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[FeFe]-Hydrogenases: maturation and reactivity of enzymatic systems and overview of biomimetic models

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While hydrogen plays an ever-increasing role in modern society, nature has utilized hydrogen since a very long time as an energy carrier and storage molecule. Among the enzymatic systems that metabolise hydrogen, [FeFe]-hydrogenases are one of the most powerful systems to perform this conversion. In this light, we will herein present an overview on developments in [FeFe]-hydrogenase research with a strong focus on synthetic mimics and their application within the native enzymatic environment. This review spans from the biological assembly of the natural enzyme and the highly controversial discussed mechanism for the hydrogen generation to the synthesis of multiple mimic platforms as well as their electrochemical behaviour.

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I Introduction

Hydrogen plays an ever-increasing role in our modern society and is anticipated to serve as a green and sustainable energy carrier as well as storage in future. While already produced on a large scale, current production of hydrogen is industrially realized by reforming of fossil fuels. Notably, only a small fraction is currently generated by water splitting. Contrary to the industrial generation of hydrogen and the political as well as societal demands to use more hydrogen, nature has almost perfected the handling of this small molecule. In a small number of eukaryotes (green algae) but more importantly in specialised anaerobe microorganisms (bacteria as well as archaea) hydrogen can act as the primary energy carrier. Among the enzymes that allow for hydrogen transformation, [FeFe]-hydrogenases are the most competent. The active site of these enzymes commonly comprises a hexanuclear Fe-cofactor, consisting of a [4Fe-4S]- and a [2Fe-2S]-cluster. The most commonly investigated [FeFe]-hydrogenases are from *Clostridium pasteurianum* (*Cp*), *Desulfovibrio desulfuricans* (*Dd*) and *Chlamydomonas reinhardtii* (*Cr*). Notably, these enzymes can be regarded as "fuel and electrolysis cells" and allow for the reversible interchange of protons to hydrogen with a turnover



Julian T. Kleinhaus

Julian T. Kleinhaus received his MSc in Chemistry in 2019 from the Ruhr-University Bochum. Under the supervision of Prof. Dr U.-P. Apfel, he focused on establishing synthesis routes for new [FeFe]-hydrogenase mimics. During his PhD studies in the Apfel group, he investigates novel electrocatalytic applications for biomimetic and bioinspired catalysts.



Florian Wittkamp

Florian Wittkamp joined the group of Ulf-Peter Apfel in 2014 for his undergraduate studies and changed to the field of [FeFe]-hydrogenases in 2015. He got his PhD from the Ruhr-University Bochum in 2020. In his thesis he discussed the chemistry of the [FeFe]-hydrogenases' active site und is currently still interested in designing new biomimetic models as possible replacement for the native H-cluster.

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frequency of up to over 10 000 µmol (H₂) min mg⁻¹ (enzyme) under mild conditions (-0.413 V vs. standard hydrogen electrode, pH 7).3 It is thus very plausible that this enzyme system received increased attention and scientists all over the world have taken the active centres of hydrogenases as a template to design mimetics which display a comparable activity for the hydrogen evolution reaction. We will herein attempt to provide a complete picture on developments in this field in the last two decades since the structure of the active site of [FeFe]-hydrogenases was revealed. Starting out from recent advances in artificial maturation of fully functional enzymes, we will continue in describing the reactivity of the natural H-cluster. Going further, we will present synthetic pathways towards [FeFe]-hydrogenase mimics, show the plentiful chemical alterations and their impact on the structure as well as their electrochemical properties. As a subject of growing interest, our discussion will furthermore shed light on the possibility of photocatalytic hydrogen evolution using hydrogenase mimics.



Shanika Yadav

Shanika Yadav received her BSc (2015) and MSc (2017) degree in Chemistry from University of Pune, India. In 2019, she received the "DAAD-Graduate School Scholarship" and joined the group of Prof. Ulf-Peter Apfel as a PhD student. Her current research focuses on synthesis of active site mimics of [FeFe]-hydrogenases and investigation of oxygen sensitivity of the enzyme.

Part A: the chemistry of [FeFe]-hydrogenases

II Maturation of natural and semi-artificial [FeFe]-hydrogenases

2.1 Native in vivo maturation

The biosynthesis and assembly of the complete active site of [FeFe]-hydrogenases, called H-cluster, requires the interaction of several maturase proteins HydG, HydE and HydF (see Fig. 1 for an overview). Its whole structure is rather uncommon in biology and consist of two individual iron-sulphur clusters, which are linked by a cysteine sidechain. The first is a [4Fe-4S]cluster, herein abbreviated with [4Fe]_H, which is responsible for electron delivery and serves as electron reservoir by switching between an oxidized and reduced state during the catalytic cycle. The second iron-sulphur cluster is a [2Fe-2S]-cluster. This subsite will be abbreviated [2Fe]H and represents the actual active centre, being the site of catalytic turnover. Depending on the position relative to [4Fe]_H, the single iron atoms are termed proximal iron (Fe_p) and distal iron (Fe_d), respectively. Fe_p is octahedrally coordinated by the cysteine's thiolate, a terminal CO and CN⁻ ligand each, two bridging sulphides that form the [2Fe-2S]-cluster and an additional μ-CO ligand, which is in a bridging binding mode between both iron atoms. Fe_d shows identical ligands but lacks the thiolate of the cysteine therefore showing a square-pyramidal coordination sphere. At the open binding site substrates, e.g. H⁺ in the H_{hvd} state and H₂ in the Hox-H2 state, as well as inhibiting diatomic gases like CO (Hox-CO) and O_2 (H_{ox} - O_2) may bind. [2Fe]_H is further coordinated by a secondary amine via the bridging sulphides, why this ligand is mostly called adt (azadithiolate, precisely: bis(sulfidomethyl)amine). The whole [2Fe]_H cluster, bearing the adt ligand, is therefore casually called ADT. For a complete graphical representation of the H-cluster see Fig. 1, red box.



Daniel Siegmund

Daniel Siegmund obtained his PhD at the University of Bochum for his work on the development of novel organometallic antibiotics with Prof. N. Metzler-Nolte. In 2018 he joined the Fraunhofer Institute for Environmental, Safety and Energy Technology (UMSICHT) in Oberhausen where he is currently a group leader for electrocatalysis in the department of energy. His research interests focus on the development of precious metal-free electrocatalysts for hydrogen

evolution and CO_2 -reduction as well as the establishment of sustainable electrocatalytic synthesis processes for organic commodity and fine-chemicals.



Ulf-Peter Apfel

Ulf received his PhD from the Friedrich-Schiller University Jena. After a postdoctoral stay at MIT (2011/2012), he started his independent career at the Ruhr University Bochum funded by the "Fonds der Chemischen Industrie" and the DFG as an Emmy Noether group leader. He holds a professorship at the Ruhr University Bochum since 2019 and is leading the department Electrosynthesis at Fraunhofer UMSICHT. His research interests

are in the field of technical electrochemistry with a special emphasis on the electrochemical reduction of CO_2 and protons and catalyst design.

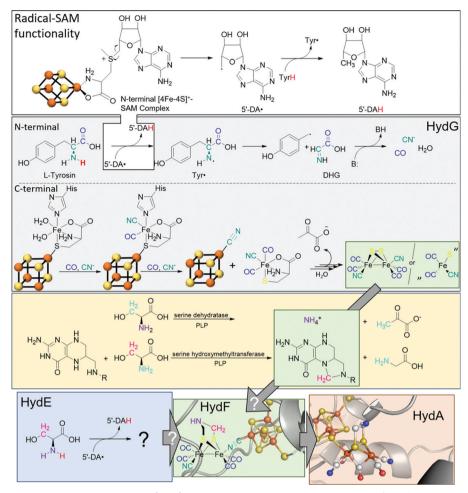


Fig. 1 Overview of biological maturase machinery of [FeFe]-hydrogenases. White: N-terminal radical SAM functionality in HydG to initiate the degradation of tyrosine. Grey: reactions performed by HydG yielding a Fe₂S₂(CO)₄(CN)₂ core or a FeS(CO)₂CN synthon. Yellow: putative PLP dependent conversion of serine by serine dehydratase and serine hydroxymethyltransferase to ammonia and 5,10-methylene tetrahydrofolate. Blue: putative reactivity of HydE. Green: possible substrates of HydF and assembled [2Fe]_H-precursor on HydF. Red: completely maturated HydA. PDB entries: HydA: 4XDC, apo-HydF: 3QQ5.2.

In 2010, Mulder and coworkers showed that without HydG, HydE and HydF, the H-cluster contains only the [4Fe]_H-cluster (apo-HydA) leading to a change of the quaternary structure of the enzyme. This change results in a positively charged channel leading directly to the active centre, which is used to incorporate [2Fe]_H and is closed in the presence of the complete H-cluster. HydG is part of the radical S-adenosyl-L-methionine (SAM) enzyme superfamily and accordingly has the usual reactivity:⁵ SAM chelates an iron atom of a [4Fe-4S]⁺-cluster via the carboxy and amine function of methionine. The remaining iron atoms are bound to the protein environment by cysteine residues. The Fe-S cluster induces a reductive cleavage of the bound SAM by an inner-sphere electron transfer, resulting in a highly reactive 5'-deoxyadenosyl radical (5'-DA*) and methionine remains on the now oxidised [4Fe-4S]²⁺-cluster. 5'-DA* abstracts a hydrogen radical of an enzyme specific substrate, forming 5'-DAH and enabling various downstream reactions. In the case of HydG, 5'-DA* abstracts one of the hydrogens of a tyrosine amine group.6 The resulting tyrosine radical (Tyr*) undergoes a homolytic bond cleavage between

 C_{α} and C_{β} and decomposes into a 4-hydroxybenzyl radical (4-HOB*) and dehydroglycine (DHG). DHG can subsequently undergo a base-assisted decomposition to form CO as well as CN and thus serves as a potential source of the biologically unusual ligands for [2Fe]_H.^{7–9}

Besides the N-terminal radical SAM functionality, HydG has another Fe-S cluster in C-terminal position: An auxiliary [5Fe-5S]-cluster, which was investigated by EPR spectroscopy and X-ray crystallography. 10 The g-values of 9.5, 4.7, 4.1, and 3.7, which are unusual for biological Fe-S clusters, represent an S = 5/2 spin. This unusual observation is caused by a ferromagnetic coupling between a $[4Fe-4S]^+$ -cluster (S = 1/2) and an additional high-spin Fe^{2+} (S = 2). Both are connected by a bridging sulphide of a nonproteinic cysteine (Cys). The additional iron is further coordinated by a histidine (His) residue and two water molecules.11

The CO and CN⁻ ligands obtained by the radical SAM functionality first substitute the aqua ligands of the additional iron resulting in a [4Fe-4S][(Cys)Fe(CO)(CN)(His)] complex. Subsequently, histidine can also be exchanged by a further CO

$$\begin{array}{c} \text{OC} \quad \text{S-S} \quad \text{CO} \\ \text{OC} \quad \text{Fe} \quad \text{ICO} \\ \text{OC} \quad \text{CO} \end{array} + \text{LiB}(\text{Et})_3\text{H} + \text{TFA} + \text{HO} \begin{bmatrix} H \\ -C \\ -O \end{bmatrix}_{\text{H}} + (\text{NH}_4)_2\text{CO}_3 \\ \text{B-100} \end{array} + (\text{NH}_4)_2\text{CO}_3 \\ \begin{array}{c} \text{THF} \quad \text{OC} \quad \text{S-S} \quad \text{CO} \\ \text{OC} \quad \text{NEt}_4\text{CN} \\ \text{OC} \quad \text{RT} \end{array} + \begin{array}{c} \text{OC} \quad \text{S-S} \quad \text{CO} \\ \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{OC} \quad \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text{3} \end{array} = \begin{array}{c} \text{OC} \quad \text{NET}_4\text{CN} \\ \text{NC} \quad \text$$

Fig. 2 Synthesis of Fe₂(adt)(CO)₆ (2) and Fe₂(adt)(CO)₄(CN)₂ (3) presented by Li and Rauchfuss in 2002.

with the remaining second cyanide binding to the [4Fe-4S]⁺cluster and liberating the [(Cys)Fe(CO)₂(CN)]⁻ complex which serves as a synthon for [2Fe]_H. This cyanide-induced release mechanism explains the 4:2 CO:CN ratio of the putative [2Fe]_H-precursor. Four tyrosine molecules are required to assemble the putative [2Fe]_H-precursor and converted into four CO and four CN⁻, two of which are cyanides responsible for the release of the synthon [(Cys)Fe(CO)₂(CN)]⁻.^{6,11}

The role of the synthon has recently been further investigated by Britt and Rauchfuss. 12 Therein, a biomimetic synthon [FeI₂(CO)₃CN]⁻ together with cysteine was added to a HvdA maturation solution consisting of apo-HydA, HydE and HydF only. In the absence of HydG, this mixture was able to completely activate HydA. Furthermore, with ¹³C and ¹⁵N labels and by using selenocysteine, cysteine was unequivocally shown to be the source of the bridging sulphides within the [2Fe-2S]cluster but does not provide the NH(CH₂)₂ bridge.¹²

Like HydG, HydE is an enzyme of the radical SAM family with two [4Fe-4S]-clusters, as demonstrated by EPR spectroscopy, ¹³ or one [4Fe-4S]- and one [2Fe-2S]-cluster, according to X-ray studies. 14 Notably, due to the one-week duration of crystal growth in the X-ray study conducted by Fontecilla-Camps, a degradation of the C-terminal [4Fe-4S]-cluster may have occurred and potentially results in the observed [2Fe-2S]cluster. However, the [4Fe-4S]-cluster can be removed by mutation without loss of maturase-specific activity and is therefore considered functionally irrelevant for in vivo maturation.¹⁴

Notably, the role of HydE in the HydA maturase machinery has not yet been finally clarified. However, due to its C-S bond formation activity15 it is assumed that the enzyme is involved in the biosynthesis of the adt ligand.¹³ Serine has recently been identified as the source of the NH(CH₂)₂ moiety. More specifically, 13C and 15N labels in combination with EPR, HYSCORE- and ENDOR spectroscopy showed that the NH₂ and β-CH₂ groups are incorporated into the adt bridge. ¹⁶ It could, however, not be clarified if a further substrate is involved, since possibly only one of the CH2 groups is derived from serine, or whether two serine molecules are needed for the complete construction of the bridge. HydE might therefore use serine to assemble the adt bridge. Here, the serine dehydratase and serine hydroxymethyltransferase were also considered as potentially involved enzymes that convert serine to pyruvate and NH₄⁺ and subsequently with tetrahydrofolate (H₄folate) to glycine and 5,10-methylene-H4folate, respectively. Glycine and pyruvate were excluded as possible building blocks of the H-cluster. 12,16 However, NH4+ and 5,10-methylene-H4folate, a biological methyl group donor, came into consideration as potential intermediates. With NH₄⁺ and 5,10-methylene-H₄folate as reagents, the adt moiety could potentially be introduced into the

precursor, $[Fe_2S_2(CO)_4(CN)_2]^{2-}$, comparable to the artificial establishment of the [2Fe]H-cluster by Li and Rauchfuss in 2002 (Fig. 2).17

Furthermore, apo-HydA is activated by HydF, if the latter was expressed together with HydE and HydG. 18,19 As EPR and IR spectroscopy as well as XRD and XAS studies have shown, HydF already contains a [2Fe-2S]-cluster alike [2Fe]_H. Thus, this maturase enzyme is at the end of the activation chain and passes along the almost completed cluster to HydA.20 HydF serves as a scaffold for the iron-sulphur synthon, which was inferred from co-purification of HydE and HydG with HydF and confirmed in vitro through a combination of surface plasmon resonance and co-purification experiments using recombinant proteins of C. acetobutylicum. 18,21-25

2.2 In vitro maturation with artificial H-clusters

In 2013, the groups of Happe, Lubitz and Fontecave showed that chemically synthesised [2Fe-2S]-models with modified bridging dithiols can replace the native [2Fe]_H and were successfully incorporated into the apo-enzyme (Fig. 3).26,27 In contrast to the biological process (see above), maturation was achieved utilising a mixture of apo-CrHydA1 and HydF from T. maritima, which was first incubated with the synthetic cluster $[Fe_2(SCH_2XCH_2S) (CO)_4(CN)_2]^{2-}$ (X = NH (adt), CH₂ (pdt), O (odt)). The successful incorporation into HvdA1 of Chlamydomonas reinhardtii was demonstrated by the specific CO and CN⁻ bands in the enzyme's IR spectrum. Notably, only the variant with X = NH revealed enzyme-specific hydrogen evolution in the presence of methyl viologen and sodium dithionite at 37 °C.27 With this study, two major uncertainties in [FeFe]-hydrogenases were finally resolved. First, X equals NH in the native [FeFe]-hydrogenase. Previous XRD studies were only capable to narrow down the options to adt, pdt and odt due to the identical electron count. 3,28 Thus, early suggestions by Fontecilla and coworkers from 2001 and results obtained from 14N-HYSCORE measurements by the group of Lubitz in 2009 were once and for all proven right.^{29,30} Second, it was shown that although the [FeFe]-hydrogenases were obtained from different organisms, T. maritima and C. reinhardtii, the maturase enzymes are identical in function.

