

Peroxodiferric Intermediate of Stearoyl-Acyl Carrier Protein Δ^9 Desaturase: Oxidase Reactivity during Single Turnover and Implications for the Mechanism of Desaturation[†]

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ABSTRACT: Combined optical and resonance Raman studies have revealed the formation of an O₂-adduct upon exposure of 4e[−] chemically reduced stearoyl-acyl carrier protein Δ^9 desaturase to stearoyl-ACP and 1 atm O₂. The observed intermediate has a broad absorption band at 700 nm and is remarkably stable at room temperature ($t_{1/2} \approx 26$ min). Resonance Raman studies using ¹⁶O₂ gas reveal vibrational features of a bound peroxide [$\nu_s(\text{Fe}-\text{O}_2)$, 442 cm^{−1}; $\nu_{as}(\text{Fe}-\text{O}_2)$, 490 cm^{−1}; $\nu(\text{O}-\text{O})$, 898 cm^{−1}] that undergo the expected mass-dependent shifts when prepared in ¹⁶O¹⁸O or ¹⁸O₂. The appearance of two Fe–O₂ vibrations, each having a single peak of intermediate frequency with ¹⁶O¹⁸O, proves that the peroxide is bound symmetrically between the two iron atoms in a μ -1,2 configuration. The same results have been obtained in the accompanying resonance Raman study of ribonucleotide reductase isoform W48F/D84E [P. Moënne-Loccoz, J. Baldwin, B. A. Ley, T. M. Loehr, and J. M. Bollinger, Jr. (1998) *Biochemistry* 37, 14659–14663], thus making it likely that other members of the class II diiron enzymes form related peroxodiferric intermediates. Study of the reactivity of peroxodiferric Δ^9 D revealed that this intermediate underwent 2e[−] reduction leading to an oxidase reaction and recovery of the resting ferric homodimer. In contrast, biological reduction of the same enzyme preparations using ferredoxin reductase and [2Fe-2S] ferredoxin gave catalytic desaturation with a turnover number of 20–30 min^{−1}. The profound difference in catalytic outcome for chemically and enzymatically reduced Δ^9 D suggests that redox-state dependent conformational changes cause partition of reactivity between desaturase and oxidase chemistries. The Δ^9 D oxidase reaction represents a new type of reactivity for the acyl-ACP desaturases and provides a two-step catalytic precedent for the “alternative oxidase” activity recently proposed for a membrane diiron enzyme in plants and trypanosomes.

Enzymes containing oxo- or hydroxo- and carboxylato-bridged diiron sites are a functionally and structurally diverse group (1–3). Sequence alignments, catalytic properties, spectroscopic studies, and crystal structures support the division of this group into at least three, and perhaps four, different classes (4–8). The class II diiron enzymes include ribonucleotide reductase, RNR R2¹ (9), bacterial hydrocarbon monooxygenases (10, 11), and acyl-ACP desaturases (12). These enzymes have a similar protein fold that contains two

copies of the amino acid sequence (Asp/Glu)X_{~40}GluXXHis separated by ~100 amino acids, and catalyze O₂ activation as part of their respective reaction cycles. Although the protein folds are different, ferritins (7) and rubrerythrin (8) also contain diiron centers that are closely related to the other class II diiron enzymes, and catalyze O₂ activation as part of the ferroxidase reaction (13, 14).

Stearoyl-ACP Δ^9 -desaturase (Δ^9 D) is the best characterized acyl-ACP desaturase (15). This soluble, homodimeric enzyme ($M_r \approx 84$ kDa) catalyzes the NAD(P)H- and O₂-dependent introduction of a cis double bond between carbons 9 and 10 of stearoyl-ACP (18:0-ACP), yielding oleoyl-ACP (18:1-ACP) with a turnover number of 20–30 min^{−1}. Optical, Mössbauer (16), RR (4), and EXAFS (17) studies have shown that resting Δ^9 D contains 4 mol of iron in two oxo-bridged diferric centers, while an X-ray structure revealed that 4e[−] reduced Δ^9 D produced by photoreduction had lost the oxo bridge (12). Furthermore, RR studies of azido Δ^9 D revealed a μ -1,3 bridged species and the absence of the oxo bridge (18), which suggested that a μ -1,2 peroxy intermediate could be formed during Δ^9 D catalysis. The structural variations revealed by these² and related studies

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¹ Abbreviations: ACP, acyl carrier protein; 18:0-ACP, stearoyl-ACP; 18:1-ACP, oleoyl-ACP; Δ^9 D, 18:0-ACP Δ^9 desaturase; resting Δ^9 D, form of Δ^9 D containing all ferric sites; 4e[−] Δ^9 D, chemically reduced form of Δ^9 D containing all ferrous sites; MMOH, hydroxylase component of methane monooxygenase; RNR R2, iron-containing component of ribonucleotide diphosphate reductase; RR, resonance Raman.

of other diiron enzymes (10, 11, 19) suggest substantial flexibility in the carboxylate ligands (20).

The ability of the carboxylate ligands to permit rearrangements at the metal center concomitant with changes in redox state has emerged as a consistent feature of the proposed catalytic mechanisms of diiron enzymes (10, 18–23). Furthermore, reaction intermediates (24) including peroxodiferic (13, 25–27), bis- μ -oxo diiron(IV) (21, 27, 28), and ferric-ferryl mixed valence (29–32) have been recently identified. The previous paper in this issue describes the RR characterization of a peroxodiferic intermediate of RNR R2 isoform W48F/D84E (26). The vibrational data clearly demonstrate that this intermediate has a symmetric μ -1,2 structure, while other work has shown that the corresponding intermediate observed in the related RNR R2 isoform D84E decays with $k' \approx 0.59\text{--}0.93\text{ s}^{-1}$ to the yield the natural product, a radical at Y122 (25).

In this work, optical, RR, and catalytic studies of both biologically and chemically reduced $\Delta 9D$ are reported. During multiple turnover, a biological electron-transfer chain consisting of ferredoxin reductase and [2Fe-2S] ferredoxin is required to introduce minimally $2e^-$ to the resting ferric enzyme. No evidence for a mixed valence [Fe(II)Fe(III)] state has been obtained from either Mössbauer or EPR studies, suggesting that the biological electron-transfer chain will give $2e^-$ reduction of one diferic center of the resting enzyme to yield the diferrous state, while the other diiron center will remain in the diferic state. In contrast, chemical reduction produces an enzyme containing all ferrous sites as shown by Mössbauer spectroscopy (16), which will herein be called $4e^- \Delta 9D$. The present work reveals that $4e^- \Delta 9D$ can form a symmetric μ -1,2 peroxodiferic intermediate whose structure is remarkably similar to that of RNR R2 D84E (25) and W48F/D84E (26). However, the $\Delta 9D$ intermediate, which has only been detected by starting the reaction with $4e^- \Delta 9D$, 18:0-ACP, and O_2 , decays via an unanticipated $2e^-$ reduction of the peroxodiferic center (or a subsequent, undetected intermediate derived from this species) to restore the resting, completely ferric enzyme. The overall stoichiometry, $O_2 + 4e^- + 4H^+ \rightarrow 2H_2O$, corresponds to an oxidase reaction, which represents a new mode of catalysis for the diiron desaturases.

