

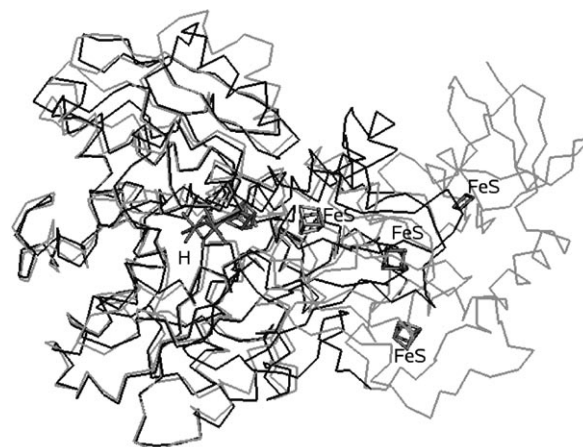
# Hydrogen-Activating Enzymes: Activity Does Not Correlate with Oxygen Sensitivity\*\*

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Hydrogenases ( $H_2$ ases) catalyze a reaction that is essential in the energetics of many bacteria and has promising technological applications: the reversible conversion between dihydrogen and protons. These enzymes are studied in various contexts, including bioenergetics and inorganic catalysis, but the main motivation is that the knowledge we shall acquire by studying them will prove useful for designing the catalysts we need to produce hydrogen from water in a clean process and to oxidize it in affordable fuel cells. However, the fact that  $H_2$ ases are inhibited by  $O_2$  hinders technological developments and much effort has been devoted to understanding the molecular basis of  $O_2$  sensitivity.<sup>[1,2]</sup> Herein, we show that a very active  $H_2$ ase, which was believed to be highly oxygen sensitive, actually reacts very slowly with  $O_2$  as a result of simultaneous reactions which we were able to resolve and quantify.

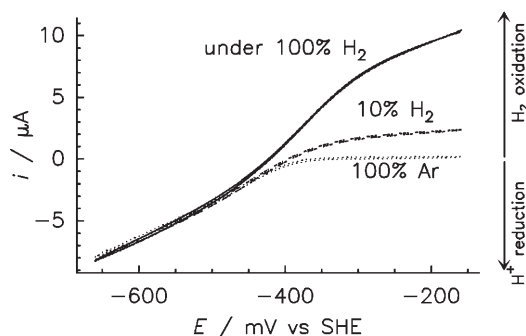
Hydrogen-activating enzymes are classified as NiFe, Fe, and FeFe  $H_2$ ases according to the metal content of their active site. Clostridial-type FeFe  $H_2$ ases house four iron–sulfur (FeS) clusters in addition to the active-site “H-cluster”. *Clostridium acetobutylicum* (Ca) FeFe  $H_2$ ase oxidizes  $H_2$  faster<sup>[3]</sup> than the thoroughly characterized enzyme from *Desulfovibrio desulfuricans* (Dd), which has a similar structure except that the electron-transfer (ET) chain consists of two FeS clusters rather than four (Figure 1).

We have used the procedure described in reference [3] to express homologically and purify by affinity chromatography to apparent homogeneity (as determined by sodium dodecyl-sulfate PAGE and elemental analysis) the strep-tagged FeFe  $H_2$ ase (HydA1) from Ca (see the Supporting Information).



**Figure 1.** Structures of *D. desulfuricans* (Dd)  $H_2$ ase (Protein databank (PDB) file: 1hfe; dark gray) and *C. pasteurianum*  $H_2$ ase (Cpl; PDB file: 1feh; light gray). The latter is highly homologous to *C. acetobutylicum* (Ca)  $H_2$ ase (HydA1).<sup>[4]</sup>

This enzyme spontaneously adsorbs onto graphite in a configuration that allows direct ET to and from the electrode. The steady-state voltammograms in Figure 2 show that Ca



**Figure 2.** Steady-state cyclic voltammograms for Ca  $H_2$ ase adsorbed onto graphite. Conditions: pH 7,  $T = 30^\circ\text{C}$ ,  $V = 20\text{ mVs}^{-1}$ ,  $\omega = 2\text{ krpm}$ .