Even more remarkable – the very same groups showed that CrHydA1 can be activated without the use of the maturase HydF and is spectroscopically indistinguishable from naturally produced enzymes.26 Herein, the apo-hydrogenase itself is incubated with the synthetic [2Fe]_H-precursor (Fig. 4).

This completely artificial process further simplifies the complicated biological maturation process and enables the production and isolation of significantly larger amounts of mature enzymes.^{9,26} Especially for spectroscopic applications,

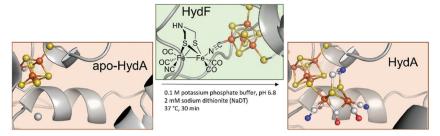


Fig. 3 Overview of HydF depended semi-artificial maturation of HydA. PDB entries: Apo-HydA: 4XDD, HydA: 4XDC, apo-HydF: 3QQ5.

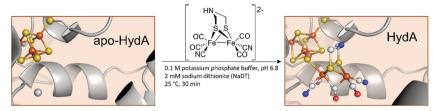


Fig. 4 Schematic representation of the artificial maturation of HydA. PDB entries: Apo-HydA: 4XDD, HydA: 4XDD.

large quantities of high-purity and fully matured enzymes are required. The development of semi-artificial hydrogenases therefore enabled the deciphering of the catalytic mechanism and mode of action of [FeFe]-hydrogenases and thus the possibility to gain information for a new generation of biomimetic [FeFe]-hydrogenase catalysts.

Using this artificial approach, however, only 14 different non-native diiron sites were tested so far for their ability to mature apo-HydA1 from C. reinhardtii.31 Although mimics of the binuclear subcluster with an altered bridging dithiolate ligand (propanedithiolate, oxadithiolate, thiadithiolate, N-methylazadithiolate (adt^{Me}), dimethyl-azadithiolate) and three variants containing only one CN ligand were successfully inserted into the apo-enzyme, the activity of those semi-artificial enzymes was below 1% of the native enzyme. In all cases, the insertion process was followed by IR-spectroscopy and the incorporation of the [2Fe]_H-mimics is visible by significant line narrowing of the CO/CN bands compared to measurements of the sole clustermimics in solution. This narrowing indicates a loss of vibrational freedom of the ligands and also implies interaction with the protein backbone. This effect was likewise observed for apo-HydA1 maturated with adt-loaded HydF or solely adt. 26,27 Upon isolation of the corresponding adt maturated enzyme, multiple signals in the CO region are present indicating a mix of Hox, Hred, Hred, Hsred and minor amounts of Hhvd. Contrary, enzymes maturated with odt, adt Me and sdt show only Hox and odt is present in H_{hvd} directly indicating varied H-cluster reactivities of the respective semi-artificial enzymes.

Using the same approach, artificial active sites that were modified at the metal or chalcogenide positions were introduced to apo-*Cr*HydA1 and apo-*Cp*1. In 2017, Kertess *et al.* presented the semi-artificial enzymes *Cr*HydA1 and *Cp*I that were maturated with the selenium derivative of the native cluster, ADSe (4). Remarkably, these enzymes showed up to native-like activity regarding proton reduction, but less stability

against O_2 -degradation and cannibalisation. Thus, significant amounts of H_{ox} -CO were found directly after maturation, which influenced the activity in F-cluster bearing CpI (see Section 3.9) more than in HydA1.³²

Later in 2018, Sommer *et al.* presented a hybrid-enzyme with a [RuRu]-analogue of ADT, which was investigated due to the interesting reactivity of the noble metal towards hydrogen. Both versions, $[Ru_2(adt)(CO)_4(CN)_2]^{2-}$ (5) and the protonated species bearing a bridging hydride $(5-\mu H)$ were found to be in the same state, namely H_{hyd} , after incorporation into the protein environment.³³ Especially the isomerization of the hydride shows the remarkable influence of the protein backbone on the structure of the diiron subsite, stabilizing the thermodynamically less stable terminal hydride (Fig. 5).

2.3 In vivo maturation with artificial H-clusters

A more recent approach regarding [FeFe]-hydrogenase maturation and, especially, its modification and in vivo investigation was presented by Berggren and coworkers in 2017.34 The researches transferred the results of the in vitro maturation experiments (Section 2.2) to an in vivo system consisting of apo-CrHydA1 that was heterologously overexpressed in E. coli (Fig. 6). In opposition to former experiments, the hydrogenase was not extracted from its host but left inside the living cells. Since E. coli lacks the maturation machinery HydEFG, the hydrogenases inside the cells remain inactive. However, in analogy to the in vitro experiments, addition of 1 mg 3 to the cell cultures (O.D. = 0.2 ± 0.02) resulted in a 35- to 40-fold increase over the background H2 evolution activity, indicating successful activation of the apo-enzymes.34 The described results represent the first intracellular activation of an apoenzyme not including improved cellular import functions and opens up the field for in vivo spectroscopic investigation of [FeFe]-hydrogenases by e.g. EPR and FTIR.35,36 Even more remarkable is the follow-up study as a joint research of the

Fig. 5 Synthetic subsite models with chalcogen or metal exchange that has been implemented into apo-enzymes.

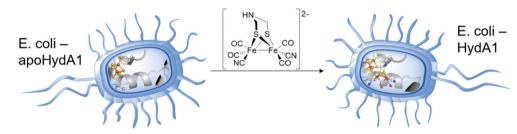


Fig. 6 Graphical representation of the artificial in vivo maturation of bacteria e.g. E. coli.

groups of Berggren and Lindblad, who targeted the modification of the photoautotrophic cyanobacterium Synechocystis PCC 6803. This bacterium harbours a bidirectional [NiFe]-hydrogenase for energy household. However, for biotechnological energy applications, high rates of hydrogen evolution are wanted. Here, the [NiFe]-hydrogenase is by far surpassed by [FeFe]-hydrogenases. Therefore, the hydrogenase CrHydA1 was expressed in a hydrogenase deficient mutant of Synechocystis PCC 6803 (Ahox) as well as in the wild type (WT) organism, containing the native [NiFe]- and the additional [FeFe]-hydrogenase. Upon addition of compound 3, Synechocystis Ahox CrHydA1 and WT-CrHydA1 showed a hydrogen evolution activity of approx. 62 and 48 (nmol O.D.⁻¹ mL⁻¹), respectively, whereas the organisms without 3 showed almost zero activity (Synechocystis ∆hox CrHydA1) and 17 (nmol O.D.⁻¹ mL⁻¹) for WT-CrHydA1. This nice work of bio-engineering underlines that the hydrogen production rates of Synechocystis can be increased by enzyme optimization and opens a new field of artificially improved enzymes for biotechnological hydrogen production.³⁷

III Reactivity of the H-cluster within the enzyme

The hydrogenase activity of green algae was investigated already 80 years ago by Gaffron and coworkers, who found *S. obliquus* to metabolize H_2 upon reduction of CO_2 in photosynthesis. 38,39 Among all algae, *C. reinhardtii* with an *in vitro* activity of 200 nmol H_2 ($\mu g \ Cr \ a \ h)^{-1}$ bears by far the most investigated [FeFe]-hydrogenase HydA1. $^{40-42}$ This hydrogenase contains solely one Fe–S cluster assembly, called H-cluster, and no additional accessory Fe–S cluster for *e.g.* electron transport as in hydrogenases from *D. desulfuricans* and *C. pasteurianum*. $^{43-47}$ Its simplicity thus makes HydA1 most convenient for researching the molecular proceedings of H_2 turnover during catalysis.

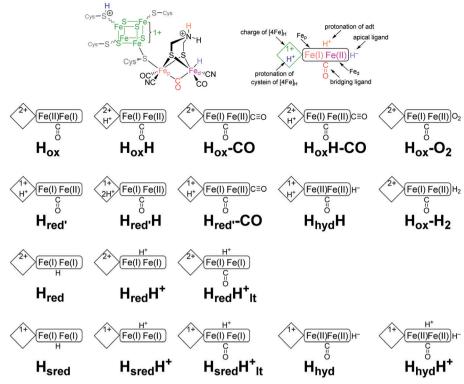
The H-cluster consists of a [4Fe-4S]- and a [2Fe-2S]-cluster, which are linked and electronically coupled *via* a cysteinyl

thiolate. 1,3,28,48,49 The cubic iron cluster, embedded into the protein by three additional cysteine residues, is part of the electron chain and, more importantly, is the midpoint of a proton coupled electron transfer (PCET) at the beginning of the catalytic cycle. 50-52 The diiron subsite conducts the catalytic proton reduction and is the focal point on mimics of the [FeFe]hydrogenases as we will discuss in Section IV. The diiron site consists of a proximal iron that has an octahedral ligand environment and a distal iron in a square pyramidal coordination with an open binding site for substrates (H+/H2) and exogenous ligands such as CO and O2 (Fig. 8). This geometry of the H-cluster in its H_{ox} state is called "rotated state", which refers to the rotation of the distal iron relative to the C_{2v} symmetric Fe₂S₂(CO)₆ core. This special geometry opens a vacant binding site for catalytic turnover and is a unique feature of this active site and was a dominant motif for the design of H-cluster mimics for hydrogen evolution (Fig. 8, Section 5.1).53

We now want to review the reactivity of the H-cluster within the native protein environment. This will include the natural [FeFe]-hydrogenase HydA1 from *C. reinhardtii*, but also hydrogenases from other organisms like *C. pasteurianum*, *D. desulfuricans*, the sensory hydrogenase HydS from *T. maritima* as well as the half-synthetically obtained hydrogenases. Besides different organisms, we will especially highlight the man-made alterations within the [FeFe]-hydrogenases, ranging from different [2Fe-2S]-cluster to mutants, for spectroscopic or stability reasons.

3.1 ADT-bridged [FeFe]-hydrogenase from C. reinhardtii

*Cr*HydA1(ADT) is the active enzyme version of the [FeFe]-hydrogenase from *C. reinhardtii* (therefore denoted *Cr*), in which all intermediates of the catalytic cycle are generally available. Note that more than ten different redox states of the enzyme are nowadays accessible, in which protonation and reduction may occur at different moieties of the H-cluster



Overview of all considered H-cluster states within this review

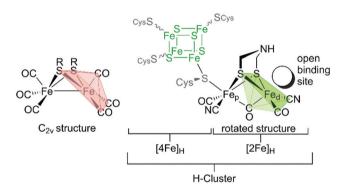


Fig. 8 H-cluster of [FeFe]-hydrogenases with indicated rotated motif of the distal iron atom (green) vs. the $Fe_2S_2(CO)_6$ core in C_{2v} symmetry (red).

during catalysis (Fig. 7). As shown in the experiments on CrHydA1(ADT) with NaDT (sodium dithionite) or H₂, selective enrichment of the intermediate states is challenging, often resulting in blurred spectroscopic results, which impede the exact determination of the nature of each redox state. Using difference spectra is one potential option to handle this problem in IR-spectroscopy. However, for this approach specific equipment is required, which is not accessible in all laboratories.

Therefore, hybrid enzymes, having a bridgehead moiety that differs from adt, as well as mutants providing selectively exchanged amino acids within the peptide backbone are promising options to influence the activity of the enzyme and therefore the accessibility of the H-clusters' redox states. In the next section, we therefore discuss those modifications and

highlight the differences towards CrHydA1(ADT) and their opportunities for spectroscopic applications.

3.2 Reactivity of CrHydA1(ADT) towards oxidising conditions

Hox is the oxidised resting state of the H-cluster and therefore starting point of most conducted experiments. This state can be enriched by treating the enzyme with mildly oxidising reagents such as thionine buffered at pH 8 ($E^0 = 60$ mV vs. SHE at pH 7). However, due to cannibalisation under these conditions, a mixture of Hox and Hox-CO (Section 3.3) is achieved. 54,55 The cannibalisation process is based on the degradation of a fraction of the enzyme sample under influence of e.g. light or oxygen. 54,56,57 Thereby, the released CO binds to the intact H-cluster from a non-degraded enzyme and blocks the active site while forming H_{ox}-CO. 58 Therefore, H_{ox} can be better enriched by auto-oxidation under inert conditions (e.g. N2), which results in a near quantitative enrichment of this state. 55,59

The electronic structure of the enzymes resting state Hox from CrHydA1 was investigated by different techniques. EPR spectroscopy (Table 1) on native CrHydA1 which was not treated with any oxidant or reductant (termed "as-isolated") shows a rhombic 2.1 signal (g = 2.100, 2.037, 1.996) and an axial 2.05 signal (g = 2.052, 2.007). 55,57,60 The former signal resembles the EPR signal of Hox that is known from measurements on hydrogenases from D. desulfuricans (DdH).49,54 The axial signal accounts for the presence of Hox-CO in the as-isolated samples, likewise known from DdH, 56,62,63 and is absent in auto-oxidised samples. 55 ENDOR-spectroscopy (ENDOR = electron nuclear double resonance) on native DdH in combination with

Table 1 EPR values of known H-cluster states in different organisms

	CrHydA1		DdH		СрІ		CpII		TmHydS	
state	g-Value	Ref.	g-Value	Ref.	g-Value	Ref.	g-Value	Ref.	g-Value	Ref.
H _{ox} H _{ox} -CO H _{red} H _{sred} H _{red} ' H _{hyd}	2.10, 2.037, 1.996 2.052, 2.007 Silent 2.076, 1.943, 1.868 Silent Broad signal centred between 2.3 and 2.07	60 60 60 55	2.100, 2.040, 1,999 2.065, 2.007, 2.001 Silent a Silent	49 49	2.10, 2.04, 2.00 2.07, 2.01, 2.01 Silent a Silent	49 49	2.078, 2.027, 1.999 2.032, 2.017, 1.997 Silent a Silent	49 49	2.113, 2.045, 2.001 2.045, 2.918, 2.007 Silent a Silent	61 61

a Not reported.