MATERIALS AND METHODS

Sample Preparations. $\Delta 9D$ was obtained as previously described (33), the protein concentration was determined by dye binding (34), and the iron content was determined by complexation with tripyridyl-*S*-triazine (35). Spinach holo-ACP isoform I (>95% phosphopantetheinylation) was produced by coexpressing a synthesized gene composed of codons favored for high-level expression in *Escherichia coli* with the gene for *E. coli* holo-ACP synthase (36). Holo-ACP was converted to 18:0-ACP (>95%) using *E. coli* acyl-ACP synthase (37) modified to contain a C-terminal His₆

tag. Holo- and acyl-ACP were quantitated by reaction of Ellman's reagent with the phosphopantetheine thiol group of either holo-ACP or holo-ACP formed by treating acyl-ACP with NH_2OH . Further details of the ACP preparations will be reported elsewhere. Reduction of anaerobic solutions of $\Delta 9D$ was accomplished by titration with concentrated solutions of buffered sodium dithionite (16). After reduction, anaerobic 18:0-ACP was added, and then the headspace of the anaerobic cuvette was replaced with O_2 (1 atm). Samples for Raman spectroscopy were also prepared with $^{16}O^{18}O$ (50 atom % ^{18}O , ICON, Summit, NJ) and $^{18}O_2$ (98 atom % ^{18}O , ICON). Raman analysis of the $^{16}O^{18}O$ gas showed the expected distribution of 25% $^{16}O_2$, 50% $^{16}O^{18}O$, and 25% $^{18}O_2$. Samples were frozen in liquid nitrogen immediately after preparation, and Raman spectra were obtained at room temperature within 10 min.

Spectroscopic Measurements. Optical spectra were obtained using a Hewlett-Packard 8452 diode array spectrometer at 20 °C. The rate of decay of peroxodiferic $\Delta 9D$ was determined by nonlinear least-squares fitting using the equation $A(t) = A(0)\exp(-k_{app}t) + A_{\infty}$. A_{∞} was determined by Kedzy-Swinbourne analysis (38). RR spectra were obtained using a custom McPherson 2061 spectrograph (0.67 m, 1800 groove grating) using a Kaiser Optical holographic super-notch filter with a Princeton Instruments liquid N_2 -cooled (LN-1100PB) CCD detector with 3 cm^{-1} spectral resolution. Excitation at 647.1 nm (40 mW) was provided by a Coherent Innova 302 Kr laser. All spectra were collected using a 90° scattering geometry on samples in capillaries cooled by N_2 gas that had been passed through an ice bath. Absolute frequencies were obtained by calibration with aspirin and CCl_4 and are accurate to $\pm 1\text{ cm}^{-1}$. Isotope shifts, obtained from spectra recorded under identical experimental conditions, were evaluated by abscissa expansion and curve resolution of overlapping bands and are accurate to $\pm 0.5\text{ cm}^{-1}$.

Product Determinations. Multiple turnover desaturation reactions were performed at 20 °C in Teflon-sealed vials containing 250 μL of 50 mM HEPES buffer, pH 7.8, 50 mM NaCl, 50 nmol of 18:0-ACP, 0.5 nmol of $\Delta 9D$, 2 nmol of *Anabaena* 7120 vegetative ferredoxin, 1.5 nmol of ferredoxin NADPH oxidoreductase, and 600 nmol of NADPH. After either multiple or single turnover reactions, acyl thioesters were converted to acyl alcohols using sodium borohydride (39) and extracted with chloroform. The acyl alcohols were converted to silyl derivatives using (*N,O*-bis-[trimethylsilyl]acetamide) (Alltech Associates, Deerfield, IL). Silyl alcohols were quantitated by gas chromatography using an Econo-Cap SE-30 column (15 m \times 0.53 mm, 1.2 μm film thickness, Alltech). The column was maintained at 200 °C, the injection port at 250 °C, and the flame ionization detector at 300 °C. Helium carrier gas flow was 2.9 mL/min. For these conditions, the following elution times were observed: stearyl alcohol, 15.5 min; oleoyl alcohol, 14 min. H_2O_2 was determined spectrophotometrically at 650 nm by using horseradish peroxidase to oxidize the chromogenic dye 2,2'-azino-bis(3-ethylbenzthiazoline-6-sulfonic acid).

RESULTS

Although steady-state kinetic studies have been performed on the acyl-ACP desaturases (15), comparable studies of

² An MCD study of $4e^- \Delta 9D$ and the 18:0-ACP complex of $4e^- \Delta 9D$ revealed substantial changes in the diiron center upon complex formation that are consistent with both diiron centers containing five-coordinate sites in $4e^- \Delta 9D$ and both diiron centers containing one four-coordinate and one five-coordinate site in the complex (Y. Yang, J.A.B., B.G.F., and E. I. Solomon, manuscript submitted for publication).

Table 1: Rates of Reoxidation for Diiron Enzymes in the Presence of Various Substances

enzyme and substance	product	rate (min ⁻¹)
autoxidation ^a		
4e ⁻ reduced Δ 9D alone ^b		<0.002
+ pantotheinate, CoA		<0.002
+ methyl stearate		<0.002
+ apo-ACP, holo-ACP		<0.002
+ 18:0-ACP	18:0-ACP, 2 H ₂ O	0.027
RNR R2 Y122F ^c		~10
MMOH alone ^d		1.3
+ component B		3.0
catalytic oxidation ^e		
Δ 9D + 18:0-ACP ^f	18:1-ACP, 2 H ₂ O	20–30
RNR R2 Y122 ^c	Y122 ^g	42–60
MMOH + component B, CH ₄ ^d	CH ₃ OH	10.2

^a Rate for reappearance of the resting enzyme in the absence of a natural substrate. ^b Autoxidation monitored by the appearance of the μ -oxo charge transfer band at 350 nm after addition of the appropriate compound to chemically reduced Δ 9D exposed to 1 atm O₂. ^c Rate for conversion of the indicated RNR R2 isoforms to the diferric state (42, 43). ^d Rate for conversion of reduced MMOH to diferric MMOH in the absence or presence of component B and methane (24). ^e Rate of catalytic oxidation leading to the formation of the standard product: for Δ 9D, 18:1-ACP; for RNR R2, Y122^g; for MMOH, CH₃OH. ^f Turnover number calculated from Δ 9D multiple turnover at V_{\max} in the presence of NADPH, ferredoxin reductase, *Anabaena* [2Fe-2S] ferredoxin, and 18:0-ACP.

single turnover catalysis have not been reported. In part, this arises from the difficulty in preparing sufficient quantities of 18:0-ACP to support these material-intensive experiments (40, 41). However, our recent efforts to produce recombinant 18:0-ACP (23) have permitted the present examination of the single turnover reactivity of 4e⁻ reduced Δ 9D (4e⁻ Δ 9D).

