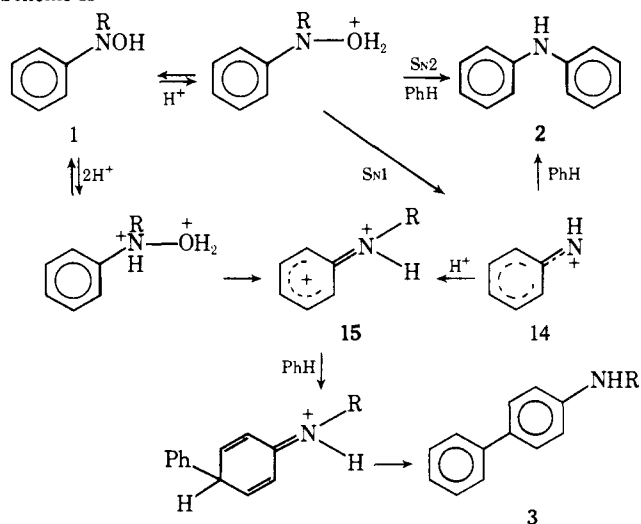
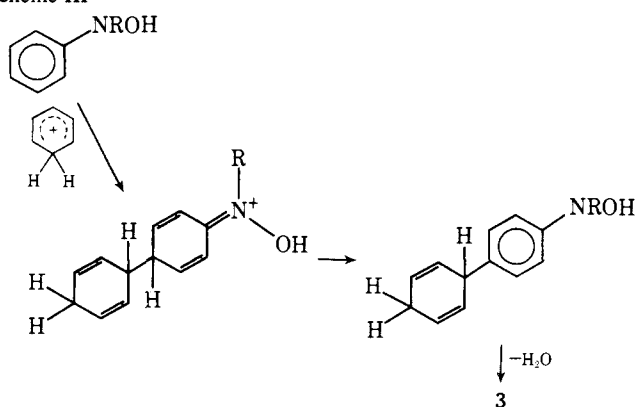


Scheme II



Scheme III



ble species generated by anodic oxidation in a strongly acidic medium,<sup>10</sup> supports the participation of the intermediate ion in this reaction.

Another mechanism which may lead to aminobiphenyls should be considered (Scheme III). This involves an electrophilic attack of benzenium ion on phenylhydroxylamine, followed by aromatization and dehydration. However, this mechanism hardly explains the formation of **8** and **11** from 4-methylphenylhydroxylamine (**7**).

## References and Notes

- (1) K. Shudo and T. Okamoto, *Tetrahedron Lett.*, 1839 (1973).
- (2) The products identified were **2** (12%), **3a** (6%), **4a** (2%), **3b** (5%), **4b** (1%), and azoxybenzene (5%). *N-tert*-Butylidiphenylamine could not be identified.
- (3) Both **2** and **3** are stable under these reaction conditions.
- (4) H. C. Brown and K. L. Nelson, *J. Am. Chem. Soc.*, **75**, 6295 (1953); L. M. Stock and H. C. Brown, *Adv. Phys. Org. Chem.*, **1**, 35 (1963).
- (5) It is possible to explain the formation of **8** by nucleophilic attack of benzene on **13**.
- (6) G. A. Olah and J. Welch, *J. Am. Chem. Soc.*, **97**, 208 (1975).
- (7) Products from these reactions are 4-aminobiphenyl, 4-amino-*p*-terphenyl, 4'-amino-*m*-terphenyl, and probably 4'-amino-*o*-terphenyl. Identification of the last compound has not been done yet.
- (8) A simple discussion that  $S_N2$  reaction leads to **2** and  $S_N1$  reaction leads to **3** does not explain the role of the acids used, though the slow reaction of **1b** in the presence of the TFA may suggest the importance of bimolecular mechanism in the TFA-catalyzed reaction which gives **2**. An alternative pathway, which cannot be eliminated, involves phenylhydroxylamine *O*-ester formed in the presence of TFSA. The ester solvolyzes to anilinium ion **14**, which reacts with benzene. Even in this case, protonation of **14** is very probable. The formation of azoxybenzene (run **4**), which is suppressed by the presence of ascorbic acid, is explained by the homolytic cleavage of the N-O bond, probably the Loeffler-Freytag type homolysis of  $PhNH_2^+-OH$ . This and the reactions of *para*-substituted phenylhydroxylamines eliminate the homolytic process in the present reactions.
- (9) P. G. Gassman, *Acc. Chem. Res.*, **3**, 26 (1970); P. G. Gassman, G. A. Campbell, and F. C. Fredrick, *J. Am. Chem. Soc.*, **94**, 3884 (1972).

