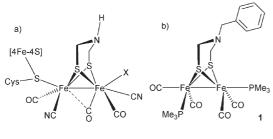
Iron hydrogenase active site mimic holding a proton and a hydride

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Received (in Cambridge, UK) 11th October 2005, Accepted 25th November 2005 First published as an Advance Article on the web 20th December 2005 DOI: 10.1039/b514280f

The first model of the iron hydrogenase active site has been prepared which concomitantly carries a proton and a hydride; the title species was characterized by IR and NMR spectroscopy and is reduced at more positive potential than any other mimic of this kind.

Iron hydrogenases (Fe-H₂ase) are enzymes which catalyze the reversible reaction of protons and electrons to molecular hydrogen.¹ Whereas the crystal structure elucidation of two Fe-H₂ases provided a detailed structural picture of their active sites (H-cluster, Chart 1a),^{2,3} small synthetic model complexes are valuable aids to compare and to understand the spectroscopic signatures of the enzyme.^{4–7} Furthermore, functional studies on the model complexes have contributed to the understanding of the enzyme's mechanism.^{8–10}





The active site is comprised of two unusual low-valent diiron nuclei which are coordinated by a cysteine-linked [Fe₄S₄] cluster, carbon monoxide and cyanide ligands and by a non-proteic azadithiolate (adt = S-CH₂-NH-CH₂-S) bridging the two iron centers.^{11,12} As the reduction of protons or the oxidation of hydrogen follows an ionic mechanism, the H-cluster inevitably has to pass through a state where it carries a proton and a hydride. Using computational techniques, the central nitrogen atom of the dithiolate bridge was identified as a potential basic site,¹³ whereas the cyanide ligands increase the electron density at the iron centers and facilitate the formation of a hydride at this position. Although the hydride has to reside at least transiently in a terminal position to engage in an intramolecular reaction with the proton,^{14,15} a bridging hydride cannot be excluded as an intermediate state of the Fe-H₂ase. In fact, a bridging hydride has recently been identified in

^aDepartments of Organic and Structural Chemistry, Arrhenius Laboratory, Stockholm University, 10691, Stockholm, Sweden ^bDepartment of Chemistry, Organic Chemistry, BMC, Uppsala University, P.O. Box 599, 75124, Uppsala, Sweden. E-mail: sascha.ott@fki.uu.se; Fax: +46 (0)18 4717340 ^cDepartment of Physical Chemistry, BMC, Uppsala University, P.O. Box 579, 751 23, Uppsala, Sweden. E-mail: reiner.lomoth@fki.uu.se; Fax: +46 (0)18 4713654 the functionally related Ni-Fe H_2 ase.¹⁶ We therefore set out to synthesize a novel Fe- H_2 ase active site mimic which contains electron donating ligands at the diiron core as well as an adt linker (Chart 1b), anticipating that an unprecedented species carrying a hydride at the diiron core and a proton at the adt nitrogen may be realized. Such a structure can be regarded as a snapshot of the H-cluster just before the formation of the H–H bond.

17-19 Hence, a solution of readily synthesized $[Fe_2(\mu-adt)(CO)_6]$ 2 in hexane was treated with trimethylphosphine to afford complex 1 in excellent yield.²⁰ Complex 1 is rather air sensitive in solution and attempts to purify the complex by column chromatography resulted in complete degradation of the product. However, addition of a little toluene and removal of the solvents and of excess trimethylphosphine at high vacuum spontaneously produced crystalline material suitable for X-ray analysis.† Singlecrystal X-ray diffraction analysis of 1 (Fig. 1) reveals the usual edge-bridged bi-square-pyramidal geometry around the iron centers which are at a distance of 2.55 Å.²¹ The two trimethylphosphine ligands prefer an unsymmetric geometry around the iron centers in the solid state with one in an apical and the other in a basal position. This conformation is similar to that preferred by an ethyldithiolate-bridged diiron complex where a minimum of steric hindrance at the bridge has been suggested to facilitate this arrangement.20

