

## Eosin-Modified Reconstituted Co(II) Protoporphyrin IX Myoglobin: A Semisynthetic Photoenzyme for H<sub>2</sub> Evolution and Hydrogenation

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Photoactivation of proteins<sup>1,2</sup> and specifically of enzymes<sup>3,4</sup> has recently been a subject of intensive research efforts. Light-induced electron transfer in photosensitizer-modified redox proteins provides a general method to probe intra-protein electron transfer rates. Recently, we demonstrated that chemical modification of a redox enzyme such as glutathione reductase by the eosin chromophore turns the native enzyme into a photoactive biocatalyst.<sup>5</sup> That is, the enzyme is activated by light rather than by the native cofactor toward its specific biotransformation. Here we wish to report on a novel approach to designing photoenzymes that includes the semisynthetic modification of the myoglobin apoprotein with catalytic and photoactive sites.<sup>6</sup>

The eosin-modified Co(II)-reconstituted myoglobin, Eo<sup>2-</sup>-Mb-Co(II), was prepared by the sequence of reactions outlined in Scheme 1. The Fe(II) heme center was extracted from native myoglobin, and the Co(II) protoporphyrin IX was implanted into the apoprotein.<sup>7</sup> The resulting Co(II) protoporphyrin IX-reconstituted myoglobin was reacted with eosin isothiocyanate to yield Eo<sup>2-</sup>-Mb-Co(II). From the absorbance of the resulting protein ( $\lambda = 526$  nm,  $\epsilon = 83\,000$  M<sup>-1</sup>cm<sup>-1</sup>), the loading of Eo<sup>2-</sup> corresponds to 2 units/protein unit. Time-resolved laser flash photolysis experiments reveal that the eosin triplet is quenched by the Co(II) protoporphyrin IX site in the Eo<sup>2-</sup>-Mb-Co(II) assembly by an electron transfer pathway, eq 1. The resulting photoproducts (Eo<sup>•-</sup>,  $\lambda = 460$  nm, and Co(I) protoporphyrin IX,  $\lambda = 404$  nm) decay by the thermodynamically favored back-reaction, eq 2. Addition of Na<sub>2</sub>EDTA to the system eliminates the back-reaction by competitive oxidation of the sacrificial electron donor, and the Co(I) protoporphyrin IX is accumulated in the system, eq 3.<sup>8</sup>

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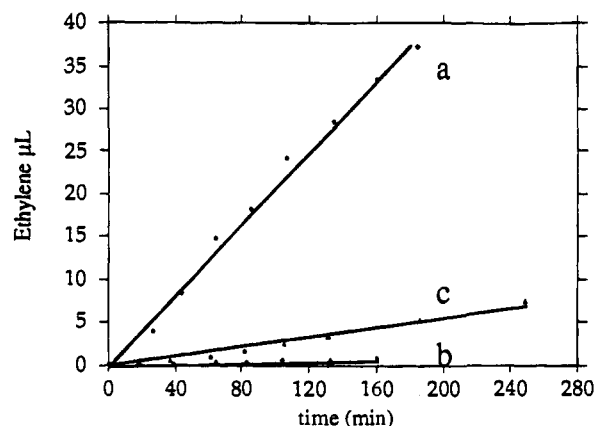
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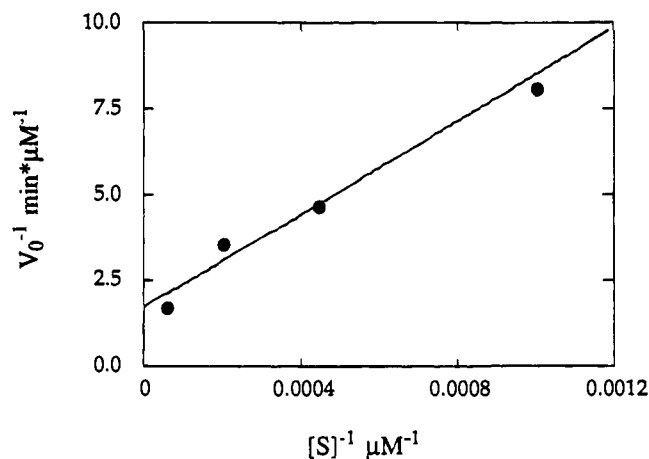
(6) For recent examples of electron transfer in reconstituted semisynthetic myoglobin, cf.: (a) Hamachi, I.; Tanaka, S.; Shinkai, S. *J. Am. Chem. Soc.* **1993**, *115*, 10458. (b) Hamachi, I.; Tajiri, Y.; Shinkai, S. *J. Am. Chem. Soc.* **1994**, *116*, 7437.

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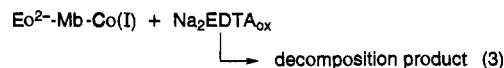
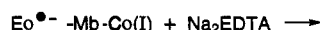
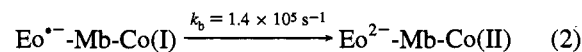
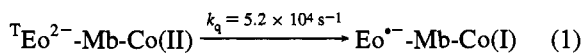
(8) The time-resolved laser experiments reveal that the second <sup>1</sup>Eo<sup>2-</sup> unit, associated with the protein, is self-quenched by ground-state Eo<sup>2-</sup> formed by the back electron transfer. The detailed analysis of these results will be provided in a later report.



