

Unusual Peroxide-Dependent, Heme-Transforming Reaction Catalyzed by HemQ

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Supporting Information

ABSTRACT: A recently proposed pathway for heme *b* biosynthesis, common to diverse bacteria, has the conversion of two of the four propionates on coproheme III to vinyl groups as its final step. This reaction is catalyzed in a cofactor-independent, H₂O₂-dependent manner by the enzyme HemQ. Using the HemQ from Staphylococcus aureus (SaHemQ), the initial decarboxylation step was observed to rapidly and obligately yield the threepropionate harderoheme isomer III as the intermediate, while the slower second decarboxylation appeared to control the overall rate. Both synthetic

harderoheme isomers III and IV reacted when bound to HemQ, the former more slowly than the latter. While H₂O₂ is the assumed biological oxidant, either H₂O₂ or peracetic acid yielded the same intermediates and products, though amounts significantly greater than the expected 2 equiv were required in both cases and peracetic acid reacted faster. The ability of peracetic acid to substitute for H₂O₂ suggests that, despite the lack of catalytic residues conventionally present in heme peroxidase active sites, reaction pathways involving high-valent iron intermediates cannot be ruled out.

eme, the complex of protoporphyrin IX with iron, is fundamentally important for life. A diverse set of metalchelating cofactors, including F430 (Ni), vitamin B12 (Co), siroheme (Fe and S), and chlorophyll (Mg), additionally share a common tetrapyrrole scaffold and hence biosynthetic and evolutionary roots with heme. In humans and other eukaryotes, the biosynthesis of chemically unfunctionalized heme b occurs via eight well-known enzymatic steps. ^{1,2} A little more than a decade ago, however, a manual survey of the available prokaryotic genomes indicated that genes encoding many of these steps were absent from diverse species, suggesting the evolutionary history of the pathway may be complex.

Sophisticated bioinformatics methods and a substantially expanded pool of genomic data in conjunction with experimental approaches have since allowed many holes in the pathways to heme and related tetrapyrrole cofactors to be filled (Scheme 1). A complete route to heme biosynthesis in Archaea and sulfatereducing bacteria has been identified.⁴ Recently, a new ending for the pathway in Gram-positive bacteria from the Actinobacteria and Firmicutes phyla, a large group of organisms characterized by their lack of an outer membrane, was proposed. This and the canonical pathway effect the same three chemical transformations on the metabolic intermediate coproporphyrinogen III. However, the reactions in the two pathways occur in a different order, yield distinct intermediates, and invoke distinct enzymes for the oxidative decarboxylation step.⁵

In the Gram-positive bacteria, this step occurs at the terminus of the pathway, after a step analogous to the well-known ferrochelatase reaction. 6 It is catalyzed by HemQ, an enzyme that

uses coproheme III as both a substrate and a cofactor in the H₂O₂-dependent oxidative decarboxylation of the propionate side chains at β -pyrrole positions 2 and 4 (Scheme 2). Cofactorindependent, O2-dependent enzymes have been described, including coproporphyrinogen III oxidase (HemF).^{7,8} However, this is to the best of our knowledge the first example of a cofactorindependent transformation involving H2O2. In contrast with coproporphyrinogen III, the metallosubstrate used by HemQ offers the possibility of mechanistically distinct, iron-mediated activation of H₂O₂ for the oxidative decarboxylation step. Interestingly, the conversion of coproheme to heme b is also catalyzed anaerobically by AhbD, a radical S-adenosylmethionine- and iron-sulfur cluster-dependent enzyme at the terminus of the biosynthetic pathway in Archaea and sulfate-reducing bacteria (Scheme 1).^{4,9} Homologues are also found in some Gram-positive and possibly other bacteria along with hemQ genes, affording a possible anaerobic alternative for this step.

HemQs are part of the chlorite dismutase family (Clds, Pfam 06778). This family is best known for the minority of its members that use heme b to catalytically convert ClO_2^- to $Cl^$ and O2 at the terminus of perchlorate respiratory pathways. 12 The reaction of HemQs, by contrast, has not been well-studied. It has been previously shown that HemQ is not highly active toward H₂O₂ in its heme b-bound form but instead fairly readily undergoes heme degradation. Further, in its ferric coproheme

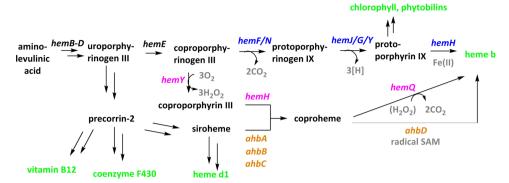
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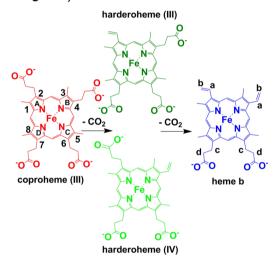
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Scheme 1. Biosynthetic Pathways Leading to Heme and Other Tetrapyrroles^a



"The genes encoding catalysts for the canonical steps common to eukaryotes and many Gram-negative bacteria are colored blue. The pathway used by Archaea and sulfur-reducing bacteria are colored orange. The recently proposed terminus of the pathway, found in Gram-positive members of Actinobacteria and Firmicutes, is colored purple. Note that HemF and HemN can catalyze the indicated step in different organisms under aerobic and anaerobic conditions, respectively. Similarly, HemJ, -G, or -Y catalyzes the removal of two H atoms ([H]).

Scheme 2. Substrate, Possible Intermediates, and Product of the HemQ-Catalyzed Reaction^a



^aHeme rings are labeled A–D and substituted pyrrole carbons 1–8. Vinyl and propionate carbons are labeled a–d.

III-bound state, the HemQ from Mycobacterium tuberculosis was shown to react with H_2O_2 rather than various O_2 /reductant combinations, fully converting coproheme III to heme b.⁵

Here, using the enzyme from *Staphylococcus aureus* (*Sa*-HemQ), we sought to identify the likely tetrapyrrole intermediates on the reaction pathway to narrow the possible reaction mechanisms. We aimed to determine whether the reactions with the propionates occur in an ordered fashion and with what time course, and whether a specific three-propionate harderoheme isomer could be identified as a reaction intermediate. Finally, to set limits on possible iron intermediates, we examined whether the same intermediates and products would be generated with organic peroxides and peracids and to establish the stoichiometry of the reaction in each case.

■ EXPERIMENTAL METHODS

Reagents and Stocks. Ferric coproporphyrin III chloride (coproheme III, Frontier Scientific) and hemin chloride (Sigma) were obtained in 10 mg ampules and used to generate 5–10 mM stock solutions in dimethyl sulfoxide (DMSO). Harderohemes III and IV prepared in house were rigorously dried in tared glass

ampules and weighed on an analytical balance. Stocks (5 mM) were prepared in DMSO in the ampules on the basis of the measured masses.