Mössbauer spectroscopy on hydrogenase II from C. pasteurianum $(CpII)^{48}$ concluded that H_{ox} is best described as a mixed-valent paramagnetic [4Fe-4S]²⁺-[Fe_p¹⁺Fe_d²⁺] complex with the net spin density on the proximal iron atom. 29,49 Due to close similarity of EPR signals from DdH and CrHydA1, Hox was assigned to a [4Fe-4S]²⁺-[Fe_p¹⁺Fe_d²⁺] state.^{55,60} The development of artificial H-cluster maturation (Section 2.2) enabled access to higher amounts of pure CrHydA1, which is especially advantageous for spectroscopic applications and crystallisation experiments. Likewise, the site-selective labelling with ⁵⁷Fe, *i.e.* labelling either [4Fe]_H or [2Fe]_H, became possible with the in vitro approach for the first time. Based on this artificially maturated CrHydA1, recent studies using site-selective X-ray absorption and emission spectroscopy (XAE-spectroscopy) came to opposing results as compared to ENDOR spectroscopy and suggested a [4Fe-4S]²⁺-[Fe_p²⁺Fe_d¹⁺] cluster with the net spin density at the distal instead of the proximal iron.⁶⁴ As a result, XAE- and EPR spectroscopy remain inconclusive regarding the oxidation states of iron within the diiron site. However, it cannot be concluded, if this discrepancy is a result of technical insufficiencies or is even related to the different enzymes used (DdH and CpI/II exhibit additional Fe-clusters besides the H-cluster that might account for the inconclusive results).

Since the ligand environment of H_{ox} is build up from three CO and two CN⁻ ligands, 3,28 IR spectroscopy is yet another strong method for its characterisation and the exact position of ligand vibrations is very sensitive to the electron density and cluster geometry. ^{26,27,29,65} Typically, vibrations of terminally bound carbonyl ligands are found between 2020 cm⁻¹ and 1940 cm⁻¹, whereas their bridging relatives show less intense signals between 1850 cm⁻¹ and 1750 cm⁻¹. In addition, terminally bound cyanide ligands characteristically reveal bands between 2120 cm⁻¹ and 2020 cm⁻¹.

For example, Hox has a very characteristic IR-spectrum revealing two cyanide bands at 2088 cm⁻¹ and 2072 cm⁻¹, two terminal CO bands at 1964 cm⁻¹ and 1940 cm⁻¹ and the bridging carbonyl at 1800 cm⁻¹. 55,56,60,64,66 In 2016, Stripp and coworkers presented a method to selectively label the H-cluster with ¹³CO by controlling the hydration of a protein film and exposing it to ¹³CO gas under light irradiation. ^{56,59} This enabled the CO band assignment to the specific ligands. Furthermore, DFT calculations on all isotope labelled H-cluster variants further suggested largely uncoupled CO vibrations of the Fe-Fe bridging carbonyl (μ -CO, band α , 1800 cm⁻¹) as well as the terminal ligands

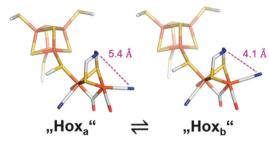


Fig. 9 Conceivable isomers of H_{ox} . Out of two structural isomers of the active-ready oxidised state, "Hoxa" represents the crystallized geometry while " $H_{ox}b$ " is characterized by a partly rotated, distal CN⁻ ligand. Reprinted from ref. 67 with permission from the American Chemical Society, Copyright 2019.

at the distal ($_d$ CO, band β , 1940 cm⁻¹) and proximal ($_p$ CO, band γ , 1964 cm⁻¹) iron atoms.⁵⁹

While the experimentally supported DFT-model and the observed structure from protein crystallography are well in line, 3,28 a second structure with an apical CN was likewise found to be a suitable state (Fig. 9, Hoxb). 59,67 However, while Hoxa was obtained based on XRD measurements at low temperature (approx. 80 K), 3,28 the structure $\mathbf{H_{ox}b}$ stems from a DFT calculation/IR spectra analysis conducted at room temperature. Thus, $H_{ox}b$ might be a higher energy state of H_{ox} . While the reason for this discrepancy is not known, H-cluster flexibility might be an important feature in order to stabilise redox states with additional ligands at the distal iron (i.e. H_{ox} -CO and H_{hvd} , Section 3.5).67,68 The exact assignment of the spectroscopic bands furthermore enabled insight into the electronic structure as well and allowed to resolve the discrepancy from EPR and XAE-spectroscopy, which remained inconclusive regarding the oxidation states of the diiron site. According to IR-spectroscopy, the [Fe_p²⁺Fe_d¹⁺] configuration should be favoured, since the CO band of the proximal ligand is shifted to higher wavenumbers compared to the signal of dCO thus supporting the EPR analysis and hints towards a decreased electron density at the proximal iron centre.

3.3 Reactivity of CrHydA1(ADT) towards CO

As mentioned in the previous section, cannibalisation by treatment with thionine leads to H_{ox} -CO besides H_{ox} . Intentionally, this state can be enriched by treatment of Hox with exogenous CO gas. 62,66 Hammerström, Lubitz and coworkers reported that

the additional carbonyl within Hox-CO can be released by a laser pulse with an energy of 355 nm. Using time-resolved FTIR spectroscopy, a half-life of $t_{1/2}$ = 13 \pm 5 ms of the dissociated CO could be determined, before rebinding occurs. 69 The molecular structure of the H-cluster in its Hox-CO state was likewise investigated by X-ray crystallography of the CO inhibited hydrogenase I from C. pasteurianum (CpI). As expected, electron density that arises from a diatomic ligand was found in apical position of the distal iron, which is accompanied by an elongation of the Fe-Fe distance from 2.56 \mathring{A} in H_{ox} to 2.71 \mathring{A} in H_{ox}-CO. 49,62 The IR spectrum of H_{ox}-CO likewise accounts for the extra CO by an additional band in the region of terminal carbonyl ligands (Table 2, d_2 CO, band δ , 2012 cm⁻¹), while bands β and γ shift to 1962 cm⁻¹ and 1968 cm⁻¹, respectively. 56,58-60,64,70 These band positions reveal a vibrational coupling of the carbonyl ligands, as already reported by Albracht and coworkers.⁶⁶ A pronounced vibrational coupling between all terminal CO ligands was later uncovered by DFT calculations.⁵⁹ The nature of the diatomic apical ligand, however, cannot be determined by XRD analyses due to the close geometric and electronic resemblance of CO and CN-. While an apical CO is favoured by most research groups, IR spectroscopy accompanied by DFT analysis suggested a rotation of the distal iron, enabled by the cluster flexibility (see above) and resulting in an apical CNinstead of an apical CO. 59,64 The electronic structure of H_{ox} -CO is thus still under rigorous debate. The [4Fe-4S]-cluster, consisting of two antiferromagnetically coupled hs-Fe^{II}Fe^{III} (S = 9/2) subclusters, is overall in a +2 (S=0) state, similar to H_{ox} . Interestingly, although for singlet spin states such as observed for [4Fe]_H no hyperfine coupling (hfc) should be observed, a pronounced spin exchange between [4Fe]_H and [2Fe]_H results in a strong hfc that differentiates H_{ox} -CO from H_{ox} , where no hfc is observed. Mössbauer, 48,71,72 EPR, 49,71 and X-ray spectroscopy as well as computational studies, 64,73 agree well with this finding and suggest a $S \neq 0$ state for [2Fe]_H. However, the electronic structure has to be different than in H_{ox} due to the observed hfc of [4Fe]_H. Here, early Mössbauer studies by Popescu and Münck favoured a paramagnetic Fe^{II}Fe^{III} state, whereas a Fe^IFe^{II} could not be excluded. 48 The latter description was, however, favoured by Lubitz and coworkers performing ENDOR measurements on ⁵⁷Fe enriched *Dd*H and suggesting an electronic configuration

Table 2 IR band frequencies of CO and CN⁻ ligands in *Cr*HydA1

State	$\nu (\mathrm{CN^-})/\mathrm{cm^{-1}}$	ν (CO)/cm ⁻¹	Ref.
H _{ox}	2088, 2070	1964, 1940, 1802	75
H _{ox} H	2092, 2074	1970, 1946, 1812	75
H _{ox} -CO	2091, 2081	2012, 1968, 1962, 1808	51
H _{ox} H-CO	2094, 2086	2006, 1972, 1966, 1816	51
$\mathbf{H}_{\mathbf{red}'}$	2084, 2066	1962, 1933, 1792	51
H _{red} 'H	2086, 2068	1966, 1938, 1800	51
H _{red} '-CO	2086, 2076	2002, 1967, 1951, 1793	70
H _{red}	2070, 2033	1961, 1915, 1891	75
$H_{red}H^{\dagger}$	2071, 2032	1968, 1917, 1891	76
H _{red} H ⁺ lt	2079, 2041	1916, 1894, 1810	77
H _{sred}	2068, 2026	1953, 1918, 1882	75
$H_{sred}H^{+}$	2067, 2027	1953, 1917, 1881	76
H _{sred} H ⁺ lt	2070, 2026	1919, 1882, 1803	77
H _{hyd}	2082, 2068	1978, 1960, 1860	75

with a paramagnetic Fe^I in proximal position. ⁴⁹ Although most of the spin density is at Fe_d, a substantial spin delocalisation over the whole cluster was reported as well, which is induced by the binding of the additional CO.⁶³ Due to the spin coupling between [4Fe]_H and [2Fe]_H and the resulting spin distribution, the EPR spectrum of H_{ox} -CO differs from H_{ox} although the redox state seems to be the same. This coupling results in an axial 2.07 signal (g = 2.065, 2.007, 2.001). ^{49,54,63} The strong spin distribution was also detected by XAE spectroscopy on HydA1 from C. reinhardtii that assigned H_{ox} -CO as $[4Fe-4S]^{2+}-[Fe^{1.5+}Fe^{1.5+}]^{3-}$, corroborated by DFT calculations.64

3.4 Reactivity of CrHydA1(ADT) towards reducing reagents - A: sodium dithionite (NaDT)

Relative to Hox, multiple single and double reduced H-cluster redox states are known. These can be accessed by treatment of the as isolated or Hox enriched species with chemical reducing agents. At first, we want to consider treatment of Hox with sodium dithionite (NaDT), which has a potential of -660 mV vs. SHE at pH 7 and is one of the most frequently used reducing agents in the hydrogenase community. In these assays, NaDT compensates the lack of a physiological electron donor and enables the formation of hydrogen.

Treatment of as-isolated CrHydA1 with a 17-fold concentration of NaDT at pH 8, followed by direct freeze quenching of the samples in liquid N_2 results in the loss of H_{ox} specific IR bands in favour of multiple signals (Fig. 10, entry A).⁶⁰ Those signals were originally attributed to the single reduced [4Fe-4S]²⁺-[Fe^IFe^I] complex and a double reduced species that is called H_{sred} (= "super reduced"). However, due to the presence of a second bridging CO signal, the existence of H_{ox} in this assembly cannot be excluded. Especially the corresponding EPR spectrum (Q-band, FID detected) points to the presence of unreacted Hox, showing the characteristic rhombic 2.1 signal. In addition, a broad signal with lower g values (g = 2.076, 1.943, 1.868) is present in the resulting spectrum. This signal resembles a reduced [4Fe-4S]-cluster, which has to stem from a double reduced H-cluster, since [4Fe-4S]²⁺-[Fe^IFe^I] $(\mathbf{H_{red}})$ and $[4\text{Fe}-4\text{S}]^+$ - $[\text{Fe}^{\text{II}}\text{Fe}^{\text{I}}]$ $(\mathbf{H_{red}}')$ are EPR silent.⁷⁴ Therefore, the double reduced H_{sred} has to be denoted as [4Fe-4S]⁺-[Fe^IFe^I] rather than a [4Fe-4S]²⁺-[Fe^IFe⁰].⁶⁰

Contrary, King and coworkers observed different results in their IR and EPR spectroscopic investigation under reducing conditions. While a 2-fold NaDT concentration and incubation at 4 °C overnight led to the same states as observed by Lubitz and coworkers (Fig. 10, entry B), a 10- to 20-fold NaDT concentration and incubation at room temperature for one minute led to decreasing bands at 1933 cm⁻¹, 1883 cm⁻¹ and emerging bands at 1979 cm⁻¹, 1803 cm⁻¹ as well as 1861 cm⁻¹ (Fig. 10, entry C), which were not observed in the spectra of Lubitz and coworkers.55

Subsequently, EPR spectroscopy was used to disentangle the mixture of states found by IR spectroscopy. Samples reduced with 2 equiv. NaDT at 4 °C afford the H_{red}, H_{red} and H_{sred} states according to IR spectroscopy. The corresponding EPR spectrum shows a broad signal between g = 2.3 and 2.07. Since H_{sred}

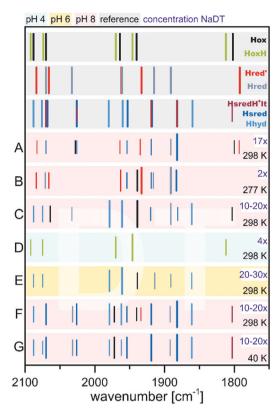


Fig. 10 Schematic representation of IR band positions of NaDT reduced CrHydA1 (colored background) in comparison with currently known states (grey background). The concentration factor (x) of NaDT is related to the protein concentration. IR signatures of known states are taken from Table 3. References for entries A-G: A, 60 B and C, 55 D, 51 E, 78 F and G. 77

should be the only EPR active state, this broad signal must occur from this double reduced species. In addition, the temperature and power properties of the broad EPR signal was found to fit best to a reduced [4Fe-4S]-cluster consistent with the findings of Lubitz and coworkers for H_{sred}. Contrary, utilizing 10- to 20-fold concentrated NaDT reduced samples (Fig. 10, entry C), signals of a hitherto unknown state were observed. The EPR spectrum shows a 90% contribution of a broad rhombic signal of g = 2.077, 1.935, 1.880 and minor contributions from rhombic 2.1 signal (H_{ox}). Again, a reduced [4Fe-4S]-cluster was suggested to be the origin of this signal. Notably, according to a post hoc IR analysis, the dominant species in this sample is H_{hvd}, which was postulated as an intermediate species within the native reaction cycle, e.g. a [4Fe-4S]⁺-[Fe^IFe^I]-H⁺ or a [4Fe-4S]⁺-[Fe^{II}Fe^{II}]-H⁻.⁵⁵ Therefore, H_{hyd} seems to accumulate under strongly reducing conditions, whereas H_{red} cannot be found. Both possible H_{hvd} species are uncharacterized intermediates at the beginning of the H2 splitting cycle or at the very end of the H⁺ reduction cycle, respectively.