$H_2$ ase efficiently catalyzes both  $H_2$  oxidation and production. The reductive current is independent of  $H_2$  pressure, which shows that  $H^+$  reduction is not inhibited by the product,  $H_2$ . The Michaelis constant ( $K_m$ ) for  $H_2$  is high (about 0.8 atm or 0.6 mm). The activity decreases at potentials above 80 mV versus the SHE at pH 7, a result that reveals the reversible anaerobic formation of the inactive state of the H-cluster

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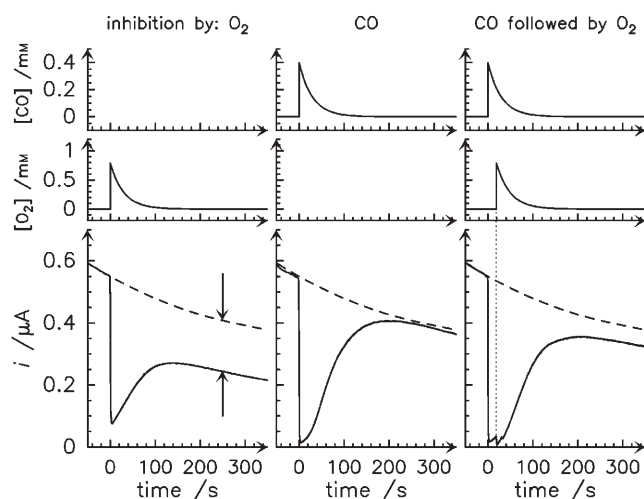
[\*\*] We acknowledge Sébastien Dementin for help and fruitful discussions, funding from the ANR, and support from the Pôle de Compétitivité Capénergies.

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called  $H_{ox}^{inact}$ . All of the above observations are similar to those made with *Dd*  $H_2ase$ .<sup>[2]</sup> However, we will show that great differences appear when the enzymes are exposed to oxygen.

We have developed a method to characterize the reaction of redox enzymes with gaseous inhibitors;<sup>[5,6]</sup> this method consists of poisoning the enzyme-coated electrode at a given potential and then injecting into the electrochemical cell an aliquot of solution saturated with inhibitor while the cell is continuously degassed by bubbling through a different gas, for example,  $H_2$ . The transient current (that is, the activity) is monitored while the concentration of inhibitor is decreasing. With this decay being exactly exponential, the quantitative analysis of the data is relatively straightforward.

In Figure 3, each column depicts a different experiment. The lower panels show the changes in activity that follow the addition of inhibitors as depicted in the upper panels. The cell

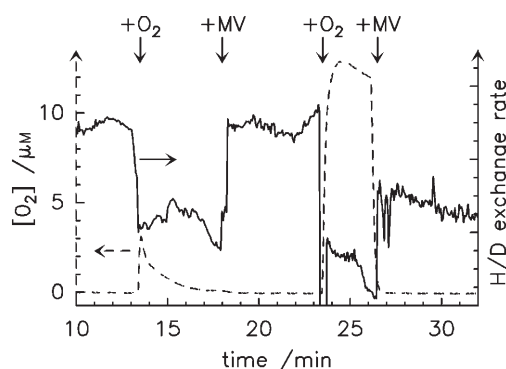


**Figure 3.** Activity transients following the addition of CO and/or  $O_2$  under  $H_2$  (1 atm). Conditions: pH 7,  $T = 30^\circ C$ ,  $E = 200$  mV,  $\omega = 2$  krpm. The dashed line is a guide for the eye. The current transients in the middle and right columns have been multiplied by 0.83 and 0.95, respectively.

was flushed with  $H_2$  and the electrode was poised at 200 mV; the enzyme slowly inactivates in the absence of inhibitor (at  $t < 0$ ) upon formation of  $H_{ox}^{inact}$ . In the experiment shown on the left in Figure 3, the oxygen concentration was stepped to  $[O_2]_0 = 0.8$  mM at  $t = 0$ , which resulted in instant inhibition. Two observations from this experiment contrast with the behavior observed with the enzyme from *Dd*. Firstly, despite the high level of  $O_2$ , inhibition is not complete: an  $O_2$  concentration that is ten times lower causes complete inhibition of *Dd* FeFe  $H_2ase$  (bottom panel of Figure 5 in reference [7]). Secondly, although a fraction of the activity is irreversibly lost after the oxygen has vanished (at the point marked by arrows), a surprisingly large part of the activity is recovered. When the inhibition was induced by CO (middle column in Figure 3), the irreversible inactivation was not observed, a result indicating that this effect is specific to the reaction with  $O_2$ . Hence, oxidative inhibition of *Ca*  $H_2ase$  consists of three distinct processes: 1) the anaerobic formation of  $H_{ox}^{inact}$  (which is reverted only by reduction),<sup>[2]</sup> 2) the