Hydrogen peroxide (H_2O_2), tert-butyl peroxide, m-chloroperbenzoic acid (mCPBA), and peracetic acid stock solutions were prepared at concentrations of 100 μ M in 50 mM potassium phosphate (KP_i) buffer (pH 7.4). H_2O_2 concentrations were ascertained by titration with freshly prepared 0.02 M KMnO₄ under acidic conditions (concentrated H_2SO_4). Titrations were monitored for loss of the characteristic pink color of permanganate as it is reduced to the nearly colorless Mn²⁺ ion. Peracetic acid stocks were analyzed with MQuant Peracetic acid test strips (EMD Millipore).

Biochemicals (lysozyme, catalase, DNase, etc.) were purchased from New England Biolabs. Reagents and media for bacterial growth, protein purification, and analysis were obtained from Fisher or GE Healthcare and used without further purification.

Synthesis and Purification of Harderoheme Isomers III and IV. Harderoporphyrin trimethyl esters were prepared as previously described^{7,8} and then metalated in an acetic acid/2% pyridine solution under argon. The resulting harderohemes were saponified using NaOH and analyzed by high-pressure liquid chromatography (HPLC) coupled to electrospray time-of-flight mass spectrometry (ESI-MS). See the Supporting Information and Scheme 1S for synthetic and related analytical details.

Expression and Purification of SaHemQ Bound to Substrate and Potential Intermediates. SaHemQ protein expression and purification were conducted as previously described⁹ (see the Supporting Information). The protein was incubated at 4 °C for 12-24 h with ferric coproheme III or harderoheme (isomer III or IV) in an approximate 1:1 (protein monomer:heme) ratio, based on the protein concentration estimated by the Bradford assay and SDS-PAGE. To remove unbound substrate and further purify the protein, coprohemebound SaHemQ was concentrated to 5 mL and loaded onto an S-200 Sephacryl gel filtration column [run at 0.4 mL/min in 50 mM KP, buffer (pH 7.4)]. Fractions were collected using an AKTA Purification System. Fractions with an $R_z \ge 0.5$ were pooled (R_z = absorbance_{Soret}/absorbance₂₈₀), concentrated, flash-frozen, and stored at -80 °C. A PD-10 desalting column (GE Healthcare) was used to remove unbound harderoheme from the protein before concentration and storage.

Extinction coefficients for the coproheme III- and heme bbound SaHemQ at the Soret absorbance band maxima ($\varepsilon_{\text{Soret}}$) were determined by the pyridine hemochrome assay. 13 Briefly, 200 μ L of heme-containing solution was mixed with 800 μ L of 50 mM NaOH containing 20% pyridine by volume. Three microliters of 0.1 M K₃[Fe(CN)₆] was added and the oxidized spectrum measured; 3-5 mg of solid Na₂S₂O₄ was then added to yield the spectra of the reduced pyridine-bound hemes. Difference spectra (reduced minus oxidized) are reported. Changes in absorbance at observed λ_{max} values were plotted against heme concentration and the data fit to a line from which extinction coefficients were obtained. Protein-bound hemes were released by suspension in a 50 mM NaOH/20% pyridine solution and their pyridine hemochrome spectra obtained similarly. Beer's law was used to estimate ε at the UV-visible absorbance maximum for the protein-associated coproheme III or heme b (Soret band λ_{max}).

UV-Visible and Resonance Raman (rR) Spectroscopy. UV-vis spectra were measured on a temperature-controlled Cary60 instrument in scanning mode at 20 °C. Samples contained 15-35 µM SaHemQ-coproheme III monomer in 50 mM KP; buffer (pH 7.4). Resonance Raman scattering was excited with 406.7 nm emission from a Kr⁺ laser. The laser beam was focused to a line parallel to the spectrograph entrance slit. Scattered light was collected in the 135° backscattering geometry (f/1) passed through a polarization filter, a holographic notch filter, and a polarization scrambler. The laser image was then focused on the entrance slit of a 0.67 m, f/4.7 Czerny-Turner spectrograph fitted with 1200 and 2400 g/mm gratings and a LN₂-cooled CCD detector (1340 \times 400 array of 20 μ m \times 20 μ m pixels, 26.8 mm × 8.0 mm image area). The spectrometer was calibrated against Raman shifts of toluene, dimethylformamide, acetone, and methylene bromide. Spectra were recorded at ambient temperature from samples prepared at $10-20 \mu M$ SaHemQ-coproheme III monomer in 50 mM KP_i in spinning 5 mm NMR tubes. UV-vis absorbance spectra were recorded from the rR samples before and after the acquisition of spectra to assess whether sample integrity had been compromised by exposure to the laser beam. The laser power at the samples ranged from 5 to 10 mW; no spectral artifacts due to photoinduced chemistry were observed at these powers.

Real-Time Monitoring of Reactions. Solutions of SaHemQ in complex with coproheme III or harderoheme III/ IV $(5-10 \ \mu\text{M})$ in 50 mM KP_i (pH 7.4) were manually mixed with oxidant from a freshly made and titrated stock (H_2O_2) , peracetic acid, mCPBA, and tert-butyl hydroperoxide). UV—vis spectra were scanned every 0.25-0.5 min. Absorbances at a given wavelength were plotted versus time and then fit by least-squares regression to single-exponential curves using KaleidaGraph. Rate constants are averages of three measurements (error = standard deviation). Resonance Raman spectra were acquired at 30 s intervals with 406.7 nm excitation. Reactions were conducted in 5 mm spinning NMR tubes at 20 °C. Difference spectra were generated by subtraction of the starting SaHemQ-coproheme III spectrum from that at each reaction time.

For discontinuous kinetic analyses, 10 μ M SaHemQ-coproheme III complex was mixed with oxidant in a 1 mL reaction mixture and aliquots were removed and quenched with 10 mM aqueous KCN [1:1 (v/v)] every 0.3 min. The quenching solution was sufficient to completely inhibit the enzymatic reaction. The heme content of quenched samples was quantified by HPLC (below).

Product Analyses by the Pyridine Hemochrome Assay, HPLC, and MS. Pyridine-hemochrome spectra were measured as described above. HPLC analyses were conducted using an Agilent 1100 series system with diode array UV—vis detection (300-700 nm); $25-50 \mu\text{L}$ aliquots of analyte were injected into a Phenomenex Luna 3c C18 (2) column (150 mm \times 4.6 mm) at a flow rate of 0.9 mL/min and 45 °C. Solvent A consisted of 1 M ammonium acetate and 10% acetonitrile; solvent B consisted of methanol and 10% acetonitrile. Elution followed a linear gradient from 35 to 95% B over 15 min. The column was washed with 95% B (2 min) and equilibrated with 35% B (6 min) between runs. ¹⁴

Standards in 50 mM KP_i (pH 7.4) were suspended 1:1 (v/v) in a 65:35 mixture of solvents A and B. For analysis of products, the same solvent was added in a 1:1 (v/v) ratio to reaction mixtures (100 µL, 10 µM protein subunit). Precipitated protein was removed by centrifugation, and supernatant was loaded into the column. Integration of the resulting peaks (400 nm) was used for construction of standard curves and quantification of the respective heme. Because of limited availability of and small impurities (<15%) in the harderoheme standards, these were not explicitly quantified; rather, integrated peak intensities from HPLC traces are reported. The exact masses of intermediates and products were verified using ESI-MS conducted on an Agilent 6538 Q-TOF instrument with a dual-ESI source (resolution of approximately 20000, accuracy of 1 ppm). Source parameters were as follows: drying gas, 12 L/min; nebulizer, 60 psi; capillary voltage, 3500 V; capillary exit, 120 V. Spectra were collected in positive mode from m/z 50 to 1000 at a rate of 1 Hz.