Comparison of Rates for Autoxidation and Catalytic Turnover. The single turnover autoxidation of 4e⁻ Δ 9D was determined from reappearance of the 350 nm absorption band arising from oxo-bridged diferric centers (4). The conversion to the ferric state was slow (Table 1, $k' < 0.002$ min⁻¹) and was not affected by the addition of $\sim 10^2$ – 10^3 -fold molar excess of pantotheinate, coenzyme A, methyl stearate, apo-ACP, or holo-ACP (Table 1), revealing that neither substrate analogues nor the protein portion of the natural substrate was able to influence the reactivity of 4e⁻ Δ 9D. Likewise, the reoxidation rate was not changed in the pH range 6–9 and was not significantly increased at 40 °C. Upon completion of the autoxidation, >95% of the intensity of the original 340 nm absorption band was observed, suggesting near-stoichiometric recovery of diiron centers in the diferric state.

For diferrous RNR R2, Table 1 shows that the overall rate for single turnover oxidation of the natural isoform Y122 is 42–60 min⁻¹ (42), while in RNR R2 isoform Y122F, representing an enzyme form lacking the residue that is the single turnover oxidation substrate, the rate of the autoxidation is ~ 10 min⁻¹ (43). Thus, despite the large difference in the stability of tyrosine and phenylalanine to oxidation, and despite the difference in the ultimate products of the reaction, the reoxidation rates for RNR R2 isoforms Y122 and Y122F differ by only 6-fold (Table 1). In the case of chemically reduced MMOH, Table 1 shows that the rate of autoxidation in the absence of the component B effector protein and substrate is 1.3 min⁻¹ (44), while the multiple turnover catalytic rate is increased by only ~ 10 -fold to 10.2

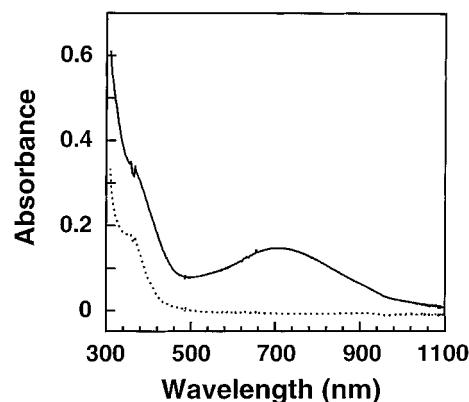


FIGURE 1: Optical spectra of 4e⁻ reduced stearyl-ACP Δ^9 desaturase exposed to 1 atm of O₂ gas in the presence (solid line) or absence (dotted line) of 18:0-ACP.

min⁻¹ upon addition of a suitable combination of the component B effector protein and methane (24, 44).

In the absence of substrate, 4e⁻ Δ 9D has an $\sim 10^4$ slower rate for single turnover autoxidation than RNR R2, RNR R2 Y122F, or MMOH (Table 1). Moreover, although the rate of reoxidation observed from 4e⁻ Δ 9D in the presence of 18:0-ACP was increased by ~ 10 -fold (Table 1, $k' \approx 0.027$ min⁻¹) as compared to 4e⁻ Δ 9D alone, this rate is still 10^3 -fold slower than that expected from the rate determined for catalytic desaturation (Table 1, $k' \approx 20$ – 30 min⁻¹). These behaviors are remarkably distinct from RNR R2 and MMOH and suggest unanticipated mechanisms for control of the Δ 9D catalytic cycle.

Peroxo-diferric Intermediate in Autoxidation. Resting Δ 9D has absorption bands at 320 and 340 nm that are typical for oxo \rightarrow Fe(III) charge transfer, whereas 4e⁻ Δ 9D has no optical absorption in the visible region (4). The dotted line in Figure 1 shows the optical spectrum of 4e⁻ Δ 9D exposed to O₂ gas (1 atm) in the absence of 18:0-ACP. No evidence for the formation of intermediates in the slow autoxidation was observed. Furthermore, no changes in absorption or autoxidation rate were observed if either apo-ACP (10-fold molar excess, lacking the functionally essential phosphopantetheine group) or holo-ACP (10-fold molar excess, lacking an acyl chain) were added to the reduced enzyme exposed to 1 atm O₂. However, when 18:0-ACP was added, a chromophoric complex with a broad absorption band at 700 nm and a shoulder at 380 nm was formed (solid line in Figure 1). The intensity of the 700 nm band was dependent on the concentration of 4e⁻ Δ 9D present and could be increased by titration of 18:0-ACP into the reaction cuvette (see below). Other experiments revealed that the same optical spectrum was formed regardless of whether 18:0-ACP was added before or after reduction or before or after addition of O₂.

Figure 2 shows RR spectra obtained by 647.1 nm excitation of Δ 9D samples prepared with ¹⁶O₂, ¹⁶O¹⁸O (50% isotopic enrichment), and ¹⁸O₂. When the O₂ adduct of the complex of 4e⁻ Δ 9D and 18:0-ACP was formed in ¹⁶O₂, two features were observed at low frequency (442 and 490 cm⁻¹) that have the expected energy for ν_s and ν_{as} of an Fe–O₂ bond, while another feature was observed at high frequency (898 cm⁻¹) that has the expected energy for ν -(O–O) of a metal-coordinated peroxide (26, 45, 46). Each of the three vibrational bands observed in the ¹⁶O₂ samples

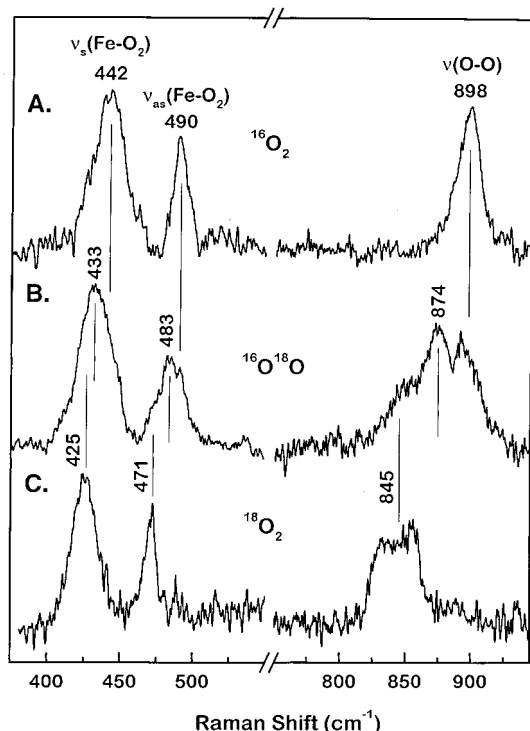


FIGURE 2: RR spectra of peroxodiferric $\Delta 9D$ generated with (A) $^{16}\text{O}_2$, (B) $^{16}\text{O}^{18}\text{O}$ (50 atom % ^{18}O), and (C) $^{18}\text{O}_2$ (98 atom % ^{18}O). Each spectrum was obtained within 10 min after thawing and was corrected for protein peaks by subtracting the 10 min spectrum obtained 50 min after thawing. The peroxo intermediate essentially disappeared after the first 10 min due to photodecomposition in the laser beam.

were symmetric, single peaks. In a sample prepared with $^{18}\text{O}_2$, the low-frequency vibrational bands were shifted by -17 and -19 cm^{-1} to 425 and 471 cm^{-1} , respectively, while the high-frequency band was shifted by -54 to 845 cm^{-1} . The magnitudes of the observed frequency shifts were consistent with the change in mass expected for substitution of isotopically labeled O_2 into the chromophore. The low-frequency peaks in the $^{18}\text{O}_2$ sample were also clearly symmetric, single peaks, whereas the $\nu(\text{O}-\text{O})$ band appeared to be a doublet. However, since $\nu(\text{O}-\text{O})$ for the $^{16}\text{O}_2$ -adduct was a single peak (898 cm^{-1}), the doublet centered at 845 cm^{-1} in the $^{18}\text{O}_2$ sample most likely arose from Fermi resonance with an unidentified mode not normally enhanced at this excitation wavelength. Fermi resonance has been previously observed in the RR spectra of other peroxodiferric model complexes (47) and in the $^{17}\text{O}_2$ complex of RNR R2 W48F/D84E, reported in the previous paper in this issue (26).