(10) U. Svanholm and V. D. Parker, *J. Am. Chem. Soc.*, **96**, 1234 (1974); D. Serve, *ibid.*, **97**, 434 (1975).

Toshihiko Okamoto,\* Koichi Shudo, Toshiharu Ohta  
Faculty of Pharmaceutical Sciences, University of Tokyo  
Hongo, Bunkyo-ku, Tokyo, 113, Japan  
Received July 11, 1975

## Oxygen Binding to Iron Porphyrins

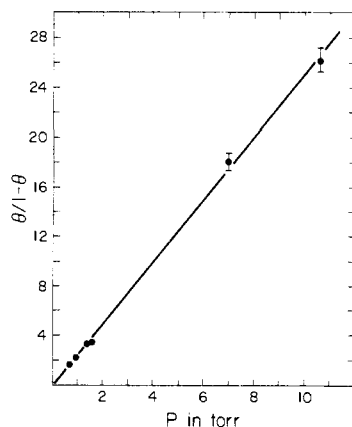
Sir:

The current literature abounds with simple synthetic models of myoglobin and hemoglobin,<sup>1-5</sup> all of which are capable of binding oxygen reversibly. In addition to reversible binding, however, a good model must be inert towards oxidation and reproduce the thermodynamic constants of the biological systems. We report here the first such comparison of a synthetic ferrous porphyrin model with myoglobin, the results of which help to delineate the role of the apoprotein in oxymyoglobin.

Using the "picket fence" porphyrin model, Fe( $\alpha,\alpha,\alpha,\alpha$ -TpvPP)(1-MeIm),<sup>4,6</sup> we have determined the enthalpy and entropy of oxygen binding in the solid state. The porosity of the crystals of this material allows the full equilibration of the solid with oxygen without the difficulties of most solution studies: the eventual irreversible oxidation of the compound and the complex equilibria with axial base. The solid-gas approach has the further advantage of a known molecular geometry as determined from x-ray diffraction data.<sup>7</sup> A simple manometric adsorption apparatus<sup>8</sup> in conjunction with an electronic manometer<sup>9</sup> was used in all experiments. The temperature of the apparatus was controlled to  $\pm 0.1^\circ\text{C}$ , while connected to a vacuum line, which could be evacuated to a pressure of  $10^{-6}$  Torr. Volumes were calibrated by expanding nitrogen from an outside gas bulb of known volume and pressure into the evacuated apparatus.

Isotherms were determined by desorption of a fully saturated<sup>10</sup> sample of Fe( $\alpha,\alpha,\alpha,\alpha$ -TpvPP)(1-MeIm)-O<sub>2</sub> into an evacuated volume. Each isotherm consists of at least six points, each point representing the extrapolation to equilibrium of 4 hr of data.<sup>11</sup> These isotherms followed the Langmuir equation at low pressures (less than 15 Torr), indicating that the iron binding sites are noninteracting in the crystal lattice. A typical plot of  $\theta/(1-\theta)$  vs.  $p_{O_2}$ , where  $\theta$  is the fraction of saturation at equilibrium, is shown in Figure 1 for the data collected at  $25.0^\circ$ . The thermodynamic enthalpy and entropy of reaction with oxygen were derived from a weighted least-squares fit to a van't Hoff plot ranging over  $50^\circ$ .<sup>12</sup> The calculated constants are  $\Delta H^\circ = -15.6 \pm 0.2$  kcal/mol and  $\Delta S^\circ = -38 \pm 1$  cal deg<sup>-1</sup> mol<sup>-1</sup> (standard state of 1 atm). The interpolated  $K_{eq}^{20^\circ}$  is 2400 atm<sup>-1</sup> or equivalently,  $p_{1/2}^{20^\circ} = 0.31$  Torr. It should be noted that some deviation from the Langmuir isotherm was observed at high pressures at  $0^\circ$ ; this is attributable to a strong physical adsorption of oxygen on the porphyrin rather than to the binding at the iron atom: the metal free ligand,  $\alpha,\alpha,\alpha,\alpha$ -H<sub>2</sub>TpvPP, physically adsorbs oxygen with an equilibrium binding constant of roughly 2 atm<sup>-1</sup> at  $0^\circ$ . It should also be noted that in contrast to solution models, our solid samples are remarkably inert to oxidation. For example, one sample has been cycled between O<sub>2</sub> and vacuum more than 200 times with no observable irreversible oxidation.