RESULTS

Characterization of SaHemQ Bound to Substrate, Product, and Potential Intermediates. Pure SaHemQ was obtained in yields of 15-20 mg/L of culture. Distinct UV—vis spectra for the protein reconstituted with coproheme III, harderohemes III/IV, and heme b are shown in Figure 1A. SaHemQ in complex with coproheme III has a slightly asymmetric Soret band with a maximum absorbance at 394 nm. Asymmetry may be due to partial conversion of coproheme III to harderoheme inside the protein environment even without the addition of an oxidant, as indicated by HPLC and MS of the SaHemQ-coproheme III complex (see below).

Values for absorption maxima and spectral extinction coefficients have been previously reported for the pyridine hemochrome of heme b ($\lambda_{\rm max} = 556$ nm; $\varepsilon_{556} = 33.3$ mM $^{-1}$ cm $^{-1}$) and coproheme III ($\lambda_{\rm max} = 546$ nm; $\varepsilon_{546} = 32.2$ mM $^{-1}$ cm $^{-1}$). Pyridine hemochrome absorption maxima for harderohemes III and IV were both at 549 nm (Figure S1A of the Supporting Information). Overlap in the pyridine hemochrome spectra for coproheme III and the harderohemes suggested that their separation and analysis by HPLC would be essential. Distinct HPLC retention times for all four hemes were obtained with the method described above (Figure S1B of the Supporting Information).

Using the pyridine hemochrome assay to quantify released coproheme III, $\varepsilon_{\rm Soret}$ for the $Sa{\rm HemQ}$ -coproheme III complex was determined to be $139.9\pm3.0~{\rm mM}^{-1}~{\rm cm}^{-1}$. Visible α/β and charge transfer (CT) bands occurred at 497/533 and 630 nm. The spectrum for $Sa{\rm HemQ}$ -heme b measured here was consistent with previously reported data. An extinction coefficient for the Soret band maximum at 406 nm was determined ($\varepsilon=76.6~{\rm mM}^{-1}~{\rm cm}^{-1}$). Visible bands were recorded at $510/527~{\rm nm}~(\alpha/\beta)$ and 630 nm (CT). Absorbance maxima for $Sa{\rm HemQ}$ -harderoheme III were 396 nm (Soret), 495/533 nm

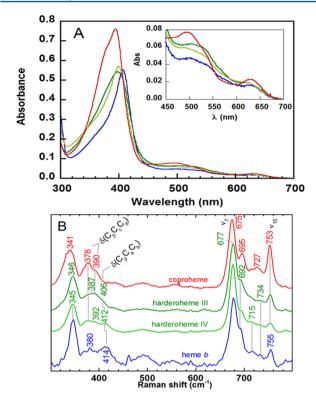


Figure 1. Characteristic spectra of SaHemQ in complex with hemes. (A) UV—vis spectra of SaHemQ in complex with ferric coproheme III (red), harderohemes III and IV (dark and light green, respectively), and heme b (blue) [5μ M coproheme, 10μ M harderoheme isomers and heme, and 50μ M KP_i (pH 7.4)]. The inset shows the visible bands on an expanded scale. (B) Low-frequency rR spectra of SaHemQ in complex with the various hemes are shown in the same colors as in panel A. Spectra were recorded with 406.7μ nm excitation of 20μ M samples in 50μ M NaP_i (pH 6.8); the propionate and vinyl bending modes are labeled.

 (α/β) , 610 nm (CT) and for SaHemQ-harderoheme (IV) 398 nm (Soret), 490/530 nm (α/β) , and 610 nm (CT). Notably, the increasingly red-shifted Soret peak maxima for the SaHemQ complexes with substrate, intermediate(s), and product are distinct.

The low-frequency rR spectra of SaHemQ complexed with coproheme III, harderohemes III/IV, and heme b are shown in Figure 1B; modes involving motions of the propionate and vinyl substituents of the various hemes are observed in this low-wavenumber region. In addition to distortions of the propionate or vinyl groups, it has been demonstrated that movement of the methyl and core atoms of the porphyrin is also involved; thus, they actually occur along with pyrrole ring deformations in these modes. ¹⁵ Nevertheless, for the purposes of this discussion, they will be termed propionate $\left[\delta(C_{\beta}C_{c}C_{d})\right]$ and vinyl $\left[\delta(C_{\beta}C_{a}C_{b})\right]$ bending.

Bands due to propionate bending modes were identified in all four spectra between 378 and 390 cm⁻¹. The four propionate groups on coproheme III present as a broad envelope with peaks at 378 and 390 cm⁻¹ in the *Sa*HemQ complex. The three propionate groups in *Sa*HemQ-harderoheme III give rise to an even less defined feature, a broad band centered at 387 cm⁻¹. The frequency of the two broad features assigned to propionate modes for *Sa*HemQ-harderoheme IV (378 and 392 cm⁻¹) is similar to those of the coproheme complex. In the case of heme *b*, the single band at 380 cm⁻¹ is assigned to a propionate bending mode for its two propionate groups. Bands associated with vinyl

bending motions appear in the harderoheme and heme b spectra between 400 and 420 cm $^{-1}$; notably, these bands are absent from the SaHemQ-coproheme III spectrum. Single $\delta(C_{\beta}C_{\rm a}C_{\rm b})$ bands are observed at 406 and 412 cm $^{-1}$ for harderohemes III and IV, respectively. While two vinyl bending bands are observed for some heme proteins, such as myoglobin, only a single feature at 414 cm $^{-1}$ is observed for SaHemQ-heme b. 16,17