In the $\Delta 9D$ sample prepared with $^{16}\text{O}^{18}\text{O}$, containing a 1:2:1 mixture of $^{16}\text{O}_2$, $^{16}\text{O}^{18}\text{O}$, and $^{18}\text{O}_2$, the center of the low-frequency vibrational bands appeared at 433 and 483 cm^{-1} , respectively, while the center of the high-frequency band appeared at 874 cm^{-1} . For each vibrational band, the frequency was nearly halfway between the frequencies observed in the $^{16}\text{O}_2$ and $^{18}\text{O}_2$ samples, which was consistent with partial change in mass. Furthermore, the use of $^{16}\text{O}^{18}\text{O}$ resulted in the resolution of all three vibrational bands [$\nu_s(\text{Fe}-\text{O})$, $\nu_{as}(\text{Fe}-\text{O})$, $\nu(\text{O}-\text{O})$] into triplet vibrational peaks. Thus, for example, the $\nu(\text{O}-\text{O})$ peak in the mixed isotope sample had a central component at 874 cm^{-1} and side components at 845 and 898 cm^{-1} whose intensities cor-

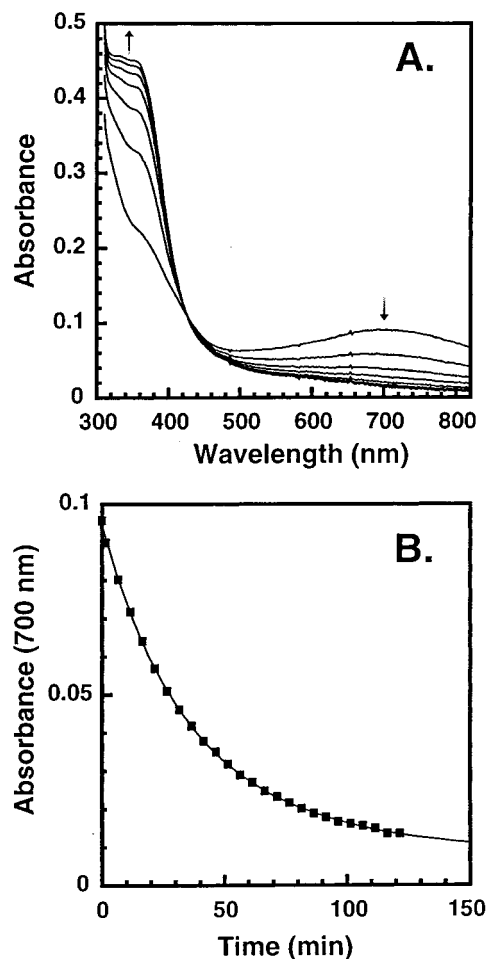


FIGURE 3: Decomposition of peroxodiferric $\Delta 9D$ formed from $\Delta 9D$ ($43\text{ }\mu\text{M}$ holoprotein) and 18:0-ACP ($140\text{ }\mu\text{M}$). (A) Optical spectra obtained at 40 min intervals. The decrease in absorbance at 700 nm and the increase in absorbance at 340 nm are as indicated. (B) The time course of the absorbance change at 700 nm . The solid line is a nonlinear least-squares fit assuming single-exponential decay of peroxodiferric $\Delta 9D$.

responded to the fractions of $^{16}\text{O}^{18}\text{O}$ (50%), $^{18}\text{O}_2$ (25%), and $^{16}\text{O}_2$ (25%) present in the gas, respectively. The presence of this triplet effectively ruled out asymmetric binding modes such as η^1 or μ -1,1, where two sets of triplets with vibrational frequencies characteristic of either ^{16}O - or ^{18}O -coordination to the metal center would be expected. Furthermore, the observed $\nu(\text{O}-\text{O})$ was considerably higher than that observed for μ - η^2 : η^2 peroxodicopper(II) complexes or oxyhemocyanin [$\sim 750\text{ cm}^{-1}$ (45, 48)]. Therefore, the vibrational spectra of Figure 2 are most consistent with a symmetric μ -1,2 peroxy structure in $\Delta 9D$,³ which also establishes the fundamental structural similarity of the peroxy level intermediates in the two closely related diiron enzymes, $\Delta 9D$ and RNR R2 W48F/D84E.

Decay of Peroxodiferric $\Delta 9D$. Figure 3A shows optical spectra obtained during the decay of peroxodiferric $\Delta 9D$, while Figure 3B shows the time course of the change in optical absorbance at 700 nm . During the decay, the loss of the 700 nm band corresponding to peroxodiferric $\Delta 9D$ and the reappearance of the oxo $\rightarrow \text{Fe(III)}$ charge-transfer

³ The *cis*- μ -1,2 and *gauche*- μ -1,2 configurations cannot be distinguished by vibrational data at present.

bands were associated with a reasonably tight isosbestic point at ~ 420 nm. After accounting for a nonzero A_∞ using Kedzy–Swinbourne analysis, plots of $\ln[(A(t) - A_\infty/A(0) - A_\infty)]$ versus time were linear for the complete decay process. Thus, the decrease in absorption at 700 nm was well fit with a single exponential, and nonlinear least-squares fitting of the experimental data gave $k_{\text{app}} = 0.027 \text{ min}^{-1}$. The stability of peroxodiferric $\Delta 9\text{D}$ was not dependent on the presence of O_2 as replacement of the headspace in the optical cuvette by repeated cycles of flushing with O_2 -free Ar caused no change in the decay rate. Furthermore, the rate was independent of $\Delta 9\text{D}$ concentration over a 4-fold range (20–80 μM), suggesting that intramolecular $2e^-$ transfer was associated with the loss of the 700 nm absorption and return of the enzyme to the resting state.

Gas chromatographic analysis revealed that 18:1-ACP was not produced by the decay of peroxodiferric $\Delta 9\text{D}$, while the stearyl chain was quantitatively recovered from total 18:0-ACP (>90%). For these catalytic assays, the detection limit for an oleoyl product relative to $\Delta 9\text{D}$ active sites present was less than 0.5%. Likewise, a spectrophotometric assay for H_2O_2 release based on horseradish peroxidase-catalyzed dye oxidation revealed that no H_2O_2 was released during decay of peroxodiferric $\Delta 9\text{D}$. In similar assays, exogenously added H_2O_2 was quantitatively recovered at less than 5% of the molar amount of $\Delta 9\text{D}$ active sites present. After the reoxidation was complete, an optical spectrum with nearly identical intensity to that of the original resting $\Delta 9\text{D}$ preparation was obtained. This reoxidation cycle was repeated for three cycles with no loss in the ability to reform peroxodiferric $\Delta 9\text{D}$ and with no decrease in the intensity of the resting $\text{oxo} \rightarrow \text{Fe(III)}$ charge-transfer band. Furthermore, redox-cycled $\Delta 9\text{D}$ had multiple turnover activity ($k' \approx 20\text{--}30 \text{ min}^{-1}$ for 18:1-ACP production, Table 1) indistinguishable from the activity of aliquots of the same $\Delta 9\text{D}$ preparation not subjected to the single turnover reoxidation cycle.