The experiments described so far were all performed at pH 8. However, the interconversion of protons and molecular hydrogen is according to the law of mass action always dependent on the pH. Therefore, accumulation of intermediate states was thought to be easier when increasing or decreasing the

proton concentration and thus, shifting the equilibrium to a specific H-cluster state. Following this approach, treatment of CrHydA1 in its Hox state with a 4-fold concentration of NaDT at pH 4 resulted in an unknown species with upshifted CO/CNfrequencies by 4 to 6 cm $^{-1}$ vs. H_{ox} instead of accumulation of a reduced species that would have been accompanied with downshifted CO and CN frequencies (Fig. 10, entry D). The formation of this species was shown to be reversible when switching back to pH 8.51 This new species was denoted as HoxH, indicating a tentative protonation event at the H-cluster but an overall similar redox state as Hox. Interestingly, no change in the IR pattern could be observed in the absence of NaDT, which points to the necessity of reducing ("turn-over") conditions. Although the transition from Hox to HoxH is not a PCET, since both, [4Fe]_H and [2Fe]_H remain in the same oxidation state according to IR-supported DFT calculations, the pH and NaDT dependent formation of HoxH suggests involvement of a PCET in its formation. Potentially, HoxH is the endpoint of the catalytic cycle, which is PCET based.⁵¹ Deprotonation of $H_{ox}H$ is thus the last step of the H_2 formation cycle and results in the regain of H_{ox}. DFT calculations along with IR spectroscopy were performed to investigate the presence of an additional proton within HoxH. The band correlation agreed best with a protonation at the cysteine S9 at [4Fe]_H, whereas protonation at one of the four bridging sulphides of the [4Fe-4S]-cluster led to strong cluster distortions and protonation at the first ligand sphere or at [2Fe]_H led to substantial stronger shifts of the CO and CN⁻ frequencies.⁵¹ Experiments at pH 8 show that H_{hvd} can be found alongside with H_{ox}, H_{red} and H_{sred} under strongly reducing conditions.⁵⁵ Drastically increasing the concentration of NaDT (20- to 30-fold) while simultaneously increasing the proton concentration to pH 6 yields almost pure H_{hyd} with minor amounts of H_{red} and Hox (Fig. 10, entry E). 78 This sample was further analysed by NRVS (nuclear resonance vibrational spectroscopy), providing vibrational information only for Mössbauer active elements, e.g. iron. The resulting spectrum shows two high energy bands at 675 cm⁻¹ and 744 cm⁻¹, characteristic for a terminal bound hydride. Upon changing the medium from H₂O to D₂O both signals are replaced by a new signal at lower energy (629 cm⁻¹), in line with H/D exchange of this terminal hydride. 78 The NRVS spectra were taken as basis for sophisticated DFT calculations to determine a possible structure for H_{hvd} . The best overall agreement between calculated and measured spectra was obtained for a [4Fe-4S]⁺-[Fe^{II}Fe^{II}]-H⁻ species, in which the amine is neutral and with the amine proton pointing towards the hydride forming an internal hydrogen bond.78 Another hydrogen bond is formed between the lone pair of the amine and the adjacent thiol group of Cys169, which is believed to be the last amino acid in a proton channel towards the H-cluster. 79 Besides the proposed H_{hyd} structure, arrangements with a protonated bridge structure (R₂NH₂⁺) were tested as well with very low agreement between the calculated and observed spectra. However, such a state was not completely ruled out but considered as potential intermediate species between H_{hvd} and the H₂ releasing/uptaking state.⁷⁸

Besides the influence of the pH value, the relevance of the temperature at which the experiments are performed was highlighted as well. Incubation with an approx. 10- to 20-fold concentration of NaDT (a 20 mM solution NaDT was used for reduction of CrHydA1; however the concentration of the enzyme is not stated) gave a mixture of H_{hvd}, H_{sred}, H_{red} and in minor amounts $\mathbf{H_{red'}}$ and $\mathbf{H_{ox}}$ both, at 280 K and 40 K.⁷⁷ However, at 40 K the intensity of the signal at 1803 cm⁻¹ increases strongly. This signal was assigned to a double reduced H-cluster state, which bears a bridging CO that is not detected in $\mathbf{H_{sred}}^{60,80}$ (bridging hydride) or $\mathbf{H_{sred}}\mathbf{H}^{+76}$ (semibridging CO and adt-H⁺). This assignment is based on the concomitant increase of H_{sred} signals at 40 K. We therefore denote this species as $\mathbf{H}_{\mathbf{sred}}\mathbf{H}^{\dagger}$ -lt that is claimed as $[4\text{Fe}-4\text{S}]^{\dagger}$ -[Fe^IFe^I] with protonation at the amine, to distinguish between low- and high-temperature states. The low-temperature IR measurements were further strengthened by NRVS measurements and DFT calculations in which both a μ-CO and a μ-H were considered. Models including μ-H produced a high-energy signal around 740 cm⁻¹ which was observed in synthetic μ-H models and experiments of [NiFe]-hydrogenases as well,81 but not found in experiments on [FeFe]-hydrogenases.⁷⁷ A μ-H ligand under these conditions was therefore rendered unlikely, while in conclusion a μ-CO ligand was favoured. However, the unchanged frequency of μ -CO compared to H_{ox} is not explained and remains inconclusive from our point of view. The reduction of [4Fe]_H within the $H_{ox} \leftrightarrow H_{red'}$ transition results in a downshift of the $\mu\text{-CO}$ frequency of 8 cm $^{-1}.^{51,74,76,82}$ The same IR band in H_{sred}H⁺-lt does not shift compared to H_{ox}, although it should result in a larger shift of the μ -CO frequency vs. H_{ox} due to the reduction of [2Fe]_H. However, it must be considered that the measurements at 40 K are under non-physiological conditions, which shows that the temperature at which [FeFe]-hydrogenases are investigated, indeed can influence the outcome of the experiment by means of trapping the H-cluster in different states compared to measurements at room temperature.

3.5 Reactivity of *Cr*HydA1(ADT) towards reducing reagents – B: hydrogen (H₂)

Notably, comparable reactivity alterations were observed upon exchange of the reducing agent – e.g. substituting NaDT as reductant with H_2 . Under physiological conditions, the former results in formation of H_2 and oxidation of an external electron donor, while the latter variant results in the final reduction of an electron acceptor and formation of protons. This interplay is of utmost importance for balancing energy levels of hydrogenases in living organisms. As an additional benefit from changing to H_2 as reducing agent is the determination of the reversibility of catalytic states. If the respective states are accessible from both approaches, H_2 formation and oxidation, theses states are most likely part of a catalytic cycle, while states that are accessible only by one method might lead to biologically less-relevant resting states or artificial, naturally non-appearing states.

According to IR spectroscopy, flushing of CrHydA1 with 100% H₂ for 15 minutes at 4–24 $^{\circ}$ C and pH 8 yields H_{sred} with

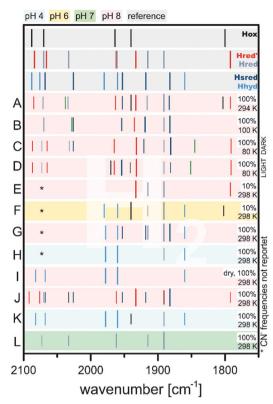


Fig. 11 Schematic representation of IR band positions of H_2 reduced CrHydA1 (coloured background) in comparison with currently known states (grey background). IR signatures of known states are taken from Table 2. References for entries: A_s^{55} B, G^{60} C and D, G^{82} E and F, G^{51} G to I, G^{75} J and G^{67} L.

minor amounts of $\mathbf{H_{red}}$, $\mathbf{H_{red}}$, and eventually very small amounts of $\mathbf{H_{ox}}$ (Fig. 11, entries A, B). ^{55,60} All reduced states can also be accessed by reduction with NaDT, rendering these three states potential intermediate candidates for a $\mathbf{H_2}$ formation cycle. It seems that, depending on the applied temperature, either $\mathbf{H_{sred}}$ (higher temperatures, entry B) or $\mathbf{H_{red}}$ (lower temperatures, entry A) are favoured. However, due to the opposed measurement temperatures, *i.e.* 100 K for samples flushed with $\mathbf{H_2}$ at 24 °C and 294 K for samples prepared at 4 °C, a qualitative analysis of this trend cannot be deduced.

It is worth mentioning that under 100% H_2 , all single and double reduced species are observed. Here, lower amounts of reducing agents, *i.e.* 10% H_2 in N_2 , result in formation of the single reduced species $\mathbf{H_{red}}$ and $\mathbf{H_{red'}}$ at the expanse of $\mathbf{H_{ox}}$, while $\mathbf{H_{sred}}$ seems to be absent in those samples according to IR spectroscopy (Fig. 11, entries E, F).⁵¹

The absence of $\mathbf{H_{sred}}$ in 10% $\mathbf{H_2}$ treated samples is an advantage compared to all NaDT reduced species. Herein, persistent contributions from $\mathbf{H_{sred}}$ crowding the IR spectrum were reported, impeding an evaluation of the resulting spectra. Moving from alkaline pH 8 to more acidic pH 6 (Fig. 11, entries E, F)⁵¹ at constant $\mathbf{H_2}$ concentrations (10%) favoured the formation of $\mathbf{H_{red}}$, $\mathbf{H_{ox}}$ and minor amounts of $\mathbf{H_{hyd}}$ over $\mathbf{H_{red'}}$ and $\mathbf{H_{sred}}$ and *vice versa*. ^{51,72} This behaviour was also observed in a redox titration experiment of CrHydA1(ADT) at different

pH values (Section 3.6) and was accounted to a "non-classical" intra H-cluster PCET, *i.e.* as transition from a [4Fe–4S]⁺–[Fe^IFe^{II}] ($\mathbf{H_{red'}}$) to a [4Fe–4S]²⁺–[Fe^IFe^I] ($\mathbf{H_{red}}$) cluster. ⁷⁶ Simultaneously to the decrease of the $\mathbf{H_{red'}}$ marker band at 1933 cm⁻¹, switching from alkaline to acidic pH decreases the signal of the bridging carbonyl at 1972 cm⁻¹ with the same rate and indicates that $\mathbf{H_{red'}}$ most likely bears a μ -CO, ligand as opposed to $\mathbf{H_{red}}$. ^{51,76} Therefore, the PCET from $\mathbf{H_{red'}}$ to $\mathbf{H_{red}}$ was suggested to be coupled to a ligand rearrangement, which has been considered challenging to merge with the large hydrogen turnover rates of [FeFe]-hydrogenases. ^{51,68,83}

The previously described conditions used by Stripp and coworkers gave small amounts of H_{hvd} upon increasing the proton concentration to pH 6. This work was later revisited by Winkler and coworkers: upon changing the pH from 8 to 4 while purging a sample of HydA1 with 100% H₂ led to IR bands at 1978 cm⁻¹, 1960 cm⁻¹, 1891 cm⁻¹ and 1860 cm⁻¹.⁷⁵ While the signal at 1891 cm⁻¹ stems most likely from H_{red} , the remaining bands were assigned to H_{hvd} (Fig. 11, entries G, H). This example shows, how Le Chatelier's principle can be applied to enrich specific catalytic states of [FeFe]-hydrogenases within the complex biological environment.^{67,75} The simultaneous increase of starting material (H2) and proton concentration (i.e. pH < 6) prevents deprotonation of the H-cluster and traps H_{hvd} . The deprotonation step is therefore presumably involved in the $H_{hyd} \rightarrow H_{red'}$ or $H_{hyd} \rightarrow H_{sred}H^{\dagger}$ conversion.

However, there is an ongoing discussion about the importance and the assignment of specific states leading to sometimes severe alterations of suggested mechanistic schemes. Nevertheless, and independent of the preferred reaction scheme, H_{hyd} was unequivocally suggested to be a key intermediate in the hydrogen cycle.

Identical results were observed using a dry H_2 stream (Fig. 11, entry I) explained by the loss of proton acceptors, *i.e.* the aqueous medium, and therefore accumulation of H^+ within the enzyme. It was suggested that the lost proton acceptors are equivalent to an increase of the proton concentration by lowering the pH, which yields H_{hyd} as well. A similar effect was observed upon impairing the proton transfer path by *e.g.* disrupting it *via* mutagenesis or exchanging the bridgehead of [2Fe]_H (Sections 3.7 and 3.8). 64,75,78,84,85

3.6 FTIR spectroelectrochemistry of CrHydA1(ADT)

As shown in the previous section, treatment of *Cr*HydA1(ADT) with reducing agents generally results in a mix of various redox states depending on the redox potential of the used reductants (Fig. 10 and 11). Importantly, preparing specific desired redox states can be controlled by using electrochemical approaches. In addition, since the redox potentials can be selectively adjusted, each redox state of the H-cluster can be enriched to almost complete purity. Especially in combination with IR or EPR spectroscopy (spectroelectrochemistry, SEC), electrochemical measurements become a powerful tool to investigate *e.g.* proton coupled electron transfers or the redox states in general. ^{52,58,70,74,76,80,86}

At -300 mV vs. NHE or more anodic potentials and broadly independent of pH, H_{ox} is populated according to SEC-IR experiments (Fig. 12, entries A, E, L). 74,76,80 First experiments on CrHydA1 at open circuit potential (OCP) and pH 8 (Fig. 12, entry J) afforded Hox-CO from the cannibalisation process besides H_{ox} . The amount of H_{ox} -CO increased upon switching to -260 mV indicating additional enzyme damage, while the concentration of Hox decreases. Interestingly, going to even more cathodic potentials (-430 mV) led to a complete loss of Hox-CO, while Hred was enriched to almost purity with only minor impurities (Fig. 12, entry N). 58 This behaviour shows that inactive Hox-CO can be reactivated by applying a sufficient reducing potential as long as no exogenous CO is added to the sample. Applying a potential of -300 mV at pH 8 in the presence of exogenous CO yields Hox-CO instead of Hox (Fig. 12, entries Q, T) as well as H_{red}-CO if more reducing conditions are applied (Fig. 12, entries R, S, U). 70,74 Nevertheless, Hox-CO can be found at potentials as low as -600 mV (Fig. 12, entry U). The midpoint potential of the H_{ox} -CO \rightarrow H_{red} -CO conversion was determined to be -360 ± 10 mV at pH 5 (Fig. 12, entries V, W) and -530 ± 30 mV at pH 8 (Fig. 12, entries T, U), 70 which is in line with earlier experiments determining a midpoint potential.⁷⁴ A similar pH dependent behaviour was likewise found for the $H_{ox} \rightarrow H_{red'}$ couple in CrHydA1(PDT) (Section 3.7). A shift of 60 mV per pH unit indicates a proton dependent formation of H_{red}-CO according to the Nernst equation (PCET).70 This protonation event was also assigned to the $H_{ox} \rightarrow H_{ox}H$ transition. It was claimed that the proton herein is located at the [4Fe]_H stabilizing cysteine S9.51 This protonation decreases the electron density of [4Fe]_H and facilitates the reduction of the [4Fe-4S]-cluster, which is in line with the more anodic midpoint potential at pH 5 (protonated Cys S9) vs. pH 8 (unprotonated Cys S9). Supposedly, the electronic structure of H_{red}-CO is therefore best described as [4Fe-4S]⁺-[Fe^IFe^{II}] comparable to the electronic state of H_{red}. The molecular structure of H_{red}'-CO is comparable to H_{ox}-CO with an apical vacancy, blocked by CO.70 According to a DFT-FTIR correlation, rotational freedom of the diiron site can lead to an apical CN ligand stabilised by the adjacent NH bridgehead. Notably, no CO-inhibited form of a reduced [2Fe]_H state (H_{red} or H_{sred}) was found under the tested conditions, i.e. 100% CO, pH 5 or 8 and -100 to -800 mV vs. NHE, which was attributed to a saturated coordination sphere of Fe_d. Therefore, the coordination sphere of the double reduced diiron sites has to be saturated by another ligand, such as a hydride.80