reversible binding of inhibitor  $O_2$ , and 3) an irreversible inactivation by  $O_2$ . To determine whether the latter process results from  $O_2$  damaging the electron-transferring FeS clusters or the active site, we ran an experiment where  $O_2$  was added while CO was present (right-hand column in Figure 3). CO is a competitive inhibitor of the enzyme, so it should bind to and protect the active site. The fact that the recovery of activity after aerobic inactivation is greater in this experiment shows that the oxygen damage is due to degradation of the H-cluster rather than the other FeS clusters.

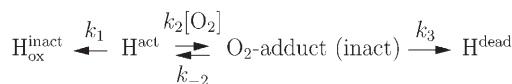
This conclusion is supported by an independent experiment in which the isotope-exchange activity (the formation of HD and  $H_2$  from  $D_2$ )<sup>[8]</sup> was monitored in the presence of  $O_2$ . The H/D activity decreased only 50 and 70 % in the presence of 3 or 13  $\mu M$   $O_2$ , respectively, and it was mostly (at low  $[O_2]$ ) or partly (at high  $[O_2]$ ) recovered when  $O_2$  was removed by adding reduced methyl viologen (Figure 4). Since isotope exchange does not involve ET through the FeS clusters, this



**Figure 4.** Inhibition of isotope exchange by  $O_2$  in solution at pH 7 and  $30^\circ C$ . The activity of the sample at time zero was  $50 \mu mol min^{-1} mg^{-1}$  for both  $H_2$  production and  $H^+/D_2$  exchange. MV: methyl viologen.

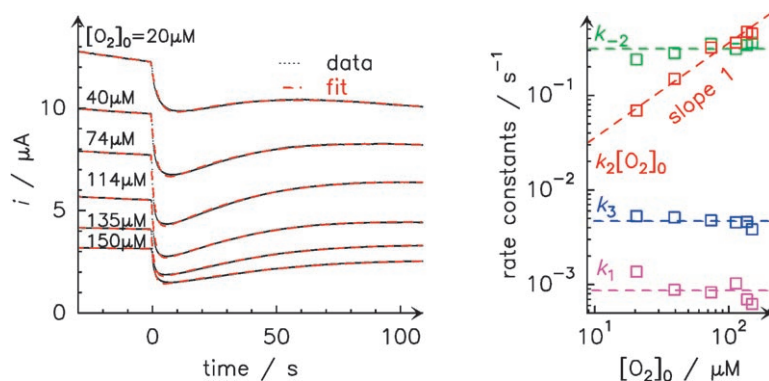
shows that  $O_2$  affects the H-cluster. Moreover, since this experiment is carried out in solution, it demonstrates that the oxygen tolerance seen in electrochemistry does not result from the enzyme being adsorbed on (and protected by) the electrode.

Hence, our experiments demonstrate that  $O_2$  inhibition proceeds by reversible formation of an  $O_2$  adduct followed by irreversible transformation of the active site, as depicted in Scheme 1.



**Scheme 1.** Oxidative inhibition of *Ca*  $H_2ase$ .

We now turn to simulations of chronoamperometric data which allow *quantitative* characterization of the inactivation processes. In Figure 5, current transients resulting from the addition of various amounts of  $O_2$  were fitted to the current equation calculated from Scheme 1, with the exponential decays of  $O_2$  concentration taken into account:  $[O_2] =$



**Figure 5.** Fit of chronoamperometric data to the solution of the equation described by Scheme 1. The adjusted rate constants are plotted against added  $O_2$  in the right panel. Conditions: 1 bar  $H_2$ , pH 7, 30°C, 2 krpm.

$[O_2]_0 \exp(-t/\tau)$ . In addition to the value of  $\tau$ , four rate constants were determined,  $k_1$ ,  $k_2[O_2]_0$ ,  $k_{-2}$ , and  $k_3$ , which span three orders of magnitude. Consistently with Scheme 1, the value of  $k_2[O_2]_0$  was found to be proportional to the concentration of added  $O_2$ , whereas all three other rate constants were independent of  $[O_2]_0$ . We determined that  $k_1 = 10^{-3} \text{ s}^{-1}$ ,  $k_2 = 3.2 \text{ s}^{-1} \text{ mM}^{-1}$ ,  $k_{-2} = 0.3 \text{ s}^{-1}$ ,  $k_3 = 4 \times 10^{-3} \text{ s}^{-1}$ .