Stoichiometry of 18:0-ACP in the Peroxo Intermediate. Figure 4A shows a titration of 18:0-ACP into a solution of $4e^- \Delta 9\text{D}$ equilibrated with 1 atm O_2 . Due to the stability of $4e^- \Delta 9\text{D}$ in the presence of O_2 , and due to the relative stability of peroxodiferric $\Delta 9\text{D}$ once formed, this titration was performed at 20°C over ~ 6 min. Addition of 18:0-ACP aliquots caused the progressive appearance of the 700 and 380 nm absorption bands assigned to peroxodiferric $\Delta 9\text{D}$. The formation of this product was maximized after the addition of ~ 2 mol of 18:0-ACP/mol of $\Delta 9\text{D}$ (Figure 4B), suggesting that one 18:0-ACP was bound per subunit of $4e^- \Delta 9\text{D}$. Other experiments performed by rapid addition of up to a 10-fold excess of 18:0-ACP revealed that no additional intensity of the 700 nm band could be formed. On the basis of the linear increase in absorption observed during the titration, a molar absorptivity of $1200 \pm 50 \text{ M}^{-1} \text{ cm}^{-1}$ was determined relative to the concentration of 18:0-ACP added. While this value is similar to those determined for other peroxodiferric intermediates, RNR R2 D84E [$\sim 1500 \text{ M}^{-1} \text{ cm}^{-1}$ (25)], MMOH [$\sim 1500 \text{ M}^{-1} \text{ cm}^{-1}$ (27)], and crystallographically characterized synthetic model complexes [$\sim 1500\text{--}3540 \text{ M}^{-1} \text{ cm}^{-1}$ (46, 49)], it does not conclusively reveal whether one or two peroxo moieties are present.

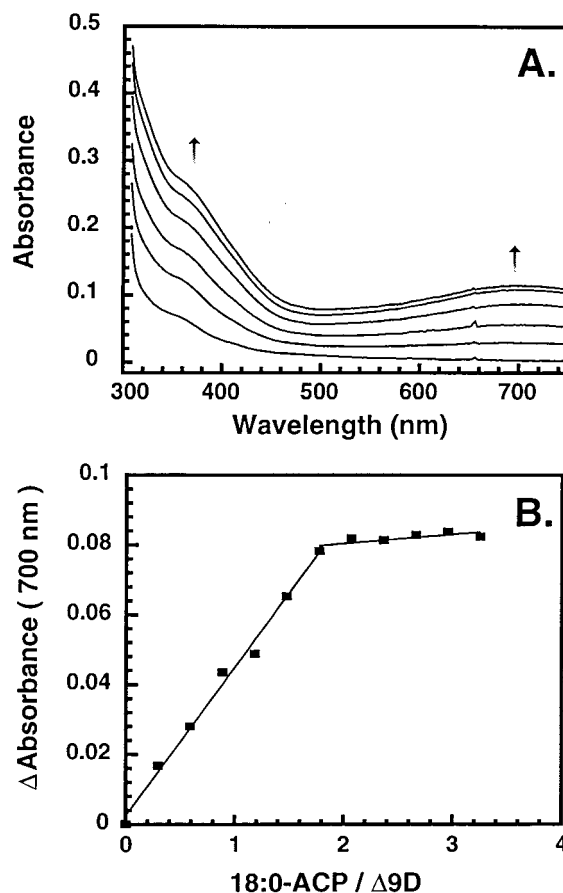


FIGURE 4: Formation of peroxodiferric $\Delta 9\text{D}$ during titration with 18:0-ACP. (A) optical spectra obtained from successive addition of 5 μL aliquots of 1.23 mM 18:0-ACP to $4e^- \Delta 9\text{D}$ (40 μM holoprotein). (B) The increase in absorbance at 700 nm plotted as a function of moles of 18:0-ACP added per mole of $\Delta 9\text{D}$ dimer.

DISCUSSION

Many new insights into biological O_2 activation have been provided by study of RNR R2, MMOH, and synthetic active-site mimics (50). Since RNR R2, MMOH, and $\Delta 9\text{D}$ have similar protein folds, ligation environments, and hydrophobic active sites, it is generally assumed that many aspects of their catalytic cycles will be closely related (2, 50, 51). This assumption has been validated by the similar Mössbauer properties of the peroxodiferric intermediates of RNR R2 D84E (25), MMOH compound P (H_{peroxo}) (27), and ferritin (13) and the similar Mössbauer properties of intermediate X of RNR R2 (52) and $1e^-$ reduced MMOH compound Q (31). In combination with the previous study of RNR R2 W48F/D84E (26), this overall structural similarity has been refined to include symmetric μ -1,2 peroxodiferric intermediates in two of the three members of the class II diiron enzymes by use of vibrational spectroscopy.⁴ However, these studies also reveal an unanticipated reactivity of $4e^- \Delta 9\text{D}$.

Structural Evidence for Peroxodiferric $\Delta 9\text{D}$. The addition of O_2 to the complex of 18:0-ACP and $4e^- \Delta 9\text{D}$ results in the formation of a quasi-stable intermediate with a broad

⁴ A resonance Raman feature initially ascribed to the putative peroxodiferric intermediate in MMOH (53) was later reported (54) to be an artifact arising, in part, from difficulties in the subtraction of the spectrum of 2-methylbutane cryosolvent used to prepare the freeze-quenched samples.

absorption maximum at 700 nm. When compared to oxyhemerythrin [$\nu(\text{O}-\text{O}) = 844 \text{ cm}^{-1}$ for η^1 coordination (55)] or oxyhemocyanin [$\nu(\text{O}-\text{O}) = \sim 750 \text{ cm}^{-1}$ for $\mu-\eta^2$: η^2 coordination (45, 48)], the relatively high $\nu(\text{O}-\text{O})$ value of 898 cm^{-1} observed for peroxodiferrous $\Delta 9\text{D}$ is indicative of a bridging peroxide structure (56), as is the observation of two $\nu(\text{Fe}-\text{O}_2)$ modes at 442 and 490 cm^{-1} . Furthermore, the accompanying paper on the peroxodiferrous intermediate of RNR R2 W48F/D84E reveals a remarkably similar RR spectrum with $\nu(\text{O}-\text{O})$ at 870 cm^{-1} , $\nu_s(\text{Fe}-\text{O}_2)$ at 458 cm^{-1} , and $\nu_{as}(\text{Fe}-\text{O}_2)$ at 499 cm^{-1} (26). For both enzymes, the observation that the $^{16}\text{O}/^{18}\text{O}$ mixed isotope produces single vibrational peaks split in a 1:2:1 pattern arising from the isotopic composition of the gas strongly supports our proposal that peroxide is coordinated to the class II diiron enzymes in a symmetric fashion, such as a $\mu-1,2$ structure. This proposal is also consistent with the ability of azide to coordinate to resting $\Delta 9\text{D}$ to form a $\mu-1,3$ azidodiferrous structure (18).