Another transition of interest is $\mathbf{H_{red'}} \to \mathbf{H_{red}}$, which was addressed via H_2 reduction experiments at different pH values by Stripp⁵¹ as well as Lubitz and coworkers. ⁷⁶ In these experiments, protein films of CrHydA1(ADT) were investigated for their IR band signatures at different pH values upon scanning the potential from -200 mV to -600 mV (Fig. 12, entries E to I). The IR signals at 1933 cm⁻¹ and 1891 cm⁻¹ were both found to have a maximum at -380 mV at pH 7, while being absent at -210 mV and -540 mV, respectively (Fig. 12, entries E, G, I). Originally, both IR bands were assigned to the same intermediate. 58,60,74 However, at pH 10 the band at

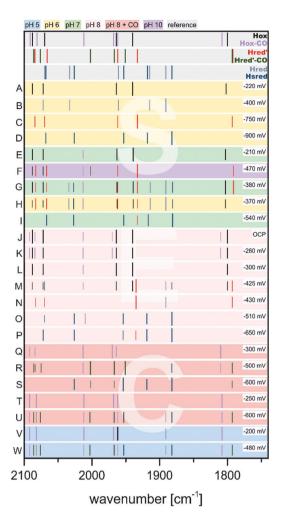


Fig. 12 Schematic representation of IR band positions of electrochemically reduced CrHydA1 (coloured background) in comparison with currently known states (grey background). IR signatures of known states are taken from Table 2. References to entries: (A to D), 80 (E to I), 76 (J, K, N, O),⁵⁸ (L, M, P to S),⁷⁴ (T to W).⁷⁰

1891 cm⁻¹ cannot be found in the IR spectra during the potential scan, while the band at 1933 cm⁻¹ occurs upon shifting to more reducing potentials (Fig. 12, entry F) and indicates a pH dependency of these IR bands. On the other hand, acidic conditions were shown to favour the species responsible for the band at 1891 cm⁻¹ (Fig. 12, entry H). Due to the acidic conditions at which the latter species was observed, it was subsequently attributed to a reduced protonated form $\mathbf{H}_{red}\mathbf{H}^{+}$. (Note: $\mathbf{H}_{red}\mathbf{H}^{+}$ and \mathbf{H}_{red} show the same IR band signature. We herein use both abbreviations to account for their different protonation state, which is not finally clarified and under severe debate within the community; for additional discussion see Section 3.10 'The catalytic cycle' at the end of this section). Consequently, the band at 1933 cm⁻¹ was assigned to an unprotonated reduced form, H_{red} (called H_{red} in the original literature). As a result of this study, the IR signatures of the single reduced H-cluster states H_{red}' and H_{red}/H_{red}H⁺ were assigned as 2084, 2066, 1962, 1933 and 1792 cm⁻¹ for H_{red} and 2070, 2033, 1961, 1915 and 1891 cm⁻¹ for \mathbf{H}_{red} , respectively.⁷⁶ \mathbf{H}_{red} , the

species assembling at alkaline pH, exhibits very small downshifts within the IR spectrum (3 to 7 cm⁻¹) of the CO and CN⁻ vibrations compared to H_{ox} . A reduction of [4Fe]_H in $H_{red'}$ is therefore more feasible than a reduced diiron site and is associated with higher shifts of the CO/CN frequencies compared to H_{ox} and H_{red} . Thus, transition from H_{red} to H_{red} is seemingly coupled to an electron transfer from [4Fe]_H to [2Fe]_H, which is orchestrated by the pH. If the proton pressure is sufficiently high to protonate the diiron site of \mathbf{H}_{red} ([4Fe-4S]+-[Fe^IFe^{II}]), the electron migrates from [4Fe]_H to [2Fe]_H, resulting in [4Fe-4S]²⁺-[Fe^IFe^I]. According to Sommer et al., this value is pH 6, whereas H_{red} dominates already at pH 8 and both are equally present at pH 7.76 This delicate behaviour might be suitable for pH sensing, inducing subtle changes within the protein backbone upon going from \mathbf{H}_{red} (alkaline) to \mathbf{H}_{red} (acidic). The midpoint potential of the $H_{ox} \rightarrow H_{red'}$ transition was found to be -375 ± 10 mV vs. SHE with a strong pH-dependency determined by the protonation event at [2Fe]_H. The latter event results in a plateau of the midpoint potential for high or low pH. For lower pH, the midpoint potential shifts by -50 mV from pH 7 to 6. This observation is also in line with the H_{ox} -CO $\rightarrow H_{red}$ -CO transition, which shows a linear behaviour with a potential shift of 55 mV pH⁻¹ between pH 5 and 8. Both processes are consistent with a PCET from H_{ox} to H_{red} . While the molecular structure of H_{red} is thoroughly discussed in literature, up to now its structure was not finally confirmed. 52,61,76,77,80,87,88

If the potential is swept to more reducing conditions as required for the $H_{red'} \rightarrow H_{red}$ transition, a set of IR bands at 2068, 2026, 1953, 1918 and 1882 cm^{-1} that is similar to the pattern of H_{red} is observed. However, this set is slightly downshifted and better resembles the H_{sred} state that is known from NaDT and H₂ reduction experiments. 55,60 The small average downshift of about 5 cm⁻¹ of the CO/CN⁻ frequencies is in line with a reduction of [4Fe]_H, as was observed for the $H_{ox} \rightarrow H_{red}$ transition. The electronic structure of H_{sred} is therefore most likely a [4Fe-4S]⁺-[Fe^IFe^I] state, which was already found by EPR spectroscopy as well (Section 3.4). 55,60 The potential needed to accumulate H_{sred} is likewise pH dependent. At pH 7 and 8, H_{sred} is obtained as the major species at potentials <-510 mV vs. NHE (Fig. 12, entries O, P). Notably, at less cathodic potentials minor H_{sred} amounts are still present next to H_{ox} and the single reduced species H_{red} and H_{red} (Fig. 12, entries G, I, M). If the proton concentration is increased to pH 5, i.e. conditions that favour the formation of H_{red}, H_{sred} is accessed more easily and found already at potentials of -480 mV. However, at pH 10, i.e. conditions that favour H_{red} over H_{red} , H_{sred} cannot be found. 76 This behaviour of $\mathbf{H_{sred}}$ indicates that it is potentially formed from H_{red} in the reaction cycle, whereas it cannot be accessed from $H_{red'}$.

3.7 Influence of alternative dithiolate bridges on the reactivity of the H-cluster

One of the most striking advantages of the (semi-)artificial maturation process (Section 2.2) is the possibility to implement H-cluster mimics that are different from native CrHydA1(ADT) enabling altered reactivity patterns of the hybrid-enzymes.31

These differences in reactivity can then be utilised to target specific H-cluster states and transitions that are otherwise not observable within the native enzymes due to rival reaction pathways, e.g. $H_{red'} \rightarrow H_{red}$ vs. $H_{red'} \rightarrow H_{hvd}$ or the simultaneous enrichment of multiple states. Such an enrichment of multiple states severely hampers a precise analysis and leads to discrepancies when putting together all the mechanistic puzzle pieces. However, only in case of a similar electronic structures of both, native ADT and the semiartificial enzymes, respectively, proper statements on the various pathways and intermediates are valid. Otherwise the spectroscopically obtained results cannot be transferred from the semiartificial enzyme to the native ADT containing enzyme. For example, a comparable electronic structure of CrHydA1(PDT) compared to the native enzyme can be anticipated due to the similar IR band positions of their CO stretching frequencies, e.g. in the H_{ox} resting state. 74 Likewise, the molecular structures of the semiartificial enzyme variants PDT, EDT, ODT and SDT with a propanedithiolate, ethanedithiolate, oxadithiolate and a thiadithiolate bridge are presented in Fig. 13. In all cases, the artificial H-clusters closely resemble the native [FeFe]-hydrogenase with a bridging carbonyl and an open binding site at the distal iron atom under cryogenic conditions of the XRD experiments.⁸⁹ The principle of CO-ligand rearrangement that occurs upon artificial maturation, i.e. loss of one carbonyl and adopting the rotated structure, is therefore independent of the bridging moiety and seems to be a general feature of [2Fe]_H mimics - at least as long as the steric bulk within the mimic does not prevent accessing the maturation channel.31

One of the most frequent alteration of [FeFe]-hydrogenases in literature concerning bridgehead variations in the enzyme and likewise in biomimetic catalysts (see Section 4) is the utilization of the PDT variant (Fig. 13A). Depending on the maturation conditions, it exclusively adopts either H_{ox} or H_{red} upon maturation. Notably, the PDT variants lack the possibility to enter the H_{red} , H_{sred} as well as H_{hyd} states due to the missing amine functionality and therefore the possibility to undergo an

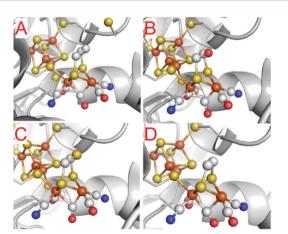


Fig. 13 Molecular structures of the H-cluster from *C. pasteurianum* artificially maturated with (A) propanedithiolate (PDT), (B) ethanedithiolate (EDT), (C) thiadithiolate (SDT) and (D) oxadithiolate (ODT) containing [2Fe]_H clusters. PDB entries 5BYR (PDT), 6H63 (EDT), 5BYQ (ODT), 5BYS (SDT).

intramolecular PCET from $H_{red'}$ to H_{red} or a classical PCET to H_{hvd} . 31,51,76,90

The H_{ox} state of PDT shows an IR spectrum that is equal to that of $\mathbf{H}_{\mathbf{ox}}^{\mathrm{ADT}}$ showing overall similar electronic situations as well as symmetry and thus support the identical structural features found by XRD experiments in their solid state. 51,52,74,90 Likewise, the EPR spectrum of $\mathbf{H}_{\mathbf{ox}}^{\mathrm{PDT}}$ resembles that of $\mathbf{H}_{\mathbf{ox}}^{\mathrm{ADT}}$, showing a rhombic signal centred at 2.1 (g = 2.094, 2.039, 1.998) further supporting the anticipated [4Fe-4S]²⁺-[Fe^{II}Fe^I] state already deduced from FTIR/DFT studies.⁷⁴ Under 100% H_2 at pH 8, H_{ox}^{PDT} undergoes a one-electron reduction and fully converts a single product, namely \mathbf{H}_{red}^{PDT} . This observation is in stark contrast to the ADT samples, in which H_{red} and H_{sred} are found as well.55,59,60,67 Furthermore, the IR spectrum of H_{red}^{PDT}/ shows CN⁻ vibrations at 2084 cm⁻¹ and 2065 cm⁻¹ as well as vibrations of the CO ligands at 1962, 1934 cm⁻¹ and 1798 cm⁻¹ which are comparable to those of the H_{red}^{ADT} spectrum.⁵¹ Increasing the proton concentration from pH 8 to 4 while keeping H2 reducing conditions led in case of ADT to the formation of H_{hyd} . To However, since the PDT analogue is not capable to adopt the H_{hyd} state, another state accumulates under these conditions that shows minor upshifted IR frequencies of all CN-/CO vibrations as compared to H_{red}, i.e. 2084, 2068, 1966, 1938 and 1802 cm^{-1} . A similar shift was observed for ADT upon reducing HydA1 at pH 4 with 2 equiv. NaDT and was denoted to a protonation of cysteine S9 at the [4Fe]_H. This state was called HoxH accounting for the additional protonation (Fig. 7). According to DFT calculations, the same protonation was suggested for H_{red}'. The new upshifted band pattern found in PDT is therefore best explained by a second protonation at one of the [4Fe]_H binding cysteines. According to DFT studies, cysteine S9 is highly favoured as potential protonation site.⁵¹ Due to the additional proton, the new double protonated state is called H_{red}'H. Notably, this state cannot be found in ADT, since then only H_{hvd} is found under otherwise identical conditions as reported for $H_{red'}H$.⁷⁵

Furthermore, H_2 reduction experiments on $\mathit{Cr} HydA1(PDT)$ show that the H_{ox} and $H_{red'}$ analogues can be very easily enriched to purity. While high purity H_{ox} samples can also be obtained in case of HydA1(ADT), $H_{red'}$ commonly comes along with H_{red} and H_{sred} , especially at pH \leq 7. Nevertheless, Lubitz and coworkers were capable to determine the potential of the $H_{ox} \rightarrow H_{red'}$ transition of HydA1(ADT) to be -375 ± 10 mV vs. SHE. 76 A linear correlation between H_{ox} and $H_{red'}$ was found for the CO inhibited species, which are not able to form a reduced $[2Fe]_{H^-}$ cluster species. 70 Contrary, utilising PDT enzyme derivatives enables a direct investigation of the $H_{ox} \rightarrow H_{red'}$ transition e.g. by SEC-IR techniques without the need of CO inhibition and without side reactions.

Fig. 14 presents the Pourbaix diagram of the transition potentials for $\mathbf{H_{ox}} \to \mathbf{H_{red'}}$ (black) and $\mathbf{H_{ox}} \mathbf{H} \to \mathbf{H_{red'}} \mathbf{H}$ (blue) as function of pH value following the peak intensity of the respective marker bands at 1941 cm⁻¹ ($\mathbf{H_{ox}}$), 1934 cm⁻¹ ($\mathbf{H_{red'}}$), 1945 cm⁻¹ ($\mathbf{H_{ox}} \mathbf{H}$) and 1938 cm⁻¹ ($\mathbf{H_{red'}} \mathbf{H}$) as well as subsequent lowering the applied potential from -100 mV to -800 mV νs . NHE.⁵² The E/pH -slopes of 55 \pm 5 mV pH⁻¹

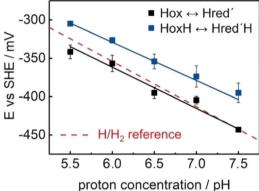


Fig. 14 Pourbaix diagram showing the transition potentials for $H_{ox} \rightarrow$ $H_{red'}$ (black) and $H_{ox}H o H_{red'}H$ (blue) as function of pH value. The slopes are 55 \pm 5 mV pH $^{-1}$ (black) and 50 \pm 3 mV pH $^{-1}$ (blue) with an approximate off-set of 50 mV that elevates the $H_{ox}H \rightarrow H_{red}'H$ potential above the H/H₂ reference (red traces, 59 mV pH⁻¹). Error bars illustrate the quality of the Nernstian fit. Figure and caption are adopted with permission from reference. Reprinted from ref. 52 with permission from John Wiley and Sons, Copyright 2017.