The high-frequency rR spectra of hemes comprise a rich array of bands that yield insight into both structural and electronic aspects of these macrocyclic complexes. Soret-excited rR spectra are dominated by bands arising from totally symmetric (A_{1g} in D_{4h}) in-plane skeletal modes $(\nu_2, \nu_3, \text{ and } \nu_4)$. Additionally, bands assigned to the non-totally symmetric B_{1g} and B_{2g} (D_{4h}) modes, ν_{10} , ν_{11} , and ν_{12} and ν_{28} , commonly appear in the Bexcited spectra due to the dynamic Jahn-Teller effect. 18 The frequencies of all these modes are sensitive to the porphyrin core size, which is responsive to metal ion radius and the position of the metal ion with respect to the 4-N_{pyrrole} plane. These metal ion parameters are functions of oxidation and spin state, and coordination number. Additionally, Soret-excited high-frequency spectra exhibit bands assigned to modes involving distortions along local vinyl coordinates. These include $=C_bH_2$ and $-C_aH$ = out-of-plane wags ($\gamma_{=C_bH_2}$, $\gamma_{C_aH=}$) and in-plane bends $(\delta_{=\text{C}_b\text{H}_2}\!,\delta_{\text{C}_a\text{H}=}\!)$, which appear between 900 and 1300 cm^{-1} (see the labeling scheme in Scheme 2).¹⁷ Finally, bands arising from $\nu_{C,=C,J}$ the vinyl C=C stretching mode, typically appear between 1615 and 1635 cm $^{-1}$. In proteins, heme b can exhibit two $\nu_{C_{\bullet}=C_{h}}$ frequencies, likely because of the 2- and 4-vinyl groups having different dihedral angles with respect to the mean porphyrin plane. 19 Although frequencies of the in-plane skeletal bands are only slightly sensitive to oxidative decarboxylation of the 2- and 4-propionate groups of coproheme III, their intensities and the intensities of the aforementioned 2- and 4-vinyl bands facilitate tracking the appearance of the vinyl groups during the HemQ-catalyzed oxidative decarboxylation of coproheme III.

Titrimetric Reactions between SaHemQ-Coproheme III and H_2O_2 . The conversion of a propionate substituent to a vinyl group plus CO2 is a net two-electron oxidation. Thus, it was expected that two molecules of H₂O₂ (one per reactive propionate) would be required for the complete conversion of coproheme III to heme b. It was further anticipated that addition of fewer than 2 molar equiv of H₂O₂ would result in the accumulation of a singly decarboxylated intermediate. However, UV-vis and rR tracking of reactions between SaHemQcoproheme III and titrimetric additions of H₂O₂ (Figure 2A,B) revealed only partial conversion of the coproheme III complex to the heme b complex with <10 molar equiv of H_2O_2 . Indeed, 10 equiv was required to affect the complete shift of the UV-vis Soret band from 394 nm (SaHemQ-coproheme III) to 406 nm (SaHemQ-heme b). UV-visible and rR spectral changes were the same regardless of whether the 10 equiv of H_2O_2 was added at once or in titrimetric increments.

Direct observation of $\rm H_2O_2$ -mediated vinyl group formation was possible by monitoring the SaHemQ-coproheme III titration by rR (Figure 2B). The appearance of a vinyl bending band at 416 cm⁻¹ with diminution in the intensity of the propionate bending bands between 378 and 390 cm⁻¹ is consistent with the expected oxidative decarboxylation reaction. Other features typically assigned to vibrations involving the porphyrin pyrrole rings also track the extent of oxidative decarboxylation, including the increase in ν_7 (pyrrole deformation mode), ν_8 (Fe-N_{pyrrole} stretch), and ν_{11} (pyrrole asymmetric folding mode) intensities

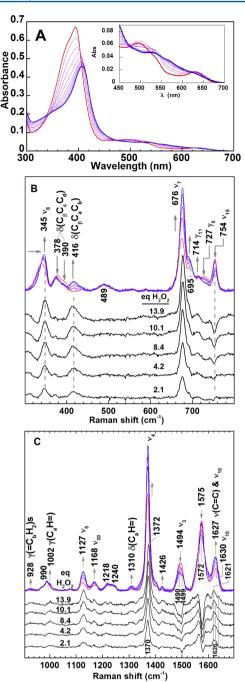


Figure 2. Spectroscopic changes upon titration of the SaHemQcoproheme III complex with H₂O₂. Reactions were monitored by (A) UV-vis spectra following the addition of 0-10 equiv of H₂O₂ (2 equiv increments) to 5 µM SaHemQ-coproheme III and 50 mM KP_i (pH 7.4). The lack of clean isosbestic points through most of the spectrum is consistent with the involvement of more than two heme complexes in the reaction. (B) Low-frequency rR spectral titration of 15 mM SaHemQ-coproheme III under the same buffer conditions as in panel A with 406.7 nm excitation. The labeling scheme used to describe the band labels in panels B and C and throughout the narrative is found in Scheme 2. (C) High-frequency range of the rR spectra shown in panel B. Molar equivalents of H₂O₂ are marked on the respective difference spectra. For panels A-C, spectra of the SaHemQ-coproheme III complexes are colored red. Partially decarboxylated species are colored purple with the final spectrum [λ_{max} (Soret) = 406 nm] colored blue. The difference spectra in panels B and C were generated by subtracting the spectrum of SaHemQ-coproheme III from the spectrum of the equilibrium reaction mixture recorded with the indicated molar equivalents of H2O2.

and the loss in intensity of ν_{15} and ν_{5} (pyrrole symmetric fold). These observations are consistent with changes occurring upon oxidative decarboxylation of propionate groups to yield vinyl substituents.

As the buildup of harderoheme was not clear in the parent rR spectra recorded with intermediate H₂O₂ stoichiometries, rR difference spectra were examined. At each point in the rR titration with H₂O₂, a single, broad band at 416 cm⁻¹ was observed, consistent with the appearance of one or more modes having vinyl bending $(\delta_{C_nC_nC_n})$ character. Given that the vinyl bending bands in the spectra of the coproheme III, harderoheme, and heme b complexes of SaHemQ (Figure 1B) are broad and within 8 cm⁻¹ of one another, they likely fall within the observed 416 cm⁻¹ difference envelope. However, when difference spectra generated by subtraction of the SaHemQ-coproheme III spectrum from those of HemQ constituted with harderohemes (Figure S2 of the Supporting Information) are compared to those observed during the reaction, the multiple features of both harderoheme spectra in this frequency range (Figure 1B) are absent. This suggests that either the SaHemQ-harderoheme complex generated via the enzymatic reaction differs from the reconstituted complex in its vinyl conformation or the harderoheme complex does not build up to a concentration that can be directly detected by these rR bands during the reaction. The latter interpretation turns out to be most consistent with the data (vide infra).

The high-frequency rR spectra also proved sensitive to structural changes caused by the oxidative decarboxylation of SaHemQ-coproheme III. Like the low-frequency spectra, the parent and difference spectra in Figure 2C can be explained largely in terms of band intensity changes, including the small apparent shifts near 1575 and 1628 cm⁻¹, which are likely due to changes in relative intensities of overlapping bands in those regions of the spectrum. Changes in the parent and difference spectra are monotonic with sequential additions of H₂O₂. Changes in the intensities of ν_2 , ν_3 , and ν_4 are largely attributed to changes in their resonance enhancements during the reaction. The increase in resonance enhancement of rR scattering by the skeletal modes is attributed to the red-shifted Soret band maximum as the constellation of peripheral substituents changes from all aliphatic ring-adjacent methyl and methylene groups to partially olefinic as the 2- and 4-vinyl groups are produced. The ν_2 region of the difference spectra results from small changes in the frequency and substantial changes in the intensity of this mode as the heme speciation progresses through harderoheme to heme *b*.