Functional Properties of Peroxodiferrous $\Delta 9\text{D}$. Despite the structural similarities of the peroxodiferrous intermediates detected in $\Delta 9\text{D}$ and RNR R2, the functional properties of these two species differ significantly. For example, peroxodiferrous RNR R2 D84E accumulates in air-saturated buffer and then decays with $t_{1/2}$ of $\sim 6 \text{ s}$ to give the expected oxidation product Y122* (25), while peroxodiferrous RNR R2 W48F/D84E is more stable by ~ 4 -fold and decays to generate a small amount of Y122* (26). In contrast, peroxodiferrous $\Delta 9\text{D}$ accumulates to high levels, but only when $4e^-$ $\Delta 9\text{D}$, 18:0-ACP, and 1 atm O_2 are present: omission of 18:0-ACP or the substitution of air-saturated buffer does not support formation of this peroxy intermediate. The titration data of Figure 4 indicate that 18:0-ACP binds tightly to $4e^-$ $\Delta 9\text{D}$, that 18:0-ACP binding linearly correlates with formation of the peroxy intermediate, and that each subunit of the $\Delta 9\text{D}$ dimer can bind a substrate molecule. Peroxodiferrous $\Delta 9\text{D}$ then decays by a first-order process with $t_{1/2}$ of $\sim 26 \text{ min}$ without detectable desaturation of 18:0-ACP, but likewise without detectable release of H_2O_2 . This decay yields resting $\Delta 9\text{D}$ in essentially quantitative recovery with respect to diferric centers as judged by optical spectroscopy and with respect to catalytic activity as judged by subsequent 18:1-ACP production using the enzymatic electron-transfer chain.

It is generally concluded that flexibility of the diiron center ligation environment has a major impact on catalysis (10, 18–23). In this regard, the stability of the peroxodiferrous intermediate described here may arise from only relatively minor alterations in conformation associated with inappropriate redox state, ligation pattern, or Fe–Fe distance. The elucidation of how, or if, these alterations permit stable intermediates to be converted into more reactive states will likely provide further insight into O_2 activation by diiron centers.

Oxidase Chemistry of $4e^-$ $\Delta 9\text{D}$. Chemically reduced $\Delta 9\text{D}$ (containing $4e^-$ in two diferrous sites) has a 10^4 -fold lower reactivity with O_2 than either RNR R2 or MMOH, suggesting that $\Delta 9\text{D}$ has a unique mechanism to prevent autooxidation. Moreover, $4e^-$ $\Delta 9\text{D}$ has $\sim 10^3$ -fold lower reactivity with O_2 in the presence of 18:0-ACP than enzymatically reduced $\Delta 9\text{D}$ (possibly containing one diferrous cluster and one diferric cluster within the same dimeric protein, since a mixed

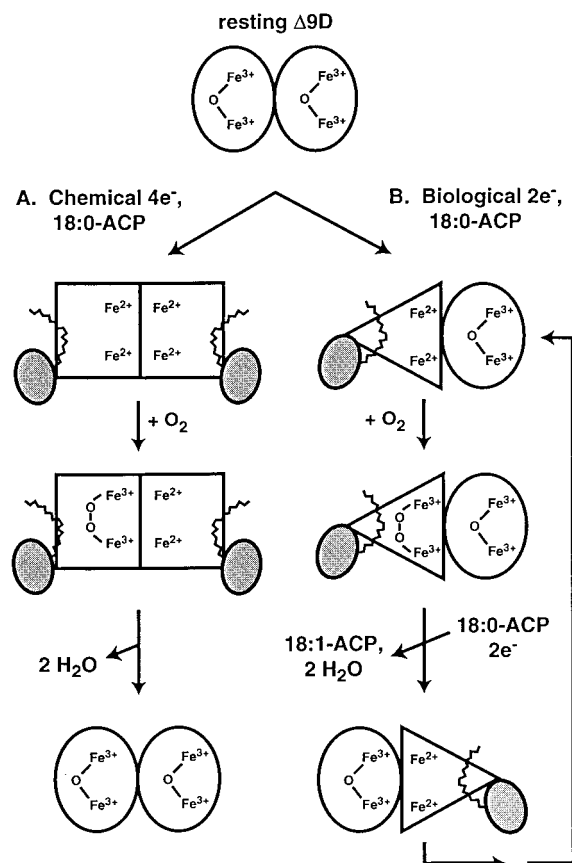


FIGURE 5: Oxidase and desaturase pathways of $\Delta 9\text{D}$ differentiated by the overall redox state. (A) oxidase pathway initiated by $4e^-$ chemical reduction. The origin of the oxo bridge (O_2 or solvent) in the resting enzyme produced after the oxidase reaction is not specified. (B) multiple turnover desaturation pathway initiated by $2e^-$ enzymatic reduction of one diiron center of the dimeric resting enzyme. The use of alternating active sites is hypothesized to coordinate the transfer of reducing equivalents, substrate binding (18:0-ACP and O_2), and product release (18:1-ACP). The kinetic order of addition of reducing equivalents and substrates is not known.

valence state has not been observed). These observations suggest that the overall redox state of $\Delta 9\text{D}$ may be an important determinant of the outcome of catalysis. Figure 5 shows proposed models accounting for the differences in catalytic properties of chemically reduced and enzymatically reduced $\Delta 9\text{D}$. We propose that $4e^-$ chemical reduction of $\Delta 9\text{D}$ produces a conformation that enforces oxidase chemistry (Figure 5A). The oxidase reaction occurs via the formation of an enzyme-bound peroxodiferrous species and ultimately leads to recovery of resting $\Delta 9\text{D}$. In these experiments, diferrous centers are the only likely source of reducing equivalents.⁵ Thus, the *in vitro* oxidase reaction may involve an intramolecular electron transfer between diferrous and peroxodiferrous centers in different subunits of the enzyme (60). In principle, the $2e^-$ reduction of either a peroxodiferrous center or a high valent intermediate produced by homolysis

⁵ The 700 nm band was quickly lost upon addition of either sodium dithionite or reduced [2Fe-2S] ferredoxin, and resulted in the reappearance of the oxo \rightarrow Fe(III) charge-transfer band of resting $\Delta 9\text{D}$ (J.A.B. and B.G.F., unpublished results). Therefore, an alternative electron-transfer pathway related to that proposed for residue W48 of RNR R2 (9, 57–59) may also be present in $\Delta 9\text{D}$, which has a structurally analogous tryptophan, W62.

of the O—O bond in the peroxidiferic center are possible mechanisms to restore resting $\Delta 9D$. These options cannot be distinguished at present.