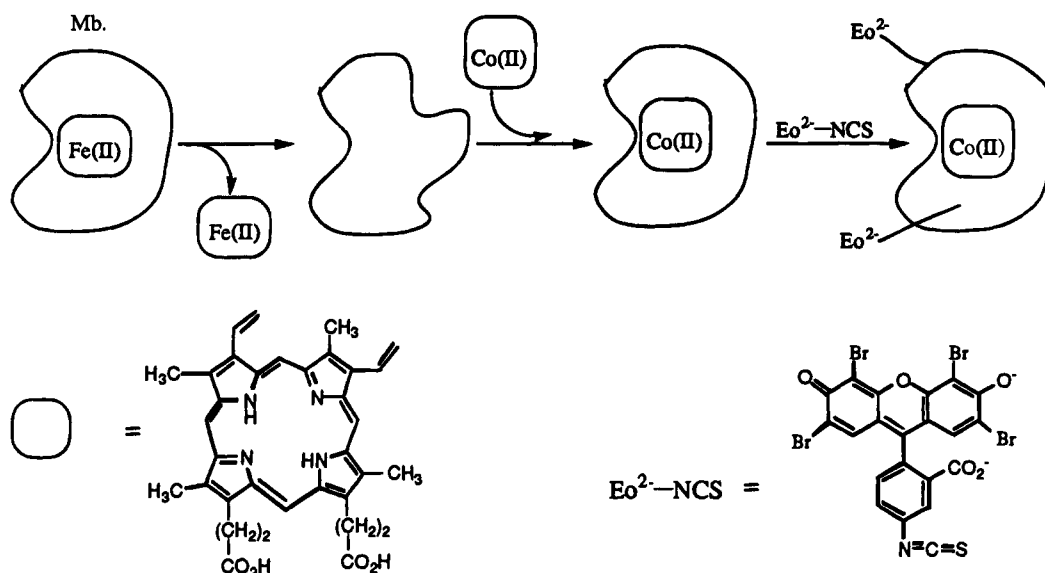
**Figure 1.** Rate of acetylene hydrogenation to ethylene at time intervals of illumination: (a) Eo<sup>2-</sup>-Mb-Co(II) system, 0.5 mgmL<sup>-1</sup> of protein; (b) Mb-Co(II) and Eo<sup>2-</sup> system, 0.5 mgmL<sup>-1</sup> of protein; (c) Co(II) protoporphyrin IX and Eo<sup>2-</sup> system. In all systems were used the bulk concentrations of Co(II), 2.5 × 10<sup>-5</sup> M and Eo<sup>2-</sup>, 5 × 10<sup>-5</sup> M. All systems consist of 3 mL of phosphate buffer, pH = 7.5, that contains Na<sub>2</sub>EDTA, 1 × 10<sup>-2</sup> M.



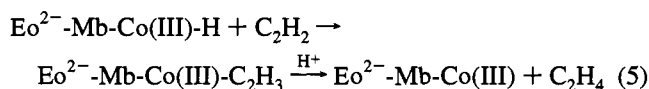
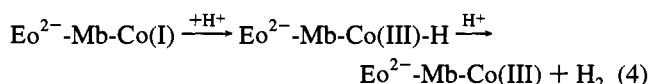
**Figure 2.** Lineweaver-Burk plot for the photohydrogenation of acetylene dicarboxylic acid in a photosystem consisting of Eo<sup>2-</sup>-Mb-Co(II) and Na<sub>2</sub>EDTA, 1 × 10<sup>-2</sup> M.



Illumination of Eo<sup>2-</sup>-Mb-Co(II) in an aqueous solution, pH = 7.5, that includes Na<sub>2</sub>EDTA, 1 × 10<sup>-2</sup> M, results in H<sub>2</sub> evolution ( $\phi = 2 \times 10^{-4}$ ). Illumination of this system under acetylene, C<sub>2</sub>H<sub>2</sub>, eliminates the formation of hydrogen, and C<sub>2</sub>H<sub>2</sub> is hydrogenated to ethylene. Figure 1 (curve a) shows the rate of C<sub>2</sub>H<sub>4</sub> evolution at time intervals of illumination,  $\lambda > 475$  nm. Control experiments revealed that H<sub>2</sub> evolution and photohydrogenation of C<sub>2</sub>H<sub>2</sub> are eliminated upon exclusion of Na<sub>2</sub>EDTA from the systems. The quantum yield for C<sub>2</sub>H<sub>4</sub> formation corresponds to  $\phi = 10^{-2}$ . For comparison, the rates of C<sub>2</sub>H<sub>4</sub> evolution from two control photosystems consisting of Mb-Co(II) and diffusional eosin (curve b) and Co(II) protoporphyrin IX (lacking the protein) and diffusional eosin (curve c) are also shown. The rate of C<sub>2</sub>H<sub>4</sub> formation in the