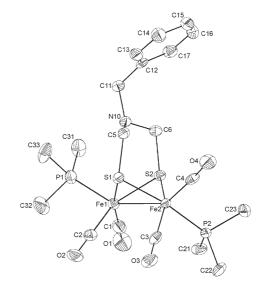


Fig. 1 ORTEP view (ellipsoids at 50% probability level) of $[Fe_2(\mu-SCH_2N(CH_2Ph)CH_2S)(CO)_4(PMe_3)_2]$ (1). Selected bond lengths (Å): Fe1–Fe2 2.5461(12), Fe1–S1 2.2663(18), Fe1–S2 2.2546(16), Fe2–S1 2.2560(17), Fe2–S2 2.2465(16), Fe1–P1 2.2174(18), Fe2–P2 2.2255(18), Fe1–C1 1.724(9), Fe1–C2 1.766(6), Fe2–C3 1.767(9), Fe2–C4 1.774(7).

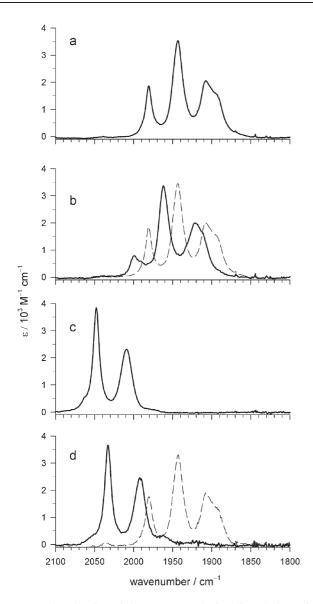
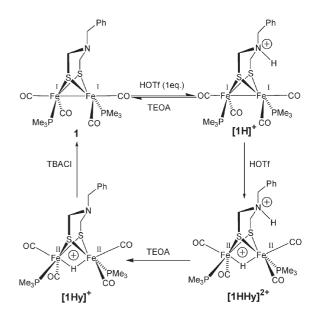


Fig. 2 Carbonyl region of the IR spectra obtained for solutions of 1 (3 mM) in CH₃CN. Protonated states were generated under the conditions indicated in parentheses. a) 1. b) $[1H]^+$ (3 mM HOTf, —); Spectrum after deprotonation with triethanolamine (TEOA, ---). c) $[1HHy]^{2+}$ (0.2 M HOTf) d) $[1Hy]^+$ (0.2 M HOTf followed by TEOA, —); Spectrum after deprotonation with tetrabutylammonium chloride (TBACI, ---).

Addition of one equivalent of triflic acid to a solution of 1 in CH₃CN results in the formation of [1H]⁺, as evidenced by the IR spectra (Fig. 2a, b) that show a shift of $\tilde{v}_{CO} = 16 \text{ cm}^{-1}$ towards higher energy. This shift is typical for a protonation of the adt nitrogen.^{8,17} Addition of triethanolamine (TEOA) reverses the adt protonation and complex 1 is quantitatively recovered. Upon addition of an excess of triflic acid, [1H]⁺ is transformed into a product characterized by the IR spectrum shown in Fig. 2c. This spectrum can be assigned to the doubly protonated species [Fe₂(µ-Hadt)(µ-H)(CO)₄(PMe₃)₂]²⁺, [1HHy]²⁺, with an additional average shift of $\tilde{v}_{CO} \approx 80 \text{ cm}^{-1}$, reflecting the substantial decrease of electron density at the diiron site as formally expressed by the Fe^{II}–Fe^{II} valence state of the hydride complex.^{10,22} [1HHy]²⁺ is the first Fe-H₂ase active site mimic which carries a proton at the adt nitrogen as well as a hydride at the diiron core. Initial attempts