Mechanistic Implications of Oxidase versus Desaturase Reactions. The observation that the oxidase reaction occurs on $4e^- \Delta 9D$, but that acyl-ACP desaturation does not, implies that multiple turnover catalysis may normally proceed via a “half-site” reactivity (Figure 5B), which has also been described for RNR R2 (61). Presently, the kinetic order for addition of reducing equivalents, O_2 and 18:0-ACP, and for the release of 18:1-ACP is not known for $\Delta 9D$. Likewise, a peroxidiferic intermediate has not been established for desaturase catalysis, but remains an attractive possibility. One possible advantage of the redox-cycle gated process summarized in Figure 5B would be that 18:0-ACP binding to the unreacted subunit could be coordinated with 18:1-ACP release from the other subunit. Furthermore, these conformational changes may be associated with activation of various O_2 adducts during the catalytic cycle. We postulate that a complete loading of electrons and 18:0-ACP onto the $\Delta 9D$ dimer by chemical reduction may disrupt this cycle, thus allowing for the observed oxidase chemistry.

Biological Implications of the Oxidase Reaction. Although the autooxidation of $4e^- \Delta 9D$ in the presence of 18:0-ACP is $\sim 10^3$ -fold slower than the desaturation of 18:0-ACP by biologically reduced $\Delta 9D$, this autooxidation chemistry provides a route to restore resting $\Delta 9D$ without yielding deleterious products such as H_2O_2 . Since $\Delta 9D$ is found in long-lived organelles such as the chloroplast and plastid, which also contain substantial levels of reducing equivalents, evolutionary selection may have provided $\Delta 9D$ a control of diiron center reactivity that has not been necessary in RNR R2 and MMOH, which are both from more rapidly growing bacteria.

$\Delta 9D$ appears to be able to perform oxidase chemistry using $4e^-$ entirely from diiron centers without oxidation of active-site amino acids. In contrast, RNR R2 performs an oxidase reaction using $2e^-$ from a diferrous center, $1e^-$ from Y122, and $1e^-$ from an exogenous source. MMOH from *Methylococcus capsulatus* (Bath) has also been proposed to convert between hydroxylase and oxidase activities as a function of component B (62). This process is of obvious relevance to the oxidase reactivity reported here, however, no enzyme intermediates were detected during these previous studies.

The $4e^-$ reduction of O_2 to $2H_2O$ without acyl desaturation and without H_2O_2 release described here represents a new mode of catalysis for $\Delta 9D$ that is further supported by the identification of a key intermediate in the reaction. This demonstration of oxidase catalysis is consistent with the rapid increase in the number of functions reported for diiron enzymes in the past few years (51, 63). In this regard, the “alternative oxidase” from fungi, plants, and trypanosomes has recently been proposed to be a diiron enzyme having repeats of the sequence motif GluXXHis (6, 64–66). Although further studies, including confirmation of the metal binding motif and the verification of a spin-coupled iron environment, are required, the catalytic and spectroscopic properties reported here for single turnover preparations of $4e^- \Delta 9D$ may indeed provide a mechanistic framework for further investigation of the “alternative oxidase” reaction by the putative membrane diiron centers.

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REFERENCES

- Feig, A., and Lippard, S. J. (1994) *Chem. Rev.* 94, 759–805.
- Wallar, B. J., and Lipscomb, J. D. (1996) *Chem. Rev.* 96, 2625–2657.
- Fox, B. G. (1997) in *Comprehensive Biological Catalysis* (Sinnott, M., Ed.) pp 261–348, Academic Press, London.
- Fox, B. G., Shanklin, J., Ai, J., Loehr, T. M., and Sanders-Loehr, J. (1994) *Biochemistry* 33, 12776–12786.
- Shanklin, J., Whittle, E., and Fox, B. G. (1994) *Biochemistry* 33, 12787–12794.
- Moore, A. L., Umbach, A. L., and Siedow, J. N. (1995) *J. Bioeng. Biomembr.* 27, 367–377.
- Harrison, P., and Arosio, P. (1996) *Biochim. Biophys. Acta* 1275, 161–203.
- deMaré, F., Kurtz, D. M., and Nordlund, P. (1996) *Nat. Struct. Biol.* 3, 539–546.
- Nordlund, P., and Eklund, H. (1993) *J. Mol. Biol.* 232, 123–164.
- Rosenzweig, A. C., Nordlund, P., Takahara, P. M., Frederick, C. A., and Lippard, S. J. (1995) *Chem. Biol.* 2, 409–418.
- Elango, N., Radhakrishnan, R., Froland, W. A., Wallar, B. J., Earhart, C. A., Lipscomb, J. D., and Ohlendorf, D. H. (1997) *Protein Sci.* 6, 556–568.
- Lindqvist, Y., Huang, W., Schneider, G., and Shanklin, J. (1996) *EMBO J.* 15, 4081–4092.
- Pereira, A. S., Small, W., Krebs, C., Tavares, P., Edmondson, D. E., Theil, E. C., and Huynh, B. H. (1998) *Biochemistry* 37, 9871–9876.
- Bonomi, F., Kurtz, D. M., Jr., and Cui, X. (1996) *J. Biol. Inorg. Chem.* 1, 67–72.
- Shanklin, J., and Cahoon, E. B. (1998) *Annu. Rev. Plant Physiol. Plant Mol. Biol.* 49, 611–641.
- Fox, B. G., Shanklin, J., Somerville, C., and Münck, E. (1993) *Proc. Natl. Acad. Sci. U.S.A.* 90, 2486–2490.
- Shu, L., Broadwater, J. A., Achim, C., Fox, B. G., Münck, E., and Que, L., Jr. (1998) *J. Biol. Inorg. Chem.* 3, 392–400.
- Ai, J., Broadwater, J. A., Loehr, T. M., Sanders-Loehr, J., and Fox, B. G. (1997) *J. Biol. Inorg. Chem.* 2, 37–45.
- Logan, D. T., Su, X.-D., Åberg, A., Regnström, K., Hajdu, J., Eklund, H., and Nordlund, P. (1996) *Structure* 4, 1053–1064.
- Rardin, R. L., Tolman, W. B., and Lippard, S. J. (1991) *New J. Chem.* 15, 417–430.
- Shu, L., Nesheim, J. C., Kauffmann, K., Münck, E., Lipscomb, J. D., and Que, L., Jr. (1997) *Science* 275, 515–518.
- Shu, L., Liu, Y., Lipscomb, J. D., and Que, L., Jr. (1996) *J. Biol. Inorg. Chem.* 1, 297–304.
- Broadwater, J. A., Haas, J. A., and Fox, B. G. (1998) *FETT/Lipid* 100, 103–113.
- Lee, S.-K., Nesheim, J. C., and Lipscomb, J. D. (1993) *J. Biol. Chem.* 268, 21569–21577.
- Bollinger, J. M., Jr., Krebs, C., Vicol, A., Chen, S., Ley, B. A., Edmondson, D. E., and Huynh, B. H. (1998) *J. Am. Chem. Soc.* 120, 1094–1095.
- Moënné-Loccoz, P., Baldwin, J., Ley, B. A., Loehr, T., and Bollinger, J. M., Jr. (1998) *Biochemistry* 37, 14659–14663.
- Liu, K. E., Valentine, A. M., Wang, D., Huynh, B. H., Edmondson, D. E., Salifoglou, A., and Lippard, S. J. (1995) *J. Am. Chem. Soc.* 117, 10174–10185.
- Lee, S.-K., Fox, B. G., Froland, W. F., Lipscomb, J. D., and Münck, E. (1993) *J. Am. Chem. Soc.* 115, 6450–6451.
- Sturgeon, B. E., Burdi, D., Chen, S., Huynh, B.-H., Edmondson, D. E., Stubbe, J., and Hoffman, B. M. (1996) *J. Am. Chem. Soc.* 118, 7551–7557.
- Willems, J.-P., Lee, H.-I., Burdi, D., Doan, P. E., Stubbe, J., and Hoffman, B. M. (1997) *J. Am. Chem. Soc.* 119, 9816–9824.