A comparison of our thermodynamic constants with those of a selection of myoglobins and of the model of Chang and Traylor<sup>3</sup> is made in Table I. It is clear that the "picket fence" is a good model for myoglobin in this respect. The close similarity of the model to the biological sys-



**Figure 1.** Isotherm of O<sub>2</sub> binding of Fe(α,α,α,α-TpivPP)(1-MeIm) at 25.0°. Error limits set from accuracy of manometer. The least-squares fit to Langmuir isotherm equation  $(\theta/(1-\theta)) = (p_{1/2})^{-1}P_{O_2}$  is shown with a slope of 2.53 Torr<sup>-1</sup> measured over the range of  $\theta = 65\%$  to  $\theta = 96\%$ .

**Table I.** O<sub>2</sub> Binding by Representative Myoglobins and Myoglobin Models

Source <sup>a</sup>	$p_{1/2}^{20^\circ}$ , Torr	$\Delta H^\circ$ , kcal/mole	$\Delta S^\circ$ , eu <sup>b</sup>
This paper	0.31	-15.6	-38
Chang and Traylor model <sup>d</sup>	0.32	...	...
Human Mb, reconst. <sup>e</sup>	0.72	-13.4	-32 <sup>c</sup>
Ox, Mb, adult <sup>f</sup>	0.55	-15	-37 <sup>c</sup>
Tuna, Mb <sup>g</sup>	0.90	-13.2	-32 <sup>c</sup>
Horse, Mb <sup>h</sup>	0.70	-13.7	-33 <sup>c</sup>

<sup>a</sup>It should be noted that some variance exists in the literature concerning these constants due to both the experimental difficulties of the myoglobin systems, as well as to the possible inherent differences between myoglobins of different species. <sup>b</sup>Standard state of O<sub>2</sub> partial pressure = 1 atm. <sup>c</sup>Calculated from reported  $p_{1/2}^{20^\circ}$  and  $\Delta H^\circ$ . <sup>d</sup>C. K. Chang and T. G. Traylor, *Proc. Nat. Acad. Sci. U.S.A.*, **72**, 1166 (1975). <sup>e</sup>A. Rossi Fanelli and E. Antonini, *Arch. Biochem. Biophys.*, **77**, 478 (1958). <sup>f</sup>A. Rossi Fanelli, E. Antonini, C. DeMarco, and S. Benerecetti, *Biochem. Hum. Genet., Ciba Found. Symp.*, **144** (1958), cited in E. Antonini and M. Brunori, "Hemoglobin and Myoglobin in Their Reactions with Ligands", American Elsevier, New York, N.Y., 1971, p 221. <sup>g</sup>A. Rossi Fanelli, E. Antonini, and R. Giuffrè, *Nature (London)*, **186**, 896 (1960). <sup>h</sup>E. Antonini and M. Brunori, "Hemoglobin and Myoglobin in Their Reactions with Ligands", American Elsevier, New York, N.Y., 1971, p 221.

tem argues that the apoprotein does not contribute significantly to the binding of oxygen,<sup>14</sup> and suggests that the primary role of the protein in myoglobin is to protect the heme from oxidation.

