

# Approaches to developing biological H<sub>2</sub>-photoproducing organisms and processes

M.L. Ghirardi<sup>\*1</sup>, P.W. King<sup>\*</sup>, M.C. Posewitz<sup>\*†</sup>, P. Ching Maness<sup>\*</sup>, A. Fedorov<sup>\*</sup>, K. Kim<sup>\*</sup>, J. Cohen<sup>‡</sup>, K. Schulten<sup>‡</sup> and M. Seibert<sup>\*</sup>

<sup>\*</sup>National Renewable Energy Laboratory (NREL), Golden, CO, U.S.A., <sup>†</sup>Colorado School of Mines, Golden, CO, U.S.A., and <sup>‡</sup>Beckman Institute, University of Illinois, Urbana-Champaign, IL 61801, U.S.A.

## Abstract

The development of efficient biological systems for the direct photoproduction of H<sub>2</sub> gas from water faces several challenges, the more serious of which is the sensitivity of the H<sub>2</sub>-evolving enzymes (hydrogenases) to O<sub>2</sub>, an obligatory by-product of photosynthesis. This high sensitivity is common to both FeFe and NiFe hydrogenases, and is caused by O<sub>2</sub> binding to their respective metallocatalytic sites. This overview describes approaches to (i) molecular engineering of algal FeFe-hydrogenase to prevent O<sub>2</sub> access to its catalytic site; (ii) transform a cyanobacterium with an O<sub>2</sub>-tolerant bacterial NiFe hydrogenase or (c) partially inactivate algal O<sub>2</sub>-evolution activity to create physiologically anaerobiosis and induce hydrogenase expression.

## Introduction

Biological H<sub>2</sub> production linked to photosynthetic water oxidation is a promising technology that may play a major role in the future of renewable energy [1,2]. Photosynthetic microorganisms, such as green algae and cyanobacteria, have the potential to store efficiently the energy of incident sunlight as high-energy H<sub>2</sub> molecules, with a maximum theoretical efficiency of approx. 13%. To achieve sustained H<sub>2</sub> production, however, photosynthetic organisms need to be able to produce H<sub>2</sub> gas directly from water at maximum photosynthetic efficiency. This presents a major challenge due to the high sensitivity of hydrogenases to O<sub>2</sub> inactivation [3]. Three approaches are being pursued at the NREL (National Renewable Energy Laboratory) to address this issue. In this overview, we describe these approaches and discuss recent advances in each.

## Engineering an O<sub>2</sub>-tolerant Fe-hydrogenase

In the green alga *Chlamydomonas reinhardtii*, H<sub>2</sub> photoproduction is catalysed by one or two Fe-hydrogenases [4,5]. Both enzymes are transcriptionally and post-translationally regulated by O<sub>2</sub> concentration and by other still unidentified metabolic factors [3,6]. These observations agree with the fact that Fe-hydrogenases are more sensitive to O<sub>2</sub> than their NiFe counterparts [7]. In the literature, one finds two main determinants of hydrogenase O<sub>2</sub> sensitivity: the structure of the catalytic site [8] and the accessibility of the catalytic site to O<sub>2</sub> gas [9]. In our work, we have focused on the latter determinant, and initiated detailed studies of gas diffusion through Fe-hydrogenases to identify the most probable pathways for

O<sub>2</sub> and H<sub>2</sub> gases. Our molecular dynamics simulations are based on the crystal structure of the highly homologous *Clostridium pasteurianum* CpI Fe-hydrogenase, due to the lack of a solved structure for the algal enzymes. Preliminary experiments have identified a few potential individual pathways for O<sub>2</sub> gas diffusion, but multiple pathways for H<sub>2</sub> diffusion (see Cohen et al. in this colloquium [9a]). These results are in contrast with the observation that there is a single H<sub>2</sub> channel that links the catalytic site to the surface of the enzyme, as proposed for the *Desulfovibrio desulfuricans* Fe-hydrogenase by probing its X-ray structure with a 0.75 Å (1 Å = 0.1 nm) probe [9]. The difference between the two observations may be due to the dynamic nature of our simulations, as opposed to the static assumptions of [9]. Nevertheless, our results underscore the possibility of selectively affecting O<sub>2</sub> diffusion into the enzyme without disrupting H<sub>2</sub> gas diffusion from the catalytic site. Further *in silico* and *in vitro* experiments are being developed to test this hypothesis, taking advantage of a recently developed system for expression of Fe-hydrogenases in *Escherichia coli*.