31. Valentine, A. M., Tavares, P., Pereria, A. S., Davydov, R., Krebs, C., Hoffman, B. M., Edmondson, D. E., Huynh, B. H., and Lippard, S. J. (1998) *J. Am. Chem. Soc.* **120**, 2190–2191.
32. Riggs-Gelasco, P. J., Shu, L., Chen, S., Burdi, D., Huynh, B. H., Que, L., Jr., and Stubbe, J. (1998) *J. Am. Chem. Soc.* **120**, 849–860.
33. Hoffman, B. J., Broadwater, J. A., Johnson, P., Harper, J., Fox, B. G., and Kenealy, W. R. (1995) *Protein Expression Purif.* **6**, 646–654.
34. Bradford, M. M. (1976) *Anal. Biochem.* **72**, 248–252.
35. Fischer, D. S., and Price, D. C. (1964) *Clin. Chem.* **10**, 21–30.
36. Lambalot, R. H., and Walsh, C. T. (1995) *J. Biol. Chem.* **270**, 24658–24661.
37. Rock, C. O., and Garwin, J. L. (1979) *J. Biol. Chem.* **254**, 7123–7128.
38. Espenson, J. H. (1981) in *Chemical Kinetics and Reaction Mechanisms*, McGraw-Hill, New York, p 218.
39. Barron, E. J., and Mooney, L. A. (1968) *Anal. Chem.* **40**, 1742–1744.
40. Beremand, P. D., Hannapel, D. J., Guerra, D. J., Kuhn, D. N., and Ohlrogge, J. B. (1987) *Arch. Biochem. Biophys.* **256**, 90–100.
41. Hill, R. B., MacKenzie, K. R., Flanagan, J. M., Cronan, J. E., Jr., and Prestegard, J. H. (1995) *Protein Expression Purif.* **6**, 394–400.
42. Tong, W. H., Chen, S., Lloyd, S. G., Edmondson, D. E., Huynh, B.-H., and Stubbe, J. (1996) *J. Am. Chem. Soc.* **118**, 2107–2108.
43. Sahlin, M., Lassmann, G., Pötsch, S., Sjöberg, B.-M., and Gräslund, A. (1995) *J. Biol. Chem.* **270**, 12361–12372.
44. Liu, Y., Nesheim, J. C., Lee, S.-K., and Lipscomb, J. D. (1995) *J. Biol. Chem.* **270**, 24662–24665.
45. Kitajima, N., Fujisawa, K., Fujimoto, C., Moro-oka, Y., Hashimoto, S., Kitagawa, T., Toriumi, K., Tatsumi, K., and Nakamura, A. (1992) *J. Am. Chem. Soc.* **114**, 1277–1291.
46. Kim, K., and Lippard, S. J. (1996) *J. Am. Chem. Soc.* **118**, 4914–4915.
47. Dong, Y., Ménage, S., Brennan, B. A., Elgren, T. E., Jang, H. G., Pearce, L. L., and Que, L., Jr. (1993) *J. Am. Chem. Soc.* **115**, 1851–1859.
48. Freedman, T. B., Sanders-Loehr, J., and Loehr, T. (1978) *J. Am. Chem. Soc.* **98**, 2809.
49. Kitajima, N., Tamura, N., Fukui, H., Moro-oka, Y., Mitzutani, Y., Kitagawa, T., Mathur, R., Heerwegh, K., Reed, C. A., Randall, C. R., Que, L., Jr., and Tatsumi, K. (1994) *J. Am. Chem. Soc.* **116**, 9071–9085.
50. Que, L., Jr., and Dong, Y. (1996) *Acc. Chem. Res.* **29**, 190–196.
51. Broadwater, J. A., Haas, J. A., Achim, C., Münck, E., Ai, J., Sanders-Loehr, J., Loehr, T. M., and Fox, B. G. Catalytic studies of acyl-ACP desaturation: The unique contributions of protein and lipid to enzymatic catalysis. Presented at the *26th Steenbock Symposium: Kinetics of Enzymatic Catalysis* (Frey, P. A., and Northrup, D., Eds.) May 28–31, 1998, Madison, WI, IOS Press, Amsterdam.
52. Ravi, N., Bollinger, J. M., Jr., Huynh, B. H., Edmondson, D. E., and Stubbe, J. (1994) *J. Am. Chem. Soc.* **116**, 8007–8014.
53. Liu, K. E., Valentine, A. M., Qui, D., Edmondson, D. E., Appelman, E. H., Spiro, T. G., and Lippard, S. J. (1995) *J. Am. Chem. Soc.* **117**, 4997–4998.
54. Liu, K. E., Valentine, A. M., Qui, D., Edmondson, D. E., Appelman, E. H., Spiro, T. G., and Lippard, S. J. (1997) *J. Am. Chem. Soc.* **119**, 11134–11136.
55. Freier, S. M., Duff, L. I., Shriver, D. F., and Klotz, I. M. (1980) *Arch. Biochem. Biophys.* **205**, 449–463.
56. Ross, P. K., and Solomon, E. I. (1991) *J. Am. Chem. Soc.* **113**, 3246–3259.
57. Rova, U., Goodtzova, K., Ingemarson, R., Behravan, G., Gräslund, A., and Thelander, L. (1995) *Biochemistry* **34**, 4267–4275.
58. Ekberg, M., Sahlin, M., Eriksson, M., and Sjöberg, B.-M. (1996) *J. Biol. Chem.* **271**, 20655–20659.
59. Parkin, S. E., Chen, S., Ley, B. A., Mangravite, L., Edmondson, D. E., Huynh, B. H., and Bollinger, J. M., Jr. (1998) *Biochemistry* **37**, 1124–1130.
60. Elgren, T. E., Lynch, J. B., Juarez-Garcia, C., Münck, E., Sjöberg, B.-M., and Que, L., Jr. (1991) *J. Biol. Chem.* **266**, 19265–19268.
61. Sjöberg, B.-M., Karlsson, M., and Jörnvall, H. (1987) *J. Biol. Chem.* **262**, 9736–9743.
62. Green, J., and Dalton, H. (1985) *J. Biol. Chem.* **260**, 15795–15801.
63. Shanklin, J., Achim, C., Schmidt, H., Fox, B. G., and Münck, E. (1997) *Proc. Natl. Acad. Sci. U.S.A.* **94**, 2981–2986.
64. Siedow, J. N., Umbach, A. L., and Moore, A. L. (1995) *FEBS Lett.* **362**, 10–14.
65. Chaudhuri, M., and Hill, G. C. (1996) *Mol. Biochem. Parasitol.* **83**, 125–129.
66. Chaudhuri, M., Ajayi, W., and Hill, G. C. (1998) *Mol. Biochem. Parasitol.* **95**, 53–68.

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