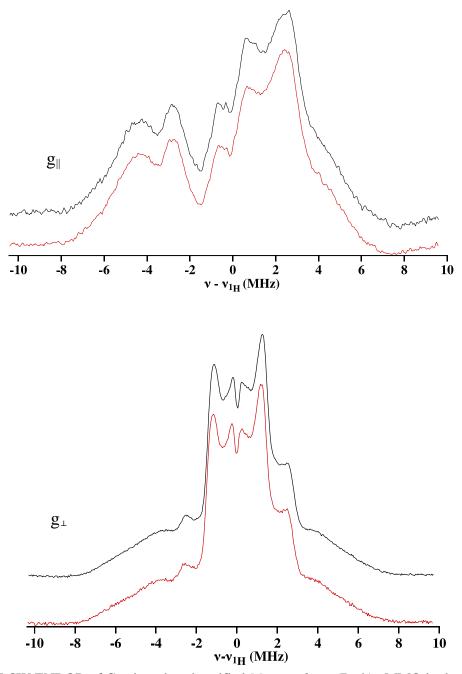


Figure S20. ¹H CW ENDOR of Cu_B in reduced purified *M. capsulatus* (Bath) pMMO in detergent (black trace) and in nanodiscs (red trace) at g|| (top) and g_{\perp} (bottom). The ENDOR spectra show no differences, indicating that Cu_B is identical within the two different environments, unlike Cu_C in which the secondary coordination sphere loses a solvent molecule (**Figure 9**). *Conditions:* 2K, 34.89 GHz MW frequency, 3.2 G modulation, 0.5 MHz/s scan rate, 30dB MW power, 13dB RF power, 32 ms time constant. Spectra were collected with RF swept low to high frequency.

References

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