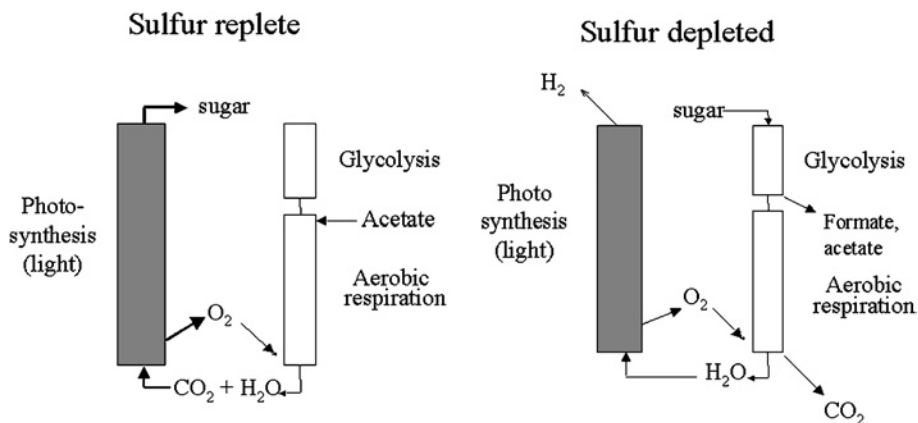
## Construction of a recombinant cyanobacterium expressing an O<sub>2</sub>-tolerant NiFe-hydrogenase

In contrast with green algae, cyanobacteria are photosynthetic microbes that contain NiFe-hydrogenases. Until recently, it was not clear whether photosynthetic reductants could be directly utilized for hydrogenase-catalysed H<sub>2</sub> production in these microbes [10]. However, Cournac et al. [11] recently demonstrated that a *Synechocystis* strain, deficient in the type I NADPH-dehydrogenase complex was able to produce a significant amount of H<sub>2</sub> in the light. This observation opens up the door to explore cyanobacteria as an additional potential system for H<sub>2</sub> photoproduction from water.

**Key words:** biohydrogen, cyanobacteria, green algae, hydrogenase, O<sub>2</sub> inactivation, sulphur deprivation.

<sup>1</sup>To whom correspondence should be addressed (email maria\_ghirardi@nrel.gov).

Figure 1 | Physiological effects of sulphur deprivation



Research conducted at NREL has uncovered an  $O_2$ -tolerant NiFe hydrogenase from the purple non-sulphur photosynthetic bacterium, *Rubrivivax gelatinosus* CBS. The *in vivo* half-life of this hydrogenase is 21 h in air, and 6 h in air when the protein is partially purified [12]. Genes encoding the CBS hydrogenase and its accessory proteins were sequenced recently. Work has been initiated to transfer and express the  $O_2$ -tolerant hydrogenase genes from CBS to *Synechocystis* so that the resultant recombinant organism can evolve  $H_2$  continuously during normal oxygenic photosynthesis.

### Green algal $H_2$ production following partial inactivation of photosynthetic $O_2$ -evolution activity

A third approach to sustain algal  $H_2$  photoproduction consists of depriving an algal culture of sulphate, required to provide proteins with sulphurylated amino acid residues. The protein most affected by sulphur deprivation in green algae is D1, the components of the water-oxidizing Photosystem II that displays the fastest rate of turnover. As the cultures move from sulphur-replete to sulphur-deprived medium, D1 turnover decreases, causing an overall reduction in photosynthesis [13]. Indeed, we have demonstrated that, within 24 h, the photosynthetic capacity of the cultures decreases below that of respiratory  $O_2$  consumption. As shown in Figure 1, as a result, the culture becomes anaerobic,  $CO_2$  fixation is interrupted and the accumulated starch is degraded. Depending on the rate of starch degradation, the cultures can either (i) oxidize all starch to  $CO_2$ , thereby consuming the residual photosynthetically evolved  $O_2$ , or (ii) resort to partial anaerobic metabolism as well, excreting formate and acetate into the medium [14].