Examination of the spectra in Figure 2C also reveals the appearance and growth of small bands that can be attributed to the production of peripheral vinyl groups ($\gamma_{=C_bH_2}$, $\gamma_{C_aH=}$, $\delta_{=C_bH_2}$, $\delta_{C,H=1}$ and $\nu_{C,=C_h}$). As the heme speciation changes with sequential additions of H2O2, the intensity of three difference features changes in the $\nu_{C_a=C_h}$ range: 1621, 1630, and 1634 cm⁻¹. On the basis of depolarization ratios in spectra recorded with B and Q excitation (data not shown), the 1630 cm⁻¹ band in the spectrum of the SaHemQ-coproheme III spectrum is assigned to ν_{10} . That depolarized band shifts and undergoes a change in intensity in the Q-excited spectra of the harderoheme and heme bcomplexes, making the overlapping difference features difficult to disentangle. Nevertheless, it is clear from the spectra in Figure 2C that, as propionate groups are oxidized to vinyl groups at the porphine periphery, bands attributable to modes involving vinyl C=C stretching appear and grow as a monotonic function of added $\rm H_2O_2.$ Moreover, the 1621 cm $^{-1}\,\nu_{\rm C_a=\rm C_b}$ band is the first to

appear in the spectrum of the equilibrium reaction mixture with 2 equiv of H_2O_2 , where harderoheme III is expected to reach its maximal concentration (*vide infra*). On the basis of this order of appearance, the $1621~{\rm cm}^{-1}~\nu_{\rm C_a=C_b}$ band is assigned to the 2-vinyl group.

HPLC analysis (Figure 3) illustrated the progress of the reaction with increasing equivalents of H₂O₂ and allowed for

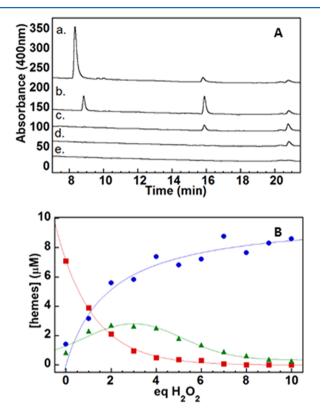


Figure 3. HPLC analysis of the products of the reaction of the SaHemQ-coproheme III complex with H_2O_2 . (A) Representative HPLC traces showing protein-free products of the reaction with 0, 2, 8, 10, and 100 equiv of H_2O_2 (a—e, respectively). Comparison of the retention times to standards (Figure S1B of the Supporting Information) identified harderoheme isomer III as the reactive intermediate. Note that the absorbance wavelength (400 nm) used for detection is closer to the $\lambda_{\rm max}$ for free coproheme III, which also has a higher extinction coefficient. (B) Quantitation of heme species involved in the HemQ reaction plotted as the percent of total heme vs equivalents of H_2O_2 added: harderoheme (green triangles), coproheme (red squares), and heme b (circles). Points are averages of three values (standard deviations all within $\pm 15\%$). Solid lines are spline curves intended to qualitatively illustrate trends in the data.

isolation and identification of the intermediate. Determination of its exact mass by ESI-MS (708.16 amu) and comparison with retention times established for authentic harderoheme standards (Figure S1B of the Supporting Information) confirmed the intermediate's identity as harderoheme isomer III (Figure 3). A plot of the integrated intensities of HPLC peaks indicated that the amount of harderoheme III reached its maximum following equilibration with 2 equiv of H_2O_2 , at which point reaction mixtures initially $10~\mu{\rm M}$ in overall heme concentration contained $2~\mu{\rm M}$ coproheme III and $5.5~\mu{\rm M}$ heme b (Figure 3B) as determined by reference to standard curves (Figure S3 of the Supporting Information). The concentration of harderoheme III at this maximum was then estimated to be $10~\mu{\rm M}$ – [coproheme III] – [heme b] = $2.5~\mu{\rm M}$. The remaining harderoheme peaks in

Figure 3B were normalized to this value. The tendency of heme b to degrade gradually in the presence of H_2O_2 and the slow catalase activity of the SaHemQ-heme b complex both suggest that these values for [harderoheme III] in Figure 3B may be modestly overestimated. Complete loss of the coproheme III HPLC peak was observed at 8 equiv of H_2O_2 , consistent with UV—vis and rR data (Figure 2). This was followed by gradual diminution of the Soret band and complete loss of the heme by 110 equiv of H_2O_2 , as measured spectroscopically and by HPLC (Figure 3 and Figure S4 of the Supporting Information). Titration of an independently generated SaHemQ-heme b complex with H_2O_2 resulted in heme destruction following addition of 100 equiv, consistent with prior work (data not shown).

Based on the quantifications in Figure 3B, the rR spectrum of SaHemQ-harderoheme III was generated using the spectra of (1) the initial SaHemQ-coproheme III, (2) the heme b product after addition of 10 equiv of H₂O₂, and (3) the equilibrium reaction mixture following addition of 2 equiv of H₂O₂ wherein all three heme complexes are present. Using the fractions of hemes determined in Figure 3B, the spectrum of harderoheme III in Figure S5 of the Supporting Information was calculated by an appropriately weighted difference. To test whether that rR spectrum is an accurate representation of the partially oxidized product, the percentages of the three hemes, quantified as described above, were used to generate linear combinations of the three component spectra and simulate the rR difference spectra for 4.2 and 8.4 equiv of H₂O₂ (Figure S5 of the Supporting Information). The striking similarity between the experimental and constructed difference spectra reveals that the spectrum of harderoheme III is indeed a component of all equilibrium reaction mixtures in the H₂O₂ titration. It therefore appears likely that, because harderoheme III does not exceed ~25% of the total heme content during the titrimetric reaction, spectral features clearly attributable to harderoheme III are not easily discerned in panels B and C of Figure 2.

Real-Time Monitoring of Reactions between SaHemQ-Coproheme III and H_2O_2 . Following addition of sufficient H_2O_2 to effect reaction (12 equiv), UV—vis spectra measured over time showed the gradual conversion of the substrate to product complex (Figure 4A). The change in absorbance at 394 nm (Soret band maximum) versus time, likely representing a composite of sequential chemical steps, could be fit to a single-term exponential function, which yielded a $k_{\rm Soret}$ of 0.30 ± 0.01 min⁻¹ ($t_{1/2} = 2.3$ min).