An independent analysis of these thermodynamic constants is possible. One can partition the entropy of an oxygen molecule into its translational, rotational, vibrational, and electronic (i.e., degeneracy) components.<sup>15</sup> Because the frequencies of the FeO<sub>2</sub> vibrations are not known, we cannot accurately calculate their contribution to the entropy of the complex; however, if we assume them to be of low frequency (400–100 cm<sup>-1</sup>), then they would contribute an additional 3 to 11 eu. If one includes the loss of the fivefold electronic degeneracy of the Fe<sup>II</sup> ( $S = 2$ ) and treats the internal rotation of the bent Fe–O–O system as a free rotor (~7 eu), then

$$\begin{aligned} \Delta S^\circ_{\text{calcd}} &= -(S_{O_2}^{\text{trans}} + S_{O_2}^{\text{rot}} + S_{O_2}^{\text{vib}} + S_{O_2}^{\text{elec}} + \\ &\quad S_{Fe}^{\text{elec}}) + S_{FeO_2}^{\text{introt}} = S_{FeO_2}^{\text{vib}} \\ &= -(36 + 11 + 0 + 2 + 3) + 7 + (7 \pm 4) \\ &= 38 \pm 4 \text{ eu (standard state of 1 atm)} \end{aligned}$$

The agreement between the calculated and experimental

values is consistent with the fit to the Langmuir isotherm, which requires independent binding sites, and with the absence of substantial systematic errors in the experimental determination. This demonstration that the "picket fence" porphyrin is a well-behaved system for solid-gas equilibrium studies sets the stage for further studies of parameters such as the nature and closeness of the axial base and the polarity of the oxygen binding site.

**Acknowledgment.** We thank Drs. Lowell Wood and Fred Aldridge of Lawrence Livermore Laboratory for the loan of the Barocell, Dr. Susan Hayes for the preparation of iron porphyrin, and the Fannie and John Hertz Foundation for fellowship support (K.S.S.). This work was supported by the National Science Foundation Grant MPS75-17018.

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- Abbreviations used in this work include: α,α,α,α-H<sub>2</sub>TpivPP, meso-tetra(α,α,α,α-o-pivalamidophenyl)porphyrin; Mb, myoglobin; 1-MeIm, 1-methylimidazole.
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- Barocell Model 523CH-12, from Datametrics, 340 Fordham Rd., Wilmington, Mass. 01887, whose ranges are 0–1, 0–10, 0–100, and 0–1000 Torr with a 3½ digit readout.
- Exposure of the sample to 300 Torr of O<sub>2</sub> for 15 hr resulted in better than 99% of stoichiometric binding of O<sub>2</sub>.
- The half-life of approach to equilibrium was roughly an hour.
- Based on  $\ln K_{eq}$  (°C): 5.34 ± 0.02 (50.0); 6.25 ± 0.05 (37.0); 7.56 ± 0.02 (25.0); 9.76 ± 0.13 (–0.1), with a standard state of 1 atm; error limits of these  $\ln K_{eq}$  are the standard deviation of the least-squares fit to the Langmuir isotherm at each temperature. Error limits of  $\Delta H$  and  $\Delta S$  derived using the extremum method of Benson<sup>13</sup> yield maximum error limits of ±0.5 kcal/mol and ±1.6 eu, respectively. It should be noted that the statistical error limits given are derived from the least-squares fit and may not necessarily reflect the actual experimental error. The reproducibility of the equilibrium constants between separately synthesized samples is less than that of a single sample;  $K_{eq}$  varied less than 10% between samples, corresponding to a variation of less than 2% in  $\ln K_{eq}$ .
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- It is not necessary to invoke hydrogen bonding from the distal histidine to O<sub>2</sub> in order to explain the stability of O<sub>2</sub> binding in biological systems, since this model system lacks that feature.
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James P. Collman,\* John I. Brauman, Kenneth S. Suslick

Department of Chemistry, Stanford University  
Stanford, California 94305

Received September 2, 1975

## Mechanism of the Cobalt Carbonyl-Catalyzed Homogeneous Hydrogenation of Aromatic Hydrocarbons<sup>1</sup>

Sir:

A variety of polycyclic aromatic hydrocarbons (PAH) are homogeneously hydrogenated in a highly selective manner<sup>2–4</sup> in the presence of Co<sub>2</sub>(CO)<sub>8</sub> and synthesis gas (CO + H<sub>2</sub>) at elevated temperature and pressure ("oxo" conditions). The operation of this catalyst, being one of the few known homogeneous catalysts for aromatic hydrogenation, has special interest. By analogy with the generally accepted mechanism of hydroformylation of olefins with the same catalyst it has been assumed<sup>5</sup> that the key reaction is the