Scheme 1. Reconstitution of  $\text{Eo}^{2-}$ -Mb-Co(II)

$\text{Eo}^{2-}$ -Mb-Co(II) system is 70-fold and 10-fold higher than that from Mb-Co(II) lacking the covalently linked  $\text{Eo}^{2-}$  system and the system consisting of diffusional  $\text{Eo}^{2-}$  and Co(II) protoporphyrin IX that lacks the protein, respectively. Time-resolved flash photolysis experiments revealed that inefficient electron-transfer quenching<sup>9</sup> of  $^1\text{Eo}^{2-}$  by Mb-Co(II) and fast diffusion-controlled recombination of the redox photoproducts in the protein-lacking system<sup>10</sup> lead to the inefficient performance of the reference systems. Thus, organization of the photosensitizer and electron acceptor within the protein provides an assembly for effective electron-transfer quenching and for the stabilization of the photoproducts against back electron-transfer. This effective charge separation enhances the subsequent catalyzed transformation as compared to the reference systems. The  $\text{H}_2$  evolution and hydrogenation of acetylene are attributed to the formation of cobalt hydride species, eqs 4 and 5.  $\text{H}_2$  evolution



originates from the protonation of the Co-H intermediate.<sup>11</sup> Insertion of  $\text{C}_2\text{H}_2$  into Co-H species kinetically competes with hydrogen evolution and leads to its hydrogenation to  $\text{C}_2\text{H}_4$ . The effective hydrogenation of acetylene as compared to the protonation of the Co-H and  $\text{H}_2$  evolution is reflected in the substantially higher quantum yield for ethylene formation. It should be noted that in a further control experiment that included Co(II) protoporphyrin IX,  $\text{Eo}^{2-}$ , and albumin as mute protein additive, the yield of photoinduced ethylene formation was similar to that observed in the protein-lacking system. This result implies that the protein additive does not affect the

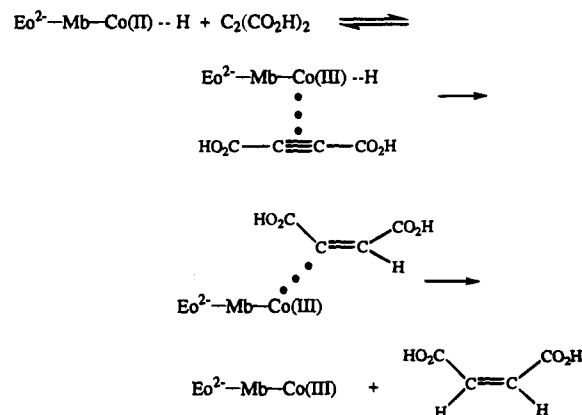
(9) The quenching rate constant of  $^1\text{Eo}^{2-}$  by Mb-Co(II) corresponds to  $K_q = 8 \times 10^6 \text{ M}^{-1}\text{s}^{-1}$ .

(10) The recombination rate constant of Co(I) protoporphyrin IX with  $\text{Eo}^{2-}$  corresponds to  $k_b = 4 \times 10^8 \text{ M}^{-1}\text{s}^{-1}$ .

(11) For other  $\text{H}_2$  evolution systems by intermediary Co-H species, cf.: (a) Brown, G. M.; Brunschwig, B. S.; Creutz, C.; Endicott, J. F.; Sutin, N. *J. Am. Chem. Soc.* **1979**, *101*, 1298. (b) Krishnan, C. V.; Creutz, C.; Mahajan, D.; Schwarz, H. A.; Sutin, N. *Isr. J. Chem.* **1982**, *22*, 98.

photoinduced hydrogenation by itself but rather the specific position of Co(II) protoporphyrin IX in the heme site of Mb controls the photocatalyzed transformation.

Photoinduced hydrogenation of acetylene dicarboxylic acid (ADA) by  $\text{Eo}^{2-}$ -Mb-Co(II) in the presence of  $\text{Na}_2\text{EDTA}$  revealed two important results: photohydrogenation of ADA produced selectively maleic acid (>92%) and only trace amounts of fumaric acid were formed (<8%). Analysis of the photohydrogenation rates at different ADA concentrations revealed that the  $\text{Eo}^{2-}$ -Mb-Co(II) system follows the Michaelis-Menten kinetic model, Figure 2,  $K_m = 4 \text{ mM}$ ,  $V_{\text{max}} = 0.6 \mu\text{M}/\text{min}$ . Thus, the Co(II)-protoporphyrin IX center associated with Mb presumably generates an active site where orientation of the Co-H insertion product is induced by the protein to stimulate the selective formation of maleic acid, eq 6.



We thus conclude that reconstitution of proteins by a chromophore and a redox-active center provides a general means to tailor photoenzymes. The activity of these assemblies takes advantage of the protein matrix in controlling charge separation and stimulating of selective chemical transformations. Preliminary studies reveal that  $\text{Eo}^{2-}$ -Mb-Co(II) acts as photocatalyst for  $\text{CO}_2$  fixation to formate and vitamin  $\text{B}_{12}$  mimicking transformations.

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