 $(H_{ox} \rightarrow H_{red'}, black)$ and 50 \pm 3 mV pH⁻¹ $(H_{ox}H \rightarrow H_{red'}H,$ blue) indicate a proton coupled reduction of the H-cluster (PCET). Furthermore, the 50 mV lowered reduction potential required for the $H_{ox}H \rightarrow H_{red}H$ transition suggests a protonation of [4Fe]_H directly affecting the transition potential. This pH dependent transition from HoxH to Hred'H is clearly located above the H/H2 reference line (red), which also explains the spontaneous reduction of HoxH in the presence of H2.52 These results are in contrast to the earlier results on the $H_{ox} \rightarrow H_{red}$ transition in CrHydA1(PDT) revealing a midpoint potential of -345 mV at pH $8.^{74}$ The respective transition potentials deviate by approximately 130 mV. This discrepancy was, however, hitherto not addressed in the literature afterwards and remains undissolved so far.

The results of these spectroelectrochemical experiments on CrHydA1(PDT) support the assumption of a proton dependent

reduction of the [4Fe-4S]-cluster in H_{red}. 51 Whereas the formation of H_{red} requires the protonation of the native adt bridge, the very same protonation is inhibited due to the absence of the amine in PDT samples. 76 In both cases, a PCET step directs the additional charge either to the [4Fe]_H or [2Fe]_H. The results on CrHydA1(PDT) are in line with the findings for a CO inhibited species of CrHydA1(ADT), which shows a 60 mV pH^{-1} linear correlation between the H_{ox} -CO $\rightarrow H_{red}$ -CO midpoint potential and pH. However, the transition to H_{red}, in case of the CO inhibited species, is overall 70 mV more cathodic, showing the influence of the additional CO ligand substitution and leads to an increased electron density at [4Fe]_H and therefore an inhibited reducibility. 64,91 The $H_{ox} \rightarrow$ H_{red}' transition in non-inhibited CrHydA1(ADT) was determined to be -353 ± 10 mV vs. SHE at pH 7. Compared to the results of CrHydA1(PDT), the midpoint potential of the native enzyme is approx. 50 mV more anodic.76 The selective conversion from Hox to Hred was further utilized to address the concentration dependency of the formation rate of H_{red}. Diluting the enzyme within bovine serum albumin (BSA) results in a severe drop of the $H_{ox} \rightarrow H_{red'}$ conversion rate.⁵¹ This behaviour was explained by an intermolecular electron transfer (disproportionation) between different [FeFe]-hydrogenase enzymes via two-electron reduced species formed upon treatment with H₂. Likewise, a comparable behaviour was found in whole cell experiments utilising CrHydA1 expressed in E. coli. Monitoring the IR signatures of the H-cluster while purging E. coli cells with 1% H₂ (99% N₂), did neither reveal the specific marker band of $\mathbf{H}_{\text{red}'}$ (1933 cm⁻¹) nor \mathbf{H}_{sred} (1882 cm⁻¹). Both states possess a reduced [4Fe-4S]-cluster, which obviously is hardly trappable in living cells.35

Contrary to such reduced states, CrHydA1(PDT) and other hybrids (e.g. CrHydA1(SDT), CrHydA1(EDT) and CrHydA1(ODT)) enabled the selective formation of Hox upon prolonged exposure to N_2 at pH \geq 8 (auto-oxidation). The CN⁻ and CO frequencies (Table 3) of EDT and SDT resemble those of the ADT and PDT variants, indicating an equal electron density at the Fe-centres. According to quantum chemical calculations at QM/MM and

Table 3 IR band frequencies of CO and CN⁻ ligands in hybrid-CrHydA1

State	$\nu (\mathrm{CN^-})/\mathrm{cm^{-1}}$	ν (CO)/cm ⁻¹	Ref.	State	ν (CN ⁻)/cm ⁻¹	ν (CO)/cm ⁻¹	Ref.
		PDT	EDT				
H _{ox}	2090, 2073	1965, 1941, 1810	52	H _{ox}	2090, 2074	1965, 1941, 1809	67
H _{ox} -CO	2094, 2083	2014, 1972, 1965, 1812		H _{ox} -CO	2094, 2081	2019, 1975, 1967, 1812	67
$H_{ox}H$	2090, 2075	1969, 1945, 1814	52	$H_{ox}H$	2094, 2076	1969, 1945, 1814	67
H _{ox} H-CO	2095, 2086	2013, 1974, 1968, 1816		H _{ox} H-CO	2098, 2087	2071, 1974, 1968, 1819	67
$\mathbf{H}_{\mathbf{red}'}$	2084, 2066	1963, 1934, 1798	52	$\mathbf{H}_{\mathbf{red}'}$	2085, 2067	1961, 1933, 1798	67
H _{red} '-CO	Not observed			H _{red} '-CO	2091, 2080	2015, 1971, 1956, 1807	67
$H_{red}'H$	2084, 2068	1966, 1938, 1802	52				
		ODT			SDT		
Hox	2086, 2070	1972, 1948, 1812	75	Hox	2088, 2070	1969, 1942, 1810	67
H _{ox} -CO	2096, 2085	2038, 1979, 1967, 1811	67	H _{ox} -CO	2094, 2081	2019, 1975, 1967, 1812	67
H _{ox} H	2093, 2078	1974, 1950, 1813	67	H _{ox} H	2091, 2076	1974, 1950, 1814	67
H _{ox} H-CO	2096, 2087	2032, 1980, 1971, 1815	67	H _{ox} H-CO	2096, 2085	2020, 1978, 1971, 1816	67
$\mathbf{H}_{\mathbf{red}'}$	2083, 2070	1964, 1943, 1804	67	$\mathbf{H}_{\mathbf{red}'}$	Not observed		67
H _{red} '-CO	2095, 2081	2011, 1978, 1930, 1806	67	H _{red} /-CO	Not attempted		67
H _{hyd}	2081, 2076	1978, 1962, 1862	75		•		

DFT levels of theory, the overall electronic configuration of all hybrid-enzymes was likewise suggested to be equal to ADT.⁶⁷

Under an atmosphere of 1% CO, all hybrid-enzymes in their oxidised forms adopt the Hox-CO state with the known IR band signature. However, while ADT, ODT and EDT immediately form Hox-CO with near quantitative yields, SDT and PDT revealed slower kinetics and adopt the Hox-CO state only in 65% and 20% yield, respectively. Even in an atmosphere of 100% CO, those two hybrids do not fully convert to the CO inhibited form. In line with those CO inhibition experiments, the decay of H_{ox}-CO to H_{ox} is very fast for SDT and PDT, while it is slow for EDT and very slow and incomplete for ODT and the native ADT forms. In Section 3.3, we already discussed the CO inhibition of CrHydA1 and mentioned the possibility of an apical CN⁻ ligand in H_{ox}-CO based on a partial rotation of _dCN⁻ in H_{ox} . ⁵⁹ The rotation of Fe_d to an apical cyanide ligand in its CO inhibited form might explain the different reactivity of the hybrid enzymes towards CO: While ADT stabilises negatively charged ligands in apical position such as CN⁻ or H⁻⁷⁸ and thus explains the fast CO inhibition and slow decay of H_{ox}-CO^{ADT}, the other hybrids lack the possibility to form this hydrogen bond. Instead, destabilisation of an apical cyanide leading to an altered kinetic was suggested. Further influences of the non-ADT bridgeheads are the steric repulsion in case of SDT and PDT or electrostatic attractions for ODT and likewise SDT. EDT seems to be unbiased due to the missing bridgehead. Therefore, no stabilising or destabilising effects occur resulting in fast CO inhibition and decelerated H_{ox}-CO decay.⁶⁷ Although obvious differences regarding the distal cyanide ligand between native ADT and the hybrid enzymes are present, the CNfrequencies within their IR spectra do not change. This observation cannot be explained from inner sphere ligand coordination and supports the necessity to also discuss outer sphere coordination, i.e. towards the protein environment. This potential influence will be discussed in a separate section concerning the proton transfer pathway (Section 3.8).

Like for CrHydA1(PDT), hybrid enzymes containing ODT, EDT and SDT bridgeheads were tested for their ability to oxidise H₂, thereby adopting the reduced H-cluster states. We already discussed, that PDT does not adopt a diiron-site-reduced form $(H_{red},\,H_{red},\,H_{hyd})$ but is trapped in $H_{red'}$ upon reduction with H_2 31,51,52,67 Whereas SDT stays in the H_{ox} state, indicating no reaction with H_2 , EDT accumulates the H_{hyd} state after initial formation of H_{red}' but returns to H_{ox} very fast in case of dwindling H2. Contrary, the ODT version accumulates Hhvd under H₂ at a very low reaction rate, which was explained by the diminished ability of the ether headgroup to heterolytically support cleavage of H_2 , while for ADT H_{hyd} was accumulated at low pH and simultaneous H₂ or NaDT reduction.^{75,78} Likewise, the regain of Hox from ODT upon switching from H2 to N2, is slower compared to ADT, but faster compared to EDT. In retrospect, it was shown that the amine base of the native H-cluster is of substantial importance, not only for the H₂ development, but also for H₂ oxidation, by stabilising the apical hydride via a hydrogen bond between NH and H⁻.^{75,78} As in H_{ox}-CO, with a proposed apical CN⁻ ligand, this stabilisation is the reason for the different formation and decay rates of ADT νs . hybrid-enzymes, in which an outer-sphere coordination was suggested to stabilise $\mathbf{H}_{\mathbf{hvd}}$ and $\mathbf{H}_{\mathbf{ov}}$ -CO.

The IR spectrum of H_{hyd} (2081, 2068, 1978, 1962 and 1868 cm⁻¹) shows blue shifted CO bands compared to \mathbf{H}_{ox}^{ODT} , which indicates a decreased electron density within the [2Fe]_Hcluster. A Fe^{II}Fe^{II} configuration and a reduced [4Fe-4S]⁺-cluster most accurately explains this finding and is further supported by the characteristic EPR signal of reduced [4Fe-4S]-clusters adopting the same redox state as found in HADT. 84 The Hhvd state is best described with a terminal H⁻ at the apical position of Fe_d and was indirectly observed by the different kinetics of hybrid-enzymes and further exploited from H/D exchange experiments revealing the bridging CO in trans position to the apical ligand of Fed. Due to the trans effect, an H/D exchange trans to µ-CO, results in a downshifted frequency of the bridging CO and indeed this shift is observed in FTIR experiments performed on CrHydA1(ODT). The bands of the terminal CO and CN⁻ did not shift upon the H/D exchange.^{75,84} Likewise, NRVS measurements on CrHydA1(ODT) displayed high energy bands at 670 cm⁻¹ and 727 cm⁻¹ for the bending of a terminally bound hydride (Fed-H), which shifts towards lower energy (564 cm⁻¹ and 625 cm⁻¹) in case of a D₂O/D₂ environment.84 Notably, the results from nuclear resonance vibrational spectroscopy (NRVS) experiments on CrHydA1(ODT) differ from those on ADT, which gave significantly different bands for Fe_d-H at 675 cm⁻¹ and 744 cm⁻¹, respectively.⁷⁸ The DFT based model of H_{hvd} from the NRVS measurements on CrHydA1(ADT) suggests a hydrogen bond between the terminal hydride and the amine headgroup, which cannot be formed in case of the ether moiety of ODT. Although serving as good model for the terminal hydride due to the accumulation of H_{hvd}, the ODT variant of the H-cluster is not able to correctly mimic the hydrogen bonding network between Cys169, NH and Fe_d-H, which is of eminent importance for the reactivity of [FeFe]-hydrogenases. 78 A further possibility to enrich pure H_{hyd} is to impair the proton transfer activity within the enzyme. This can be performed by site directed mutagenesis (SDM) of amino acids within the proton transfer pathway (PTP). 64,75,84-86,92

3.8 Influence of the proton transfer pathway and mutations thereof on the reactivity of the H-cluster

The main trajectory for protons between bulk water and the H-cluster is formed by an H-bond network between the side chains of the five amino acids R286, E282, S319, E279 and C299 (from the surface to the H-cluster, numbering corresponds to *CpI*), two water molecules Wat826 and Wat1120 (cumulated as W1 in Fig. 15, PDB-ID 4XDC, chain B) and the secondary amine of [2Fe]_H. ^{79,93,94} In order to address all amino acids to an individually adequate extend, we will start discussing the most inner located amino acid C299 (C169 in *Cr*HydA1) and its mutants and continue going outwards residue by residue. The first amino acid that undergoes a weak hydrogen bond to the NH moiety of the H-cluster is cysteine C299/C169 (*CpI*/*Cr*HydA1). This side chain is 3.5 Å away from the amine based on the crystal structure 4XDC of *CpI*. A second hydrogen

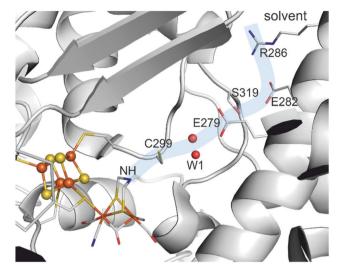


Fig. 15 Proton Transfer Pathway (PTP) of [FeFe]-hydrogenase I from Clostridium pasteurianum between bulk water and the H-cluster via R286(R148), E282(E144), S319(S189), E279(E141), water complex W1 and C299(C169). Amino acid codes in parentheses are the respective residues in CrHydA1. Protein structure from PDB entry 4XDC.

bond (3.2 Å) is formed between the thiol moiety of C299/C169 and W1.79 This inner core of the H-bond network stays rigid during proton uptake and release independent of the H-cluster redox state as shown by IR spectrocopy.95

Due to the direct interaction of C299/C169 and the amine of [2Fe]_H, this amino acid was the target of numerous mutagenesis studies. 64,67,79,85,86,92,96 All performed modifications result in a diminished hydrogenase specific activity, which reflects the importance of the interplay between the cysteine's thiol group, the adjacent water complex and the amine bridge. Most modifications at this position were performed to enrich the H-cluster redox state H_{hvd} . Since it was found that a lack of the proton shuttling ability of the [FeFe]-hydrogenases, e.g. by changing ADT to ODT^{75,84} or blocking the proton transfer pathway H_{hvd} can be enriched, C299A/C169A^{64,85} and C299S/ C169S^{86,92,96} mutants of *CpI* or *CrHydA1* were used to trap the enzyme in H_{hyd} for further spectroscopic investigations of this state. Remarkably, the crystal structure of the C299A_{CDI} mutant (Fig. 16A, green) shows an additional water molecule W_{C299A} occupying the vacant space of the cysteine's thiol. However, this water molecule is not capable to restore the complete

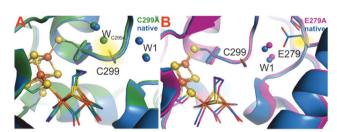


Fig. 16 Native CpI(ADT) in comparison with its mutants CpI-C299A (A, green) and Cpl-E279A (B, magenta). In case of the C299A mutant, water W* adopts the position of the absent thiol. Protein structures from PDB entries 4XDC (native), 6GLY (C299A) and 51A3 (E279A).

