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N₂ reduction: interfacing the enzyme nitrogenase with electrodes

Vincent Fourmond and Christophe Léger*

Two processes contribute equally to the humanity's colossal needs for fixed forms of nitrogen: the high-temperature, high-pressure Haber–Bosch process, [1] and the biological reduction of N_2 by the enzyme nitrogenase, which is found only in a selected group of microorganisms and played a critical role during the early expansion of microbial life. [2] Nitrogenases accomplish the great feat of activating N_2 under mild conditions, using the energy that is supplied in vivo by the hydrolysis of adenosine 5'-triphosphate (ATP).

Milton, Minteer and coworkers recently showed that this enzyme can receive from methyl viologen, and ultimately electrodes, the electrons that it uses to reduce N_2 . [3a] In an earlier work, they used cobaltocene to mediate electrons between an electrode and the MoFe protein of nitrogenase, but no N_2 reduction was observed. [3b] This new result opens the way for developing various electrochemical devices and calls for further investigations of the kinetics of the reaction of the enzyme with the versatile electron donor methyl viologen.

Nitrogenase (here we focus on the Mo-containing version) catalyses the reduction of N_2 according to a reaction that is often written as:

$$N_2 + 8e^- + 16MgATP + 8H^+ \Rightarrow 2NH_3 + H_2 + 16MgADP + 16P_i$$
 (1)

For each molecule of N_2 that is reduced (a six-electron reaction), the enzyme needs at least 8 electrons, hydrolyzes at least 16 molecules of ATP to ADP and inorganic phosphate (Pi) and produces at least one molecule of H_2 . In the absence of N_2 , the enzyme catalyzes hydrogen evolution according to

$$2H^+ + 2e^- + 4MgATP \Rightarrow H_2 + 4MgADP + 4P_i$$
 (2)

Nitrogenase is composed of two components termed Fe and MoFe proteins (fig 1), which bind and slowly dissociate during the catalytic cycle. [4a] The Fe protein contains the binding sites for ATP and a single [4Fe4S] redox cluster. The MoFe protein houses the [Fe7S9MoC-homocitrate] cofactor or "FeMo-co", which is the active site for N2 reduction, and the [Fe8S7] P-cluster, which mediates electron transfer from the Fe protein to FeMo-co. [4b]

The transformations among catalytic intermediates of the FeMoco, denoted E_0 to E_8 (where the subscript indicates the number of electrons and protons delivered to the FeMo-co), are shown in scheme 1. Reversible N_2 binding occurs at stage 3 or 4, after the accumulation of 3 or 4 electrons and protons. The mechanism proposed by Hoffman, Dean and Seefeldt supports the obligatory release of two (expensive) reducing equivalents upon release of one H_2 molecule when N_2 binds. $^{[5]}$ In this mechanism, the FeMo-cofactor cycles through only two formal redox levels of the metal-ion core as the first 4 electrons and protons are successively transferred to the FeMo-co, because the reducing equivalents are formally stored as pairs of metal-bound protons and hydrides, rather than on the metal ions. A similar

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mechanism operates in hydrogenases, the enzymes that reversibly oxidize and produce H_2 . At stage 4 in scheme 1, reductive elimination of H_2 produces a highly activated, doubly reduced site, where N_2 binds and is subsequently hydrogenated. Other chemists have favored alternative mechanisms that use only 6 electrons. [6]

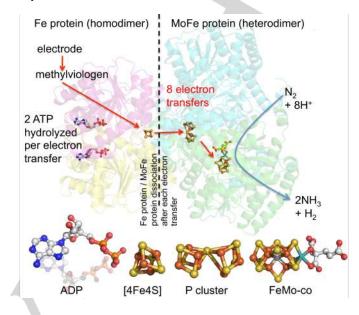
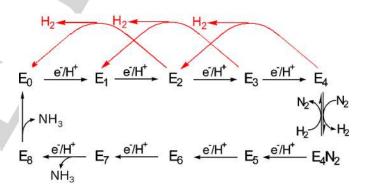


Figure 1. The structure, reaction and cofactors of the Fe protein / MoFe protein nitrogenase complex (pdb accession codes IN2C and 3U7Q).^[4]



Scheme 1. Kinetic scheme depicting the sequential delivery of electrons/protons to the FeMo-co, the futile H_2 -production cycles (in red), the reversible binding of N_2 (consistent with the hypothesis of obligatory reductive elimination of H_2 discussed in ref [5]), and the release of product NH_3 . Adapted from figs 3 and 19 and scheme 1 in ref [5].