The reaction was subsequently monitored over time via a discontinuous method in which substrate, intermediate, and product could be independently monitored as a function of reaction time [chemical quench followed by HPLC (Figure 4B)]. Loss of coproheme III as a function of time is modeled well by a single-term exponential decay function, which yields a k_c of 2.1 \pm 0.3 min⁻¹. The rates at which harderoheme III is produced $(-k_c[\text{coproheme III}][H_2O_2])$ and converted to heme b $(k_h[\text{harderoheme III}][H_2O_2])$ must be equal when [harderoheme III] is maximal (i.e., d[harderoheme III]/dt = 0 at 0.85 min). Equating these rates and using the concentrations of coproheme III (1.8 μ M) and harderoheme III (1.3 μ M) at 0.85 min (see Figure 4B) allowed estimation of k_h at 2.9 min⁻¹ (further details in the Supporting Information). Thus, consistent with the titration data in Figure 3, harderoheme III is the observed intermediate oxidative decarboxylation product. Additionally, the rates of its appearance and decay are consistent with

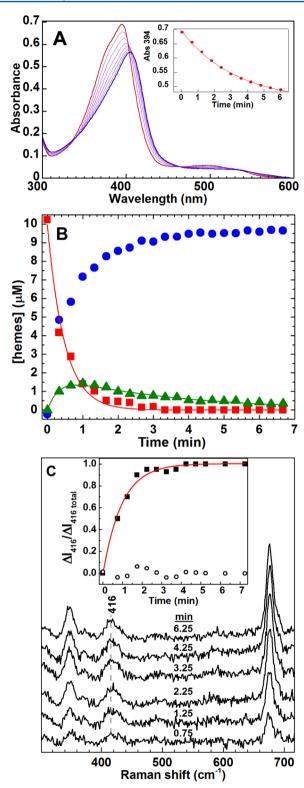


Figure 4. Time-resolved reactions between the SaHemQ-coproheme III and 12 equiv of H_2O_2 . (A) Reaction monitored by UV—vis [8 μ M enzyme and 50 mM KP_i (pH 7.4)]. For the SaHemQ-coproheme III complex (red), spectra measured every 0.25 min (purple) and the final spectrum (Soret band λ_{max} = 406 nm, blue) are shown. The absorbance at 394 nm vs time fitted to a single exponential (k_{Soret}) is shown as an inset. (B) Reaction monitored discontinuously by HPLC following a chemical quench: (red squares) coproheme III, (green triangles) harderoheme III, and (blue circles) heme b. The data describing coproheme decay were fit to a single-exponential equation (red line) to obtain a first-order rate constant (k_c). The harderoheme III progress of

Figure 4. continued

reaction was fit (green line) to a two-term exponential decay function, which was used to determine t at maximal [harderoheme III]. A first-order rate constant for conversion of harderoheme to heme b ($k_{\rm h}$) was computed as described in the text and the Supporting Information. (C) Reaction (15 μ M enzyme) monitored continuously by 406.7 nm excited rR scattering. Difference spectra recorded at the indicated times are plotted. The inset shows $\Delta I_t/\Delta I_{\rm total}$ vs time at 416 cm⁻¹ where increasing rR intensity reports vinyl formation; ΔI_t is the change in rR intensity at time t, and $\Delta I_{\rm total}$ is the total change in rR intensity over the course of the reaction. The red line shows a single-term exponential fit (first-order rate constant k_v); empty circles show residuals.

the rate of substrate, coproheme III, decay and the low maximal harderoheme III concentration during the reaction.

Formation of vinyl groups was monitored during the reaction via rR spectroscopy (Figure 4C). Assuming that the vinyl group bending modes of harderoheme III and heme b have similar resonance enhancements and frequencies, the time dependence of the 416 cm⁻¹ difference band intensity represents a composite rate constant for vinyl group appearance (harderoheme III formation and conversion to heme *b*). Thus, like the UV–visible spectra (vide supra), these data are modeled well by a single-term exponential function to yield a k_v of 1.04 \pm 0.10 min⁻¹. Despite the scatter in the rR intensities due to short acquisition times, the values of k_v and k_{Soret} which each describe the overall conversion of coproheme III to heme b, are within a factor of 4 (cf. 1.04 \pm 0.10 and 0.30 \pm 0.01 min⁻¹). Using the values of k_c and k_h estimated from the data in Figure 4B, the apparent rate constant for vinyl group formation, k_v , was estimated to be 1.6 min⁻¹ (details in the Supporting Information). Although larger than the experimentally determined values for k_v and k_{Soret} , this estimate is in sufficiently reasonable agreement to support the reaction model suggested by the titrimetric reactions in which coproheme III is converted sequentially to harderoheme III and then to

Reactions between SaHemQ Complexes with Harderohemes and H_2O_2 . SaHemQ complexes with harderohemes III and IV were subsequently titrated with H_2O_2 . Monitoring by both UV—vis and HPLC indicated conversion of each to heme b following addition of 8 and 2 equiv of H_2O_2 , respectively (Figure S6 of the Supporting Information). The rates of each reaction were monitored via changes at the Soret band maximum (Figure S7 of the Supporting Information) over time following addition of 12 equiv of H_2O_2 . The harderoheme III complex reacted with a time course comparable to that of coproheme III. The data were fitted to a single exponential with a k of 0.28 ± 0.07 min $^{-1}$, a value identical to the computed value for $k_{\rm Soret}$. The harderoheme IV isomer reacted within the mixing time.

Reaction between SaHemQ-Coproheme III and Peracetic Acid. Titration of the SaHemQ-coproheme III complex with peracetic acid (PAA) gave spectroscopic results very similar to those obtained with $\rm H_2O_2$ (Figure S8 of the Supporting Information). The completeness of each reaction and the identity of the products were verified from the HPLC and rR spectra (Figure 5 and Figure S5 of the Supporting Information). Comparison of the integrated HPLC peaks to standard curves indicated that 2 equiv of PAA resulted in decomposition of $70 \pm 2\%$ of the initially available coproheme III. Eight equivalents fully turned over the substrate, which was converted to heme b in $85 \pm 8\%$ yield. Similar behavior was observed in the rR titration. The appearance of bands assigned to vinyl modes (vide supra) was

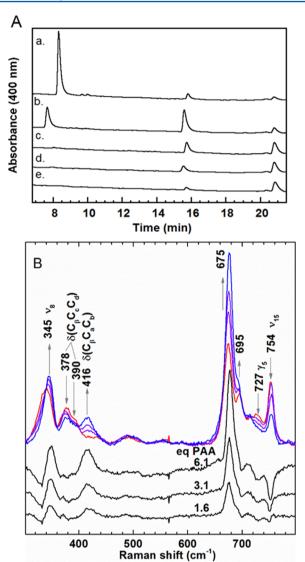


Figure 5. Titration of the SaHemQ-coproheme III complex with peracetic acid. (A) HPLC analyses of the products of the reaction of the SaHemQ-coproheme III complex with 2, 8, or 10 equiv of peracetic acid are shown. Full conversion of coproheme III to products followed addition of 10 equiv of PAA. Comparison of the retention times to those of pure standards (Figure S1B of the Supporting Information) identified harderoheme isomer III as the reactive intermediate and heme b as the final product. Note that the absorbance wavelength (400 nm) used for detection is closer to the λ_{max} for free coproheme III, which also has a higher extinction coefficient. (B) Peracetic acid titration monitored by changes in the Soret (406.7 nm) excited rR spectrum. The spectrum for SaHemQ-coproheme III is colored red; those of the partially decarboxylated complexes are colored purple, and the final spectrum is colored blue. Difference spectra were generated by subtracting the spectrum of SaHemQ-coproheme III from the spectrum obtained from the reaction mixture at equilibrium with the given molar equivalents of PAA.

consistent with the products of the reaction being the same as in the H_2O_2 reaction.