The  $H_2$ -production phase is temporary, due to the eventual effect of sulphur deprivation on all other cellular functions. We have demonstrated that the cultures can cycle between a sulphur-replete, photosynthetic  $O_2$  evolution mode and the sulphur-depleted,  $H_2$ -photoproduction mode for at least three times without significant loss of activity [15]. However,

preliminary economic analyses at NREL attributed significant costs to the cycling of cultures, which led us to develop a continuous, chemostat-based system for algal  $H_2$ -production [16]. This latter development resulted in a decrease in the cost of  $H_2$  production by a factor of three. Finally, current developments include attempts to immobilize sulphur-deprived cells on to glass fibres at high cell density, which has resulted in increased stability and increased volumetric rates of  $H_2$  production.

### Conclusion

Although still under investigation, biological  $H_2$  photoproduction represents a potentially revolutionary technology to harvest solar energy and store it in an easily transportable chemical form. The  $O_2$  sensitivity of the hydrogenase enzyme is, however, only one of the many obstacles to the development of a truly commercial system. Major developments in algal physiology and genetics, electron transport, light-harvesting optimization and photobioreactor engineering will also be required to achieve efficient coupling of photosynthetically generated reductants with  $H_2$  gas production.

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### References

- Melis, A. (2002) *Int. J. Hydrogen Energy* **27**, 1217–1228
- Levin, D.B., Pitt, L. and Love, M. (2004) *Int. J. Hydrogen Energy* **29**, 173–185
- Ghirardi, M.L., Togasaki, R.K. and Seibert, M. (1997) *Appl. Biochem. Biotechnol.* **63–65**, 141–151
- Happe, T. and Kaminski, A. (2002) *Eur. J. Biochem.* **269**, 1022–1034
- Forestier, M., King, P., Zhang, L., Posewitz, M., Schwarzer, S., Happe, T., Ghirardi, M.L. and Seibert, M. (2003) *Eur. J. Biochem.* **270**, 2750–2758

- 6 Posewitz, M.C., Smolinski, S.L., Kanakagiri, S., Melis, A., Seibert, M. and Ghirardi, M.L. (2004) *Plant Cell* **16**, 2151–2163
- 7 Frey, M. (2002) *ChemBioChem* **3**, 153–160
- 8 Bleijlevens, B., Buhrke, T., van der Linden, E., Friedrich, B. and Albracht, S.P.J. (2004) *J. Biol. Chem.* **279**, 46686–46691
- 9 Nicolet, Y., Piras, C., Legrand, P., Hatchikian, C.D. and Fontecilla-Camps, J.C. (1998) *Structure* **7**, 12–23
- 9a Cohen, J., Kim, K., Posewitz, M., Ghirardi, M.L., Schulten, M. and King, P. (2005) *Biochem. Soc. Trans.* **33**, 80–82
- 10 Tamagnini, P., Axelsson, R., Lindberg, P., Oxelfelt, F., Wünschiers, R. and Lindblad, P. (2002) *Microbiol. Mol. Biol. Rev.* **66**, 1–20
- 11 Cournac, L., Guedeney, G., Peltier, G. and Vignais, P.M. (2004) *J. Bacteriol.* **186**, 1737–1746
- 12 Maness, P.C., Smolinski, S., Dillon, A.C., Heben, M.J. and Weaver, P.J. (2002) *Appl. Environ. Microbiol.* **68**, 2633–2636
- 13 Melis, A., Zhang, L., Forestier, M., Ghirardi, M.L. and Seibert, M. (2000) *Plant Physiol.* **122**, 127–135
- 14 Kosourov, S., Seibert, M. and Ghirardi, M.L. (2003) *Plant Cell Physiol.* **44**, 146–155
- 15 Ghirardi, M.L., Zhang, L., Lee, J.W., Flynn, T., Seibert, M., Greenbaum, E. and Melis, A. (2000) *Trends Biotechnol.* **18**, 506–511
- 16 Fedorov, A., Kosourov, S., Seibert, M. and Ghirardi, M.L. (2005) *Appl. Biochem. Biotechnol.*, in the press

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