to deprotonate [1HHy]²⁺ to recover 1 with an excess of triethanolamine failed and a new spectrum emerged which in shape resembled that of [1HHy]²⁺, however shifted by $\tilde{v}_{CO} \approx$ 16 cm^{-1} towards lower frequencies. It thus seems that the adt nitrogen is selectively deprotonated, leaving the hydride portion intact and the species formed is identified as the hydride complex $[Fe_2(\mu-adt)(\mu-H)(CO)_4(PMe_3)_2]^+$, $[1Hy]^+$. The formation of $[1Hy]^+$ by deprotonation of [1HHy]²⁺ relies on the inertness of the hydride towards nitrogen bases. The sluggish deprotonation behavior with nitrogen bases has been reported for a related hydridic diiron complex,¹⁰ but could be overcome by the addition of cyanide or chloride.²³ In the case of **[1Hy]**⁺ addition of tetrabutylammonium chloride results in quantitative recovery of 1 (Fig. 2d, dashed line). Complex 1 constitutes a dibasic species which, as such, should give rise to three different protonation states. However, owing to the remarkable difference in deprotonation kinetics between the adt nitrogen and the Fe-Fe bond, complex 1 can selectively be prepared in *four* protonation states, 1, $[1H]^+$, $[1Hy]^+$, and $[1HHy]^{2+}$ (Scheme 1). 24

NMR characterization of all four states shows that protonation of the adt nitrogen renders the two Fe(CO)₂P(CH₃)₃ subunits in $[1H]^+$ and $[1HHy]^{2+}$ non-equivalent as the ring inversion of the sixmembered metalloazaheterocycle is restricted. Hence, the trimethylphosphine protons which give rise to one doublet in the ¹H NMR spectra of 1 and [1Hy]⁺ due to their coupling to the phosphorus center, are split into two doublets at $\delta = 1.56, 1.54$ ppm for $[1H]^+$ and at $\delta = 1.57$, 1.63 ppm for $[1HHy]^{2+}$. Similarly, the ³¹P NMR spectra of the adt-protonated [1H]⁺ and [1HHy]²⁺ feature two phosphorus signals at $\delta = 31.1$, 19.8 ppm for $[1H]^+$ and at $\delta =$ 25.3, 21.3 ppm for [1HHy]²⁺ whereas only one signal is observed in the spectra of 1 and $[1Hy]^+$ (Fig. 3). The hydride is visible in the ¹H NMR spectra of $[1Hy]^+$ and $[1HHy]^{2+}$ as a triplet ($J_{H-P} = 22$ Hz) at $\delta = -15.6$ ppm and as a doublet of doublets ($J_{H-P} = 21, 23$ Hz) at similar chemical shift, respectively. Since the same coupling constant can be observed in the ³¹P NMR of the respective



Scheme 1 Protonation reactions of 1 in acetonitrile solution with triflic acid (HOTf). Deprotonation of the adt nitrogen with triethanolamine (TEOA) and of the hydride with tetrabutylammonium chloride (TBACl).

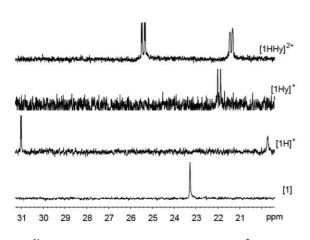


Fig. 3 31 P NMR spectra of 1, [1H]⁺, [1Hy]⁺ and [1HHy]²⁺ in CD₃CN at 25 °C. Prepared from 1 (6 mM) under conditions similar to those used in the IR experiments.

species, it is clear that the hydride in $[1Hy]^+$ and $[1HHy]^{2+}$ resides in a bridging position²⁵ and that the complexes are stable even under these drastic acidic conditions. In contrast to the solid state structure, the phosphine ligands prefer a symmetrical basal position relative to the iron centers in $[1Hy]^+$ and $[1HHy]^{2+}$ as evidenced by the relatively large coupling constants of $J_{H-P} =$ 22 Hz.²⁰

Voltammetric investigation of complex 1 reveals an irreversible reduction at *ca.* -2.2 V *vs.* Fc^{+/0} and the potential for the first reduction is shifted to -1.0 V for [1HHy]²⁺. This dramatic shift of 1.2 V is a result of the fact that [1HHy]²⁺ concurrently carries a proton and a hydride.²⁶