The reaction of SaHemQ-coproheme III following addition of 12 equiv of PAA was subsequently monitored by time-resolved UV-vis (Figure S9 of the Supporting Information) and rR spectroscopy (data not shown). The spectral changes were similar to those shown in Figure 4 but occurred with a notably larger rate constant ($k = 2.4 \pm 0.04 \text{ min}^{-1}$).

Reaction between SaHemQ-Coproheme III and mCPBA or tert-Butyl-OOH. Both oxidants were tested as potential HemQ cosubstrates. However, neither resulted in conversion of the SaHemQ-bound coproheme III to an identifiable harderoheme isomer or heme b. Instead, following addition of 10 equiv of either oxidant, the UV—vis spectrum of the bound coproheme III appeared to recede slightly toward the baseline, with larger numbers of equivalents producing a similar effect that was more pronounced in magnitude. HPLC analysis confirmed that heme b was not produced (Figure S10 of the Supporting Information).

DISCUSSION

Nature has devised at least three routes for the biosynthesis of heme b, $^{3-5}$ one of which appears to be common to a large group of bacteria. The terminal step of this process, catalyzed in a cofactor-independent manner by the enzyme HemQ, was studied here using equilibrium and time-resolved methods, synthetically prepared intermediates, and alternate oxidants with the goal of identifying possible intermediates.

Only 2 equiv of H_2O_2 is, in principle, necessary for the oxidative decarboxylation of two propionates to vinyl groups; however, 10 equiv was required here to completely convert SaHemQ-coproheme III to the -heme b complex, with two CO_2 molecules and water as the presumed coproducts (Figure 2). This requirement for a stoichiometric excess of H_2O_2 suggested that some of it reacts nonproductively. Consistent with that conclusion, a full equivalent of the product heme b was not observed with a stoichiometric addition of H_2O_2 (Figure 3), suggesting concurrent conversion of coproheme III, H_2O_2 -mediated degradation of heme b, and possible catalytic disproportionation of H_2O_2 . Heme b degradation was driven to completion with the addition of ~ 100 equiv of H_2O_2 (Figure S4 of the Supporting Information).

The observed inefficiency of the reaction and degradation of the product by reaction with exogenous H_2O_2 could point toward the upstream catalyst, HemY (Scheme 1), as an endogenous source of H_2O_2 . This idea is consistent with the previously documented interplay of HemY, HemH, and HemQ, where the kinetics of HemH and -Y are altered by the presence of HemQ.²⁴ It could also explain how a catalase positive organism like *S. aureus*, which avidly detoxifies H_2O_2 , might be able to manage and use H_2O_2 as an essential enzyme substrate.

With <10 equiv of $\rm H_2O_2$, coproheme III was converted to a species with several of the anticipated spectral properties of harderoheme (Scheme 2). These include rR bands associated with both vinyl and propionate bends (Figure 2). Mass spectrometry corroborated the identity of the product as a harderoheme, and chromatographic comparison to synthetic standards showed it to be exclusively isomer III (Figure 3 and Figure S4 of the Supporting Information). This suggests that, as in heme biosynthesis-associated coproporphrinogen and uroporphyrinogen oxidases, ^{25,26} the reaction occurs in an obligate clockwise direction, viewed from the distal heme pocket, with the ring A propionate converting to a vinyl first, and the ring B propionate second.

This reaction sequence is consistent with our proposed model for coproheme III binding, 11 which is based on published crystal structures of HemQ in the absence of bound substrate and structures for related chlorite dismutases bound to heme $b.^{12,27}$ An overlay of two representative structures illustrates the conservation of hydrogen bonding contacts to the nonreactive ring D propionate. By contrast, a key tyrosine residue that forms

hydrogen bonds to the ring C propionate in Clds [Y118 (Figure 6)] is conserved in the primary sequences of HemQs; however, in the substrate-free HemQ, this tyrosine (Y124) is part of an ~40-amino acid stretch of the HemQ monomer (residues 100–140) that is clearly out of alignment with the Cld structure. This stretch instead forms the upper part of a large opening in the subunit, which may be important for gating substrate entry and/or product egress. In addition to the conserved tyrosine, the

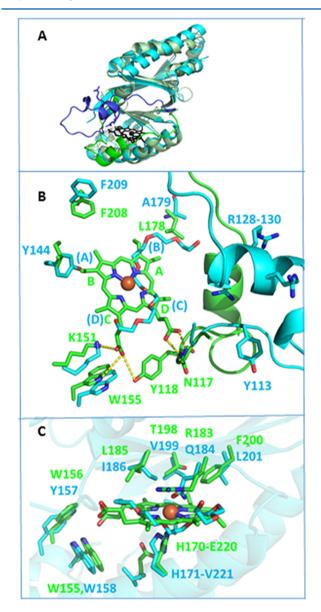


Figure 6. Overlay of the structures of a representative heme *b*-bound chlorite dismutase (carbon colored green, Protein Data Bank entry 3Q08)¹² and solvent-bound HemQ (carbon colored cyan, Protein Data Bank entry 1T0T).²⁷ (A) Monomeric subunit structures are very similar with the exception of a loop—helix region, highlighted in darker shades of blue and green. This region is located on the exterior of the HemQ homopentamer. (B) Key residues surrounding the heme *b*/coproheme III binding sites are indicated in this view overlooking the distal pocket. The placement of tetrapyrrole rings A—D in the expected orientation of the coproheme III is shown. K151, W155, N117, and Y113 (Cld numbering) are conserved in both HemQs and Clds and form contacts to the ring C/D propionates. Y113 (HemQ) is at the same sequence position as Y118 (Cld). It is part of the loop—helix region in HemQ. (C) Conserved active site Cld/HemQ residues (side view).

putative gate contains a triad of positively charged residues (arginine or lysine, R128–130 in Figure 6) that are conserved in HemQs but not Clds. These could form hydrogen bonds or salt bridges to the ring B propionate when the gate closes. A binding geometry that places coproheme III "face up" relative to the Clds' "face down" heme *b* would allow such hydrogen bonding contacts to form, while preventing potential steric clashes between a HemQ-conserved tyrosine residue (Y144).

The overall conversion of the coproheme III to heme b, monitored by changes in the UV—vis and rR spectra, is relatively slow ($t_{1/2}$ on the order of 1—2 min based on $k_{\rm v}$ or $k_{\rm Soret}$ pH 7.4, 20 °C). The time-resolved chemical quench/HPLC experiment suggests that the decarboxylation reactions have similar rate constants, with that for the reaction at ring A [2-propionate (Scheme 1)] being slightly slower than that of its ring B counterpart (4-propionate). This conclusion is consistent with the observed maximum harderoheme III of accumulation of only ~25% in the reaction between SaHemQ-coproheme III and 10 equiv of H_2O_2 (Figure 3).