The fact that purification of active clostridial  $H_2$ ases requires strictly anaerobic conditions has always been interpreted as revealing the extreme oxygen sensitivity of these enzymes. On the contrary, our data and quantitative analysis show that this enzyme is less sensitive to oxidative inactivation than most other  $H_2$ ases:

- We estimate that the rate of anaerobic oxidation ( $k_1 = 10^{-3} \text{ s}^{-1}$ ) is about an order of magnitude slower than that for the enzyme from *Dd*; this inertia is actually detrimental to *Ca*  $H_2$ ase since the  $H_{ox}^{inact}$  state is protected against  $O_2$  and can be reactivated by reduction.<sup>[2]</sup>
- Although oxygen irreversibly inactivates the enzyme from *Dd*, it binds reversibly to *Ca*  $H_2$ ase ( $k_{-2}/k_2 = 0.1 \text{ mM}$ ). This is usually typical of  $O_2$ -resistant enzymes.<sup>[2]</sup>
- The oxidation that is responsible for irreversibly inactivating the enzyme when it is purified under air is surprisingly slow ( $k_3 = 4 \times 10^{-3} \text{ s}^{-1}$ ).

- The apparent bimolecular rate constant for the reaction with  $O_2$ ,  $k_2 k_3 / (k_3 + k_{-2}) = 4 \times 10^{-2} \text{ s}^{-1} \text{ mM}^{-1}$ , is much smaller than that for prototypical NiFe  $H_2$ ases ( $32 \text{ s}^{-1} \text{ mM}^{-1}$  in reference [5]), which are usually said to be more resistant to  $O_2$ .

The molecular basis of this relative resistance must now be determined. The enzymes from *Dd* and *Ca* are very similar in the region surrounding the active site (Figure 1).<sup>[4]</sup> Yet we have shown that their active sites react very distinctively with  $O_2$ . *Ca*  $H_2$ ase is one of the very few  $H_2$ ases for which efficient molecular-biology tools are now available.<sup>[9]</sup> The precise characterization of the oxidative reactions described herein will guide our attempts to engineer the enzyme in order to

slow further the reactions that make it sensitive to  $O_2$ .

Received: September 18, 2007

Revised: November 21, 2007

Published online: February 1, 2008

**Keywords:** electron transfer · hydrogenases · inhibition · kinetics · protein film voltammetry

- [1] A. L. De Lacey, V. M. Fernandez, M. Rousset, R. Cammack, *Chem. Rev.* **2007**, *107*, 4304–4330.
- [2] K. A. Vincent, A. Parkin, F. A. Armstrong, *Chem. Rev.* **2007**, *107*, 4366–4413.
- [3] M. Demuez, L. Cournac, O. Guerrini, P. Soucaille, L. Girbal, *FEMS Microbiol. Lett.* **2007**, *275*, 113–121.
- [4] J. C. Fontecilla-Camps, A. Volbeda, C. Cavazza, Y. Nicolet, *Chem. Rev.* **2007**, *107*, 4273–4303.
- [5] C. Léger, S. Dementin, P. Bertrand, M. Rousset, B. Guigliarelli, *J. Am. Chem. Soc.* **2004**, *126*, 12162–12172.
- [6] M. G. Almeida, C. M. Silveira, B. Guigliarelli, P. Bertrand, J. J. G. Moura, I. Moura, C. Léger, *FEBS Lett.* **2007**, *581*, 284–288.
- [7] K. A. Vincent, A. Parkin, O. Lenz, S. P. J. Albracht, J. C. Fontecilla-Camps, R. Cammack, B. Friedrich, F. A. Armstrong, *J. Am. Chem. Soc.* **2005**, *127*, 18179–18189.
- [8] L. Cournac, G. Guedeney, G. Peltier, P. M. Vignais, *J. Bacteriol.* **2004**, *186*, 1737–1746.
- [9] L. Girbal, G. von Abendroth, M. Winkler, P. M. C. Benton, I. Meynial-Salles, C. Croux, J. W. Peters, T. Happe, P. Soucaille, *Appl. Environ. Microbiol.* **2005**, *71*, 2777–2781.