Hydrolysis of one mol of ATP provides about 60 kJ (the exact value depends on how far from equilibrium is the steady-state ATP/ADP ratio maintained by the bioenergetics metabolism;^[7] it is not, unlike sometimes stated, the standard free energy of the reaction of ATP hydrolysis). Considering the stoichiometry in ea 1. this implies that electrons involved in a particular electron transfer step in the nitrogenase catalytic cycle may be 'energized" by more than 1.2 V by using 2 molecules of ATP (1.2V=2×60kJ/F). Which step(s) exactly benefit(s) from this additional energy is an active subject of research. Both ATP binding to the Fe protein and Fe / MoFe proteins association decrease the potential of the [4Fe4S] cluster, [8] which implies that binding is much stronger when the Fe protein is oxidized; therefore, ATP hydrolysis may just be the price to pay for forcing the dissociation of the complex. In experiments where low potential electrons are photo-generated and transferred to the

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MoFe protein to drive catalysis (resulting in either H_2 evolution or N_2 reduction, depending on the nature of the photosensitizer), there is no need for Fe protein or ATP. [9]

In most kinetic studies, the electrons required to reduce N₂ are received by nitrogenase from the slow and complex electron donor dithionite (DT), or, less frequently, from the physiological reductants flavodoxin or ferredoxin. Here, Milton, Minteer and colleagues make a significant step forward by providing the first evidence that the non physiological electron donor methyl viologen (MV), which is often used in kinetic and electrochemical studies of other redox enzymes, can sustain N₂ reduction by nitrogenase at a significant rate. [3a] This makes it possible to interface the enzyme with electrodes and to use it in a variety of bioelectrochemical devices, the feasibility of some of which is illustrated in the paper. Indeed, the authors demonstrated that the nitrogenase cathode can be coupled to various anodes in a fuel cell configuration, including one anode that produces electrons by the catalytic oxidation of H₂ by the enzyme hydrogenase. A fuel cell was constructed that reaches an OCP of 250 mV and delivers 20 μ A/cm² at a maximal power density of 1.4 μ W/cm².

In the experiments highlighted here, ADP is continuously rephosporylated by the sacrificial consumption of creatine phosphate. The supply of ATP will remain limiting for any practical use of nitrogenase, unless it becomes possible to couple N_2 bioelectrochemical reduction to a system that regenerates ATP using the energy from e.g. light.^[10]

Methyl viologen is used here for the first time as an electron donor to nitrogenase, and this electron transfer will probably be investigated in follow up studies, knowing that changing the redox partner of nitrogenase may change the rate limiting step in catalysis [11] and the distribution of products. [9] The authors observe that the rate of N₂ reduction by MV is 75% of that when DT is used as reductant, and that the efficiency of the reduction (in terms of charged passed to NH₃ produced) is suboptimal, in the range 25% to 60%. [3a] This suggests that the enzyme wastes electrons (and ATP) to produce more H2 than just one per N2 (eq 1). According to scheme 1, H2 can be released from a number of catalytic intermediates, as the FeMo-co spontaneously returns to a two-electron more oxidized state, closing a futile H₂ production cycle. [5] If the rate of reduction of the nitrogenase complex by MV is low, it may be that the competition between H₂ evolution and ATP-dependent electron accumulation leads to the continuous reoxidation of the FeMo-co (upon H2 release) before N₂ binds (scheme 1), and decreases the overall efficiency of N₂ reduction. This is reminiscent of the earlier observation that not all photochemical electron donors can sustain N₂ reduction. [9] A final complication with MV may be the possibility of ATPdependent reverse electron transfer from the Fe protein in the complex back to oxidized MV. This may consume ATP and slow down the reduction of the MoFe protein, further increasing the probability of wasting reductive equivalents by H₂ release. Electrons may also be lost in the non-enzymatic evolution of H₂, but probably not on the carbon electrodes used by the authors.

As a final perspective, we note that a central challenge regarding the biological reduction of N_2 is that both of the Fe and MoFe proteins that compose the nitrogenase enzyme are irreversibly damaged by oxygen. The former is particularly sensitive. 12al Nitrogenase-based photobiological production of H_2 is possible in aerobic photosynthetic microorganisms such as cyanobacteria on condition that H_2 production is temporally or spatially separated from photosynthesis, and oxygen sensitivity of nitrogenase has been recognized as an obstacle for bioengineering nitrogen fixation into cereal crops. Strict anoxia will also be a requirement for using the enzyme in electrochemical cells, but the viologen mediator may come to the rescue. Indeed, there have been recent developments showing that oxygen sensitive enzymes can be protected from

oxygen upon incorporation in a hydrated film of polymer bearing viologen moieties. The latter mediate electron transfer between the enzyme and the electrode, and also prevent oxygen from penetrating the film by reducing it to water. [12b] So far the protection has proved effective only under conditions of oxidative catalysis, but redox hydrogels can also promote reductive catalysis. The objective of using nitrogenase will further motivate the search of alternative protection mechanisms that operate under reductive conditions.

Keywords: electrocatalysis • nitrogenase • N2 reduction

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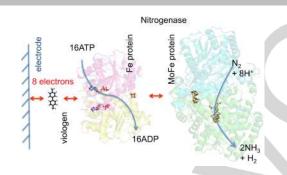
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Entry for the Table of Contents (Please choose one layout)

Layout 1:

HIGHLIGHT

Milton, Minteer and coworkers report the first evidence for the bioelectrochemical reduction of N_2 to ammonia by nitrogenase. This complex enzyme could be "wired" to an electrode by using the soluble mediator methyl viologen; this very simple approach makes it possible to develop a variety of biotechnological devices.



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