enzymatic activity of the mutant but accepts a proton from the H-cluster upon H2 oxidation also leading to an enrichment of the H_{hvd} state.⁷⁹ In contrast to alanine and serine mutants, which show less than 0.2% of the hydrogenase specific activity, the C299D/C169D mutants show 25% and 65% remaining activity, respectively (Table 4). The pH-dependent activity optimum of C299D_{CpI}, however, is shifted from pH 8 (native) to pH 6.5 and shows an altered but intact protonation equilibrium within the PTP.⁷⁹

The next amino acid of the proton transfer path is glutamic acid E279/E141, which strongly interacts with W1 (2.5 Å) via a stable trans complex. 97,98 Furthermore, it only weakly interacts with S319/S189 (3.6 Å), indicating a discontinued proton transfer in the H_{ox} state. Stripp and coworkers determined E279/E141 as the key residue upon switching from proton release to its uptake. 95 Adjusting the CrHydA1 to H_{red} requires a proton from the PTP to enter the H-cluster, i.e. proton uptake reactivity. 51,76 According to IR spectroscopy, E141 of CrHydA1 thereby forms an H-bond to S189 closing the gap between the inner and outer core of the PTP and enables a more continuous proton transport. In line with these results, E141D_{CrHvdA1} loses 90% of its activity due to the longer distances between D141 and W1 and S189, respectively (Table 4).79 The remaining activity of glutamine and alanine mutants at this position is below 1%, indicating the complete loss of the proton transfer ability. In contrast to C299A, the crystal structure of E279A does not show an additional water molecule rescuing the proton transport (Fig. 16B), which explains the low residual remaining activity. In line, the mutant E279A adopts the H_{hvd} state under H₂ oxidising conditions, since the proton transport pathway is interrupted at this position.⁷⁹

S319/S189 is the first residue of the outer PTP and its side chain is 3.6 Å apart from E279/E141 and serves as H-bond acceptor during proton uptake. It further tightly interacts with E282/E144, which is only 2.8 Å apart, as H-bond acceptor and donor during proton release and uptake, respectively.95 Both mutants, S319A of CpI and S189A of CrHydA1 show approx. 5% activity regarding H2 evolution at pH 6.8 relative to the respective native enzymes (Table 4). The crystal structure of S319A from CpI (Fig. 17A) shows an additional water molecule W_{S319A},

Table 4 H₂ production activity of SDM variants targeting the putative PT pathway in Cpl and HydA1. H₂ production activities of PT pathway variants determined at pH 6.8 are presented in % relative to the respective wild-type activity⁷⁹

CpI mutant	H ₂ -Production activity/%	<i>Cr</i> HydA1 mutant	H ₂ -Production activity/%
C299A	0	C169A	0
C299D	30	C169D	65
C299S	0.05	C169S	0.1
E279Q	0.65	E279Q	0.2
E279A	0.06	E279A	0.1
E279D	30	E279D	5
S319A	5	S189A	10
E282Q	5	E144Q	0.45
E282A	60	E144A	45
E282D	80	E144D	50
R286A	90	R148A	55
R286L	300		

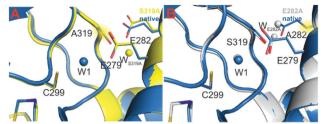


Fig. 17 Native Cpl(ADT) in comparison with its mutants Cpl-S319A (A, yellow) and CpI-E282A (B, grey). In case of the S319A mutant, water W_{S319A} adopts a position near the serine vacancy, closing the PTP. The crystal structure of E282A shows to invaded water molecules W_{E282A}1 and W_{E282A}2, one at the vacancy of the carboxylic acid and one more outwards, forming a makeshift proton pathway between bulk solvent and S319. Protein structures from PDB entries 4XDC (native), 6GM4 (S319A) and 6GM1 (E282A)

which does not exactly occupy the vacant -OH site of serine but is in proximity to E279 and E282 and closes the proton transfer path. As seen before in C299A, the makeshift water molecule changes the pH-dependent activity of the mutant to the highest activity between pH 6.5 and 7. However, in contrast to the C299A mutant, this modification does not lead to the accumulation of H_{hvd} under H₂ oxidizing conditions but shows the reduced species H_{red}, H_{sred} and H_{red} besides H_{ox}.

Due to the missing possibility of S319A to form H-bonds to its neighbours, the next amino acid residue in the PTP, glutamic acid E282, points more towards the arginine residue R286 at the edge between enzyme and solvent. In native [FeFe]hydrogenases, E282/E144 and S319/S189 interact via a hydrogen bond of 2.8 Å length. The deprotonated side chain might further form a salt bridge to the guanidine moiety of arginine R286.99 Mutants of E282/E144 show overall diminished activity regarding H₂ evolution, which is, however, less pronounced as observed for e.g. E279/E141. Like for the more inwards positioned glutamic acid, E282Q/E144Q show only 5% and 0.5% remaining H2 release activity and a pH-dependent activity maximum between pH 6 and 6.5 according to the altered pK_a value of the glutamine residue (Table 4). Remarkably, the aspartic acid mutants E282D/E144D show 80% and 50% remaining activity, respectively (Table 4). This contrasts the respective E279D/E141D mutants, which show significantly less activity. This discrepancy demonstrates the significance of the distinct amino acid residues for the catalytic PTP. However, the importance of proper amino acids within PTP appears to decrease from the H-cluster to the enzyme's surface. Interestingly, the E282A/E144A mutants show 60% and 50% residual activity, which again drastically differentiates E282/ E144 from the inner laying E279/E144 (Table 4). The remarkable activity of these mutants most likely stems from two water molecules invading from the bulk solvent into the PTP (Fig. 17B). One is located directly at the vacancy of native E282, 2.5 Å away from the hydroxyl moiety of S319, and another one is placed more outwards in proximity to the bulk solvent, as revealed by XRD studies of E282A from CpI. Therefore, those water molecules build up an H-bond network from the solvent to S319, taking care of proton transfer in the outer core of the PTP. However, as

previously described for other mutants, this is well in line with a slight shift of the maximal activity from pH 8 to pH 7.79 Besides its H-bond towards S319/S189, the side chain of the glutamic acid residue E282/E144 interacts with R286/R148, which is positively charged/protonated under physiological conditions and therefore forms a salt bridge with negatively charged/deprotonated E282/E144 of 2.8-3.1 Å. 79,95 IR spectroscopic investigations on the R148 from *Cr*HydA1 propose a neutral side chain during proton uptake (formation of H_{red}), which renders the arginine residue the first proton donor.95 A permanently neutral charge at the position of R286/R148 was achieved by the R286L mutant.⁹³ Surprisingly, under conditions of non-rate limiting electron transport, i.e. using methyl viologen as mediator, the R286L mutant from CpI surpasses its native counterpart by the factor of three and is therefore the only mutant with increased activity in comparison to the native enzyme. This interesting finding was explained by the absence of the salt bridge between R286 and E282 that neutralises the negative charge of the carboxylic acid. In R286L, this charge is still present increasing the driving force for protons to enter the PTP. 93,99 Furthermore, E282 is more exposed to the bulk solvent in R286L, as seen from Zn²⁺-inhibition experiments.

3.9 Influence of an additional F-domain on the reactivity of the H-cluster

The reactivity discussed so far is based on results from [FeFe]hydrogenases HydA1 from C. reinhardtii. Especially due to the possibility to artificially maturate this enzyme, most research concentrates on this "blueprint" for H-cluster reactivity. The main difference between HydA1 and other [FeFe]-hydrogenases is the electron supporting chain of two [4Fe-4S]-clusters, so called F-clusters, which are absent in eukaryotic CrHydA1 but present in prokaryotic hydrogenases, e.g. DdH and CpI. The F-clusters will be denoted as d[4Fe]_F (distal F-cluster, relative from the H-cluster) and $_{p}[4Fe]_{F}$ (proximal F-cluster). We will stress the resulting differences in reactivity after describing the similarities between HydA1 and prokaryotic hydrogenases. The Hox state of CrHydA1, i.e. the smallest possible [FeFe]hydrogenase, exhibits only the H-domain and is characterized by a rhombic EPR signal (g = 2.10, 2.037, 1.996), an IR signature with uncoupled vibrations for each CO/CN⁻ ligand (2088, 2070, 1964, 1940, 1802 cm⁻¹) and an electronic structure with an oxidised [4Fe-4S]²⁺ cluster and a bi-valent diiron site (Fe^{II}Fe^I). Upon CO binding, the IR pattern changes to 2091, 2081, 2012, 1968, 1962 and 1808 cm⁻¹, respectively, accounting for the additional CO ligand and the EPR spectrum changes from a rhombic to an axial signal (g = 2.052, 2.007). The IR spectra of $\mathbf{H_{ox}}$ and $\mathbf{H_{ox}}$ -CO from $Dd\mathbf{H}$ (hydrogenase from D. desulfuricans) show characteristic signatures similar to H_{ox} and H_{ox} -CO from CrHydA1 (see Tables 2 and 3).56 Likewise, the EPR spectrum of DdH in the oxidised state agrees with the observed spectrum of HydA1 (see Table 1) but shows additional signals for the F-cluster. ^{54,91} In case of H_{ox} , a broad rhombic signal (g = 2.059, 1.935, 1.877) is present in the spectrum as well. This was assigned to $_{\rm d}[4{\rm Fe}]_{\rm F}$ since a spin coupling between the H-cluster and $_{\rm p}[4{\rm Fe}]_{\rm F}$ is expected and would result in an overall EPR silent state if the

distal H-cluster is reduced. 91 The overall electronic and molecular structure of the H-cluster in Hox and Hox-CO from CrHydA1 and DdH can therefore considered as identical, which is also in line with matching 57Fe hyperfine values, examined by HYSCORE spectroscopy, from both enzymes. 49,71 However, differences occur for reduced enzymes.⁹¹ Under slightly reducing conditions, i.e. 50 μ M NaDT, the IR spectrum of DdH(PDT) shows a mixture of $\mathbf{H}_{\mathbf{ox}}^{\mathrm{PDT}}$ and $\mathbf{H}_{\mathbf{red}'}^{\mathrm{PDT}}$. However, the typical rhombic signal of $\mathbf{H}_{\mathbf{ox}}$ in the EPR disappears, whereas the broader rhombic signal (g = 2.059, 1.935, 1.877) remains in the spectrum and another broad signal around g = 2.01 appears. This result was explained by an equilibrium between a state, in which both F-clusters are reduced and the H-cluster is oxidised (F_{red}/F_{red}/H_{ox}), and a state in which the H-cluster and d[4Fe]_F are reduced while p[4Fe]_F is oxidised $(\mathbf{F}_{red}/\mathbf{F}_{ox}/\mathbf{H}_{red})$. In the former state, $_{p}[4\text{Fe}]_{F}$ and the H-cluster combine to an electronically coupled cluster pair, resulting in an altered EPR spectrum compared H_{ox} (broad g =2.01), while in the latter state both, the H-cluster and the proximal F-cluster are EPR silent. 91 To account for this equilibrium, the respective states are called $F_{red}\!/H_{ox}$ and $F_{ox}\!/H_{red'}$ with both states bearing an EPR active reduced distal F-cluster. Notably, the IR maxima of e.g. F_{red}/H_{ox} shift by approx. 1 cm⁻¹ compared to Fox/Hox, which was observed in high-res SEC-IR experiments. 91 Based on these EPR results, an effect of the cluster pairing regarding the transition potentials of the H-cluster is likely. The apparent midpoint potential of the $H_{ox} \rightarrow H_{red'}$ transition in DdH(PDT) is -500 mV vs. SHE at pH 8, which is

certainly more negative than that observed for CrHydA1(PDT) (depending on the reference: 25 mV⁵² or 155 mV⁷⁴ more negative). Furthermore, the two-state population, followed by IR spectroscopy, does not show a Nernstian behaviour. This is a result of redox anti-cooperativity, which was rationalised by simulating the resulting population curves. Therein, the extra electron upon reduction of F_{ox}/H_{ox} was allowed to equilibrate between $_{p}[4Fe]_{F}$ ($\mathbf{F_{red}}/\mathbf{H_{ox}}$) and $[4Fe]_{H}$ ($\mathbf{F_{ox}}/\mathbf{H_{red'}}$), where the ratio was determined by the respective cluster potential. Taking the small IR band deviation of 1 cm⁻¹ for reduced p[4Fe]_F into account, the new four-state population curves became strictly Nernstian. The new model includes a reduction potential of [4Fe]_H, which depends on the redox state of p[4Fe]_F. If p[4Fe]_F is already reduced, a more negative potential for [4Fe]_H was observed and vice versa, i.e. redox anticooperative behaviour.91

In CrHydA1, CO inhibition leads to increased electron density at [4Fe]_H and results in a more negative reduction potential compared to non-inhibited species (Section 3.3). The same observation was found for DdH(ADT) as well. The potential shift of [4Fe]_H is strong enough to prevent an equilibrium between $F_{red}/H_{ox}\text{-CO}$ and $F_{ox}/H_{red}/\text{-CO}$ due to the much more positive reduction potential of p[4Fe]_F. Therefore, F_{red}/H_{ox} -CO is exclusively present. To achieve H_{red} -CO, a second electron is needed to enter the Fred/Hred/-CO state, which is again, as observed for DdH(PDT), formed at much more negative potential due to anti-cooperativity (Fig. 18).

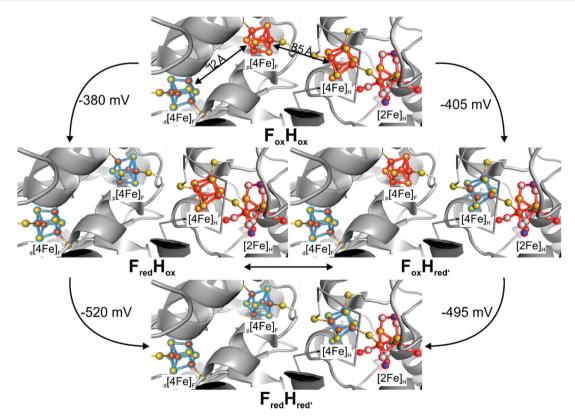


Fig. 18 Four-state model of the redox-anticooperative effect, which results from the additional F-clusters in [FeFe]-hydrogenases (except HydA1). Clusters coloured in red are oxidized, clusters coloured in blue are reduced. Model depicted from ref. 91.

These findings explain why H_{red} -CO in DdH cannot be observed under the conditions of CrHydA1.