In spite of harderoheme III being the apparent reaction intermediate, SaHemQ complexes with both harderoheme isomers III and IV were converted to heme b upon reaction with H₂O₂ (Figures S6 and S7 of the Supporting Information). The reaction with isomer III was slow, with a rate constant similar to that for the overall SaHemQ-coproheme III reaction (k_{Soret} or $k_{\rm w}$) under similar conditions. By contrast, the reaction with isomer IV was complete within the experimental mixing time, making it considerably faster than the conversion of the harderoheme III generated as an intermediate in the oxidation of SaHemQ-coproheme III. This reactivity pattern further suggests that reactivity of the 2-propionate group is governed by noncovalent interactions between the 4-propionate group and the enzyme (vide supra), which are absent in the SaHemQ complex with harderoheme IV. For example, the ring B 4propionate could be both stabilized and solvent-protected by salt bridges to conserved arginine residues (R128-130) when the active site closes around the coproheme III substrate. Thus, one possible explanation for the higher reactivity of harderoheme IV is that absence of the salt bridge could afford greater solvent access to the ring A propoionate, thereby accelerating the initial steps of the reaction of harderoheme IV with H₂O₂. This suggests a structure-based hypothesis in which loss of the carboxylate group from the 4-propionate and its nonbonded contacts with the active site gate helps to trigger opening of the active site. When that happens upon formation of heme b, it facilitates release of product. When it happens for the non-natural harderoheme IV, it accelerates its oxidative decarboxylation.

Acceleration of the reaction by water suggests the involvement of proton transfer in the rate-limiting step of the reaction. In canonical heme peroxidases, an active site base over the open (distal) coordination position, usually a histidine adjacent to an arginine, helps facilitate the Fe(III)/H2O2.28 The neutral imidazole side chain acts as a base toward H₂O₂, catalyzing formation of the initial Fe^{III}-OOH (Compound 0) intermediate. In peroxidase mutants lacking an active site base, formation of Compound 0 can be slowed by up to 5 orders of magnitude. 29,30 The HemQ active site is remarkably hydrophobic and lacks both a conserved basic residue and an identifiable route for proton entry and departure. Hence, the formation of Compound 0, the most plausible initial reaction product for H₂O₂ and either the coproheme III or harderoheme complexes of HemQ, might be expected to be slow or even rate-limiting, but potentially faster if water gains access to the active site.

Once formed, a Compound 0 intermediate can have at least three fates. First, with the input of a proton, the Fe^{III}O-OH bond can break heterolytically to form water and a high-valent iron-oxo intermediate (Scheme 3, pathway a). This intermediate

Scheme 3. Possible Reaction Pathways Following from Compound 0

(Compound I) is formally described as Fe^{IV} =O with an oxidizing equivalent on either the porphyrin or the side chain of an aromatic amino acid, such as tyrosine or tryptophan, capable of supporting a radical cation. In heme peroxidases, the protonated active site base acts as the acid in the Compound Iforming reaction. The conserved arginine polarizes the FeOOH moiety, electrostatically promoting movement of both of the electrons from the O-O bond toward the water leaving group.²⁴ On the proximal side of the porphyrin plane, the heme's histidine ligand is hydrogen bonded to an aspartate or glutamate side chain. The resulting anionic character of the proximal His ligand supports an additional electrostatic push toward heterolytic cleavage of the Fe^{III}O-OH bond.³¹ The HemQ active site lacks both the distal His-Arg and proximal His-Glu/Asp motifs of a typical peroxidase. Instead, it has a conserved distal glutamine surrounded by nonpolar residues and, as evinced by the 213 cm⁻¹ $\nu_{\rm Fe-His}$ frequency in the 441.6 nm excited rR spectrum of its fivecoordinate HS ferrous form (data not shown), a neutral proximal histidine due to the lack of a hydrogen-bonding Asp or Glu. In short, the HemQ active site appears to lack the typical bonded and nonbonded interactions between the heme and enzyme that drive Fe^{III}O-OH heterolysis.

However, peracetic acid, an oxygen atom donor, rapidly converted SaHemQ-coproheme III to -heme b with harder-oheme III as an intermediate. Because Compound I is expected to form when these ferric species react with a peracid, these results suggest that a species isoelectronic with Compound I may be able to form and affect the transformations of coproheme III to harderoheme III and then harderoheme III to heme b. Whether a similar iron intermediate proceeds from SaHemQ's reactions with H_2O_2 , however, is not clear. Notably, the larger oxygen atom donor mCPBA and the organic hydroperoxide, t-butyl-OOH were both ineffective at converting coproheme III to heme b (Figure S10 of the Supporting Information).

Alternatively, the FeO–OH bond of Compound 0 could break homolytically, generating Fe^{IV} =O, water, and a propionyl β -carbon radical in a stepwise (Scheme 3, pathway b) or concerted manner (pathway c). Transient generation of a propionyl β -carbon radical is an early step in some proposed mechanisms for coproporphyrinogen oxidase, ³² wherein the ring A and B propionates are converted to vinyls in an O₂-dependent fashion. These pathways both have the advantage of producing less charged intermediates that are more consistent with the hydrophobic active site. A stepwise homolytic pathway is

currently favored in heme oxygenases, 33 another enzymatic class in which a heme acts as both a substrate and a cofactor.

Interestingly, HemQs are evolutionarily and structurally related to IsdGs, which catalyze $\rm O_2$ -dependent heme degradation specifically in several groups of Gram-positive bacteria. The two proteins both possess hydrophobic heme binding domains characterized by a mutually conserved distal glutamine (HemQ) or asparagine (IsdG) and proximal histidine and tyrosine residues. Why nature evolved pathways for heme biosynthesis and degradation that are particular to Gram-positive bacteria is not clear.

ASSOCIATED CONTENT

S Supporting Information

(1) Details of the metalation and deprotection of harderoporphyrins III and IV, (2) details of the purification of SaHemQ₂ (3) HPLC and pyridine hemochrome analyses of the various hemes, (4) titrations of the SaHemQ-coproheme III complex with H₂O₂ and peracetic acid, (5) titrations and kinetic data describing the reaction of the SaHemQ-harderoheme complexes with H₂O₂, (6) data showing the reaction of SaHemQ-coproheme III with mCPBA and t-butyl-OOH, and (7) description of kinetic analysis. The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.bio-chem.5b00492.

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Notes

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ABBREVIATIONS

Cld, chlorite dismutase; DMSO, dimethyl sulfoxide; HPLC, high-performance liquid chromatography; *m*CPBA, *m*-chloroperbenzoic acid; LB, lysogeny broth; KP_i, potassium phosphate; rR, resonance Raman; *Sa, S. aureus*; SDS–PAGE, sodium dodecyl sulfate–polyacrylamide gel electrophoresis; UV–vis, ultraviolet–visible.

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