In summary, we have synthesized the first biomimetic model of the Fe-H₂ase active site which can be protonated on either the Fe– Fe bond or the adt nitrogen as well as on both sites simultaneously. All four protonation states are well-defined and have been characterized by IR and NMR spectroscopy. Double protonated $[1HHy]^{2+}$ is reduced at -1.0 V vs. Fc^{+/0}, a potential considerably more positive than that of any other Fe-H₂ase mimic ever reported. Reduction at such mild potential is a result of the two preceding protonations, one of which involves the adt nitrogen.²⁶ From a structural point of view, $[1HHy]^{2+}$ resembles an intermediate closer to the hydrogen formation event than other model complexes which only carry either a hydride or a proton.^{8,10} With regard to the enzyme, the question arises as to whether the catalytic cycle involves the formation of a double protonated species at the Fe^I–Fe^I level similar to our model system.

We gratefully acknowledge financial support by the Swedish Energy Agency, the Swedish Research Council and NEST, SOLAR-H (EU contr. nr. 516510).

Notes and references

† [Fe₂(μ-SCH₂N(CH₂Ph)CH₂S)(CO)₄(PMe₃)₂] (1). Trimethylphosphine (204 mg, 2.68 mmol) was added to a solution of complex 2 (160 mg, 0.335 mmol) in 5 ml hexane under nitrogen atmosphere. After 3 h of stirring, the solvent and unreacted trimethylphosphine were removed *in vacuo*. The resulting red-brown solid was re-dissolved in toluene and the solution was filtered through a plug of Celite. The solvent was removed, and the deep red solid was washed with cold hexane (174 mg, 91%). Single crystals suitable for X-ray analysis were obtained from toluene/hexane

solution upon rapid concentration *in vacuo*. Anal. calculated for C₁₉H₂₉Fe₂NO₄P₂S₂: C, 39.81; H, 5.10; N, 2.44. Found: C, 39.83; H, 5.27; N, 2.38%. ¹H NMR (400 MHz, CDCl₃): δ = 7.30–7.26 (m, 3H, Ar*H*), 7.18 (d, *J* = 6.8 Hz, 2H, Ar*H*), 3.55 (s, 2H, NCH₂Ph), 3.08 (s, 4H, SCH₂N), 1.50 (d, *J* = 9.2 Hz, 18H, *PMe*₃). ¹³C NMR (100.6 MHz, THF-d₈): δ = 217.5, 217.3, 137.9, 130.3, 129.1, 128.3, 63.8, 54.2, 20.8, 20.4 (2C). ³¹P NMR (161.9 MHz, CD₃CN): δ = 23.3. IR (CH₃CN, cm⁻¹): *v*_{CO} = 1980, 1943, 1907, 1892 (sh). Crystallographic data of 1: Monoclinic, *a* = 10.7622(14) Å, *b* = 19.773(3) Å, *c* = 11.7983(15) Å, β = 90.390(15)°, vol = 2510.7(6) Å³, *T* = 293(2) K, *P*2₁/*c*, *Z* = 4, μ = 1.474 mm⁻¹, *N*_{measured} = 16225, *N*_{unique, all data = 4834, *N*_{unique, (*I* ≥ 2 σ (*I*)). CCDC 283269. For crystallographic data in CIF or other electronic format see DOI: 10.1039/b514280f}}

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- 24 Since no interconversion between **[1H]**⁺ and **[1Hy]**⁺ could be observed on the timescale of hours, it cannot readily be determined which monoprotonated species is the thermodynamically stable form.
- 25 In addition, two-dimensional ¹H-³¹P correlation spectroscopy (HMQC) of [1HHy]²⁺ revealed the coupling of the hydride to both phosphorus centers.
- 26 Reference 10 describes a related diiron-hydride complex lacking the adt nitrogen which can be protonated at a cyanide ligand instead. This species, however, is reduced at -1 V vs. Ag/AgCl, corresponding to less than -1.4 V vs. Fc^{+/0}.