Even more interesting is the comparison of CrHydA1(ADT) and DdH(ADT) since former shows a fascinating pH-dependent behaviour regarding single-electron reduced species: at alkaline pH (i.e. pH \geq 8) H_{red} is favoured, while at acidic pH H_{red} is preferred (see Section 3.6).⁷⁶ Due to the amine bridge, such a behaviour can be expected for DdH as well but is observed to another extent. At pH 6, Hred dominates upon reducing the potential, while $H_{red'}$ is not observed. This is in line with the findings from HydA1. Interestingly, H_{sred} , which replaces H_{red} at even more negative potentials in CrHydA1 cannot be found in DdH under the same conditions. The hypothesis explaining the absence of H_{sred} in DdH is that the second electron, required to form H_{sred} and finally liberate H₂ can be stored in the F-clusters instead of the H-cluster. Due to the anticooperative effect of the additional [4Fe-4S]-cluster, the H-cluster's pK_a shifts towards a value close to that of H_{sred} in HydA1. Therefore, bacterial hydrogenases may be able to skip H_{sred} (the participation of H_{sred} in the catalytic cycle is, however, anyway under debate) and directly form H_{hvd}. H_{hvd} was found in DdH under the same conditions as in CrHydA1, being low pH and constant H₂ supply, showing the importance of this state in both organisms.80 Decreasing the proton concentration to pH 8 results in H_{red} being formed in HydA1. However, in DdH H_{red} is still the dominant species. Although the amount of both species in DdH increases simultaneously at potentials between -200 and -400 mV, H_{red} vanishes beyond this potential. According to simulations, the proximal F-cluster is reduced at approximately -400 mV, which results in an anticooperative effect and renders the reduction of [4Fe]_H less likely to occur. Apparently, the pK_a of the amine bridgehead shifts from \approx 7.7 for HydA1 to \approx 9.3 in *Dd*H due to the redox anticooperativity. 91 However, the influence of a protonation at the stabilising cysteine residue S9 of the [4Fe]_H, which might be influenced by the reduction of the proximal F-cluster, was considered as possible explanation. This should be addressed in an additional study since the formation of $H_{ox}H$ under similar conditions as in CrHydA1 was reported for DdH as well. H_{ox}H is associated with the H-cluster in oxidised form with an additional proton at cysteine S9, which causes minor (4 to 6 cm⁻¹) IR band shifts.⁵¹ The complex situation of two different protonation events (cysteine S9 and adt) besides the three different reducible clusters (p[4Fe]_F, [4Fe]_H and [2Fe]_H) makes the interplay of protons and electrons in DdH a very interesting topic to study albeit a challenging one.

Like DdH, hydrogenase I from C. acetobutylicum (CaI) bears two additional F-clusters next to the H-cluster within the protein frame. Compared to CrHydA1 and DdH, the IR spectrum from auto-oxidised CaI in the H_{ox} state shows slightly downshifted CN^- (2082 cm⁻¹ and 2070 cm⁻¹) and upshifted CO stretching frequencies (1969, 1646 and 1800 cm⁻¹). The EPR spectrum, however, closely resembles that of DdH and CrHydA1 showing a rhombic 2.1 signal (g = 2.009, 2.039, 1.999). At acidic pH, the IR bands slightly shift towards higher wavenumbers, as was observed for CrHydA1 and DdH in the $H_{ox} \rightarrow H_{ox}H$

conversion as well, which indicates a similar reactivity at this point. 51,101 Due to the accessory F-clusters, the reactivity of CaI seems to be very close to that of DdH. At pH 8, upon reducing the enzyme by NaDT or by photocatalytic electron supply, \mathbf{H}_{red} is formed indicated by the marker band at 1899 cm $^{-1}$ within the IR spectrum, whereas \mathbf{H}_{red} is dominant under these conditions in HydA1. The reduction is likewise accompanied with a seemingly broad rhombic signals within the EPR spectrum that can be assigned to the reduced F-clusters (g = 2.043, 1.941, 1.911 and g = 2.073, 1.930, 1.868), which was also observed in DdH.

At 13 K, the IR spectrum of NaDT reduced CaI shows bands at 2055, 2040, 1921, 1899 and 1801 cm $^{-1}$. More interestingly, a μ -CO band is observed under these conditions. As previously described for CrHydA1, this may arise from the cryogenic conditions at which the spectrum was recorded. ^{77,102} Changing conditions to D_2O did not yield any observable shifts, which renders a terminal bound hydride at Fe_d unlikely. Unfortunately, no room temperature data are available to discuss the existence of a bridging CO vs. a bridging H $^-$, which is proposed as one possible structure of a single-reduced H-cluster intermediate as well. ⁸⁰

Experiments on CrHydA1, in which specific amino acids were exchanged by site directed mutagenesis showed that e.g. Hhvd is enriched without the need of increasing both, proton concentration and H₂ pressure.⁷⁵ This is also true for CaI, whose IR spectrum shows the presence of two formerly unobserved species at room temperature upon changing C298 to Serine. Decreasing the temperature to 10 K results in the vanishing of bands at 2042 cm⁻¹, 2022 cm⁻¹, 1892 cm⁻¹, 1978 cm⁻¹ and 1781 cm⁻¹, which were assigned to H_{sred} . The remaining bands at 2083 cm⁻¹, 2067 cm⁻¹, 1977 cm⁻¹, 1964 cm $^{-1}$ and 1851 cm $^{-1}$ were assigned to \mathbf{H}_{hyd} . The fact that both states, \mathbf{H}_{hyd} and \mathbf{H}_{sred} , are only observed if the proton path is blocked, renders them possible tautomers and endpoints of the catalytic H₂ oxidation. As discussed above for DdH, the existence of reduced F-clusters can compensate the second reduction step from a single reduced to a double reduced H-cluster species before H_{hyd} is formed. There, H_{sred} is not observed and a proposed transient state in the $\mathbf{H}_{red}\mathbf{H}^{\dagger} \to \mathbf{H}_{hvd}$ transition. Most likely, this is also the case for CaI since the F-clusters should show the same effect here. Additionally, H/D exchange experiments showed that the H_{red}H⁺ → H_{hvd} transition is slower in case of D2O, resulting in accumulation of **H**_{red}**H**⁺. The slowed kinetics implicate an intramolecular proton transfer for the transition from $\mathbf{H}_{red}\mathbf{H}^{+}$ to \mathbf{H}_{hvd} . Both enzymes, DdH and CaI, exhibit additional F-clusters for electron relay and most likely changing the H-cluster's electronics. To determine the general function of these additional clusters, all enzymes excluding CrHydA1 should be investigated regarding their redox anti-cooperativity as performed on DdH. This includes enzymes from Clostridiae (CaI and CpI) and eventually HydS from T. maritima, which exhibits F-clusters as well and is known as sensory [FeFe]-hydrogenase. 61 Sensory-type hydrogenases exhibit a third accessory [4Fe-4S]-cluster at the C-terminus, which is ligated by a Cx2Cx4Cx16C motif. Although the H-cluster of TmHydS is structurally the same as in CrHydA1, CaI and DdH, the reactivity

of the sensory-type hydrogenase is dramatically different from the prototypical-type hydrogenases, which were described above. First indication of the altered reactivity is the 100-fold and 5-fold lower activity in H₂ production and oxidation, respectively, of TmHydS compared to CrHydA1.⁶¹ This can be explained by the altered amino acid ligation of the H-cluster compared to e.g. CpI. While Cysteine (C299) is the endpoint of the PTP in prototypical CpI, an alanine (A131) occupies this position in TmHvdS. Likewise. methionine residues M353 and M497, which are part of the H-cluster's coordination sphere in CpI, are replaced by G177 and S267 in TmHydS, respectively. 103 Mutagenesis experiments of CpI showed that variants of respective amino acids, i.e. C299S, 353L and M497L, led to diminished activity as well. 104 In addition, upon maturation of apo-HydS with an artificially synthesised H-cluster (under 2% H₂), an IR spectrum with CN⁻ bands at 2055 and 2022 cm⁻¹ and CO bands at 1894, 1871 and 1763 cm⁻¹ is observed (Table 5). While this pattern is not changed under reductive conditions (H2 or NaDT), oxidation with thionine led to an altered spectrum of 2088, 2079, 1971, 1947 and $1806~{\rm cm}^{-1}$ that resembles Hox spectra of e.g. DdH and CaI. In conclusion, HydS exclusively adopts a reduced state under already minor amounts of reducing agents, which is not observed for other [FeFe]hydrogenases. The latter shows major amounts of Hox upon maturation under the same conditions. The 30 to 80 cm⁻¹ downshift vs. H_{ox} of IR bands in the reduced state implies a reduction of $[2Fe]_H$ as in \mathbf{H}_{red} of CrHydA1. However, the observed CO/CN frequencies do not match those from H_{red} in HydA1, especially the observed µCO band in TmHydS cannot be associated to an H_{red}-like state (compare Tables 2 and 5). Therefore, the reduced state in TmHydS is called H_{red*}.

3.10 The catalytic cycle(s) of [FeFe]-hydrogenases

Despite the intensive investigation on the catalytic cycle of [FeFe]-hydrogenases in the last decades, up to today, not "one and only" working mechanism of this enzyme family is known. However, based on the reactivity described in this section, some cycles were suggested, which will be briefly described in this section. Especially, we aim at identifying similarities and major discrepancies of those cycles, which are mainly based on inconsistent findings on the reduced H-cluster states \mathbf{H}_{red} and \mathbf{H}_{sred} . It is evident from the results presented in this

section that the exact nature of the states involved in the catalytic cycles is an ongoing matter of a lively debate in the community and hence we will leave the final judgement to the reader as we do believe that all of those currently reported mechanisms have strengths and weaknesses. We anticipate that future theoretical and experimental insights will lead to continuous reassessments and changes in the catalytic cycles.

In Fig. 19–21, three proposed catalytic cycles are presented, which are adapted from a recent joint publication of the groups of Lubitz, Birrell and Dyer. 105 From sub-turnover time-resolved IR spectroscopy, the authors derived cycle C, since all contained states were identified in their experiments. However, mechanisms A and B are very similar to C, considering the main H-cluster states H_{ox} , H_{red} and H_{hyd} . All cycles start with H_{ox} , which is the overall accepted entry point of the catalytic cycle. Hox is characterised by many different techniques (see above) as a $[4Fe-4S]^{2+}$ - $[Fe_p^{2+}Fe_d^{1+}]$ -cluster with a bridging μ -CO ligand and an apical vacancy at the distal iron, with an eventually slightly apical rotated CN⁻ ligand. ^{29,49,64,67} From here, the first reduction event occurs at the [4Fe-4S]-cluster, which is especially plausible in hydrogenases with additional F-clusters as electron delivery chain but also true for HydA1, which lacks this chain of [4Fe-4S]-clusters. 41,51,57,82 However, the exact mechanisms of the electron transfer are different in cycles A versus B and C. While in cycle B and C a simple electron transfer forms H_{red} , a PCET is responsible for H_{red} formation in cycle A. H_{red} is described similar to $\mathbf{H_{red}}$, as a $[4\text{Fe-4S}]^{1+}-[\text{Fe}_p^{2+}\text{Fe}_d^{1+}]$ complex with an Hox like molecular structure, but with an additional proton at one of the [4Fe-4S]-cluster stabilising cysteines (in blue colour in Fig. 20). 51,52,88

PCETs are common in nature and advantageous for a multielectron process due to the balanced charge of the reduced moiety. In all cycles the next state, which could be trapped and thoroughly investigated by numerous different techniques is \mathbf{H}_{hyd} . However, the mechanism to get there differentiates the cycles. While in cycle A a second PCET from $\mathbf{H}_{red'}$ directly results in \mathbf{H}_{hyd} , cycles B and C follow a successive mechanism of separate proton- and electron-transfer steps. A proton transfer to \mathbf{H}_{red} results in the formation of $\mathbf{H}_{red}\mathbf{H}^+$, which electronic structure was characterized by Lubitz and coworkers as $[4Fe-4S]^{2+}-[Fe_p^{1+}Fe_d^{1+}]$ with a protonated amine bridgehead,

Table 5 IR frequencies of known H-cluster states from hydrogenases with an additional F-domain

State	ν(CN ⁻)/cm ⁻¹	ν (CO)/cm ⁻¹	Ref.	State	$\nu (\mathrm{CN^-})/\mathrm{cm^{-1}}$	ν (CO)/cm ⁻¹	Ref.
		DdH	TmHydS				
H_{ox}	2093, 2079	1965, 1940, 1802	75	H_{ox}	2087, 2079	1971, 1947, 1806	61
H _{ox} H	2097, 2082	1965, 1940, 1802	75	H _{ox} -CO	2090	2016, 1973, 1964, 1805	61
H _{ox} -CO	2096, 2089	2017, 1972, 1963, 1812	91	H _{red*}	2055, 2022	1894, 1971, 1763	61
H _{red}	2079, 2040	1915, 1892, 1962	75	H _{sred} *	2047, 2013	1900, 1861, 1751	61
H_{hyd}	2089, 2079	1980, 1963, 1860	75				
CaI						СрІ	
Hox	2082, 2070	1969, 1646, 1800	101	H_{ox}	2082, 2071	1970, 1947, 1800	75
H_{red}	2052, 2035	1914, 1894	101	$H_{ox}H$	2084, 2073	1975, 1953, 1809	75
$H_{red}H^{+}$	2055, 2040	1921, 1899, 1801	101	H_{red}	2071, 2053	1915, 1899, 1962	75
H_{hvd}	2080, 2063	1975, 1960, 1849	101	H_{sred}	2065, 2039	1922, 1894, 1958	75
.				H _{hyd}	2082, 2068	1984, 1968, 1856	75

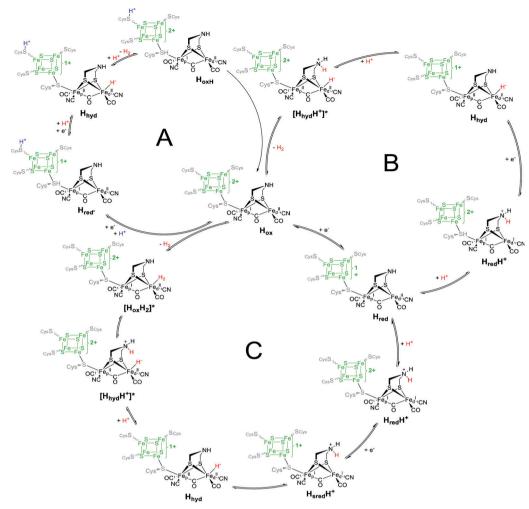


Fig. 19 Main proposed catalytic cycles A-C adapted from ref. 87 and 105

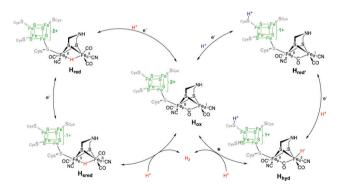


Fig. 20 Shortened catalytic cycle A from Fig. 3.11 and secondary cycle, which is based on bridging hydrides. Red protons: catalytic PTP, blue proton: accessory PTP. *HoxH might be involved in this conversion.

which is why the additional electron from the four-iron cluster has to migrate to the diiron site. ^6 In cycle C, the following electron transfer forms $\mathbf{H}_{sred}\mathbf{H}^{+}$, that isomerises to \mathbf{H}_{hyd} . In cycle B, the latter state is directly formed upon electron transfer. Considering the functionality of the amine bridge within the PTP, an intermediate with a protonated bridgehead

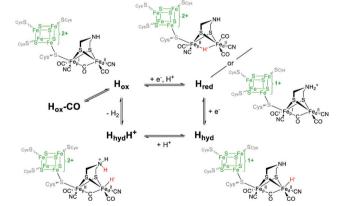


Fig. 21 Schematic representation of the catalytic cycle of [FeFe]hydrogenase, based on the states observed under whole-cell conditions including the protonated hydride species ($H_{hyd}H^+$). Figure and caption from ref. 35.

is feasible - at least as very short-lived transient state - even in cycle A. This shows the hen-and-egg problem of a PCET, which may be facilitated via an electron-proton-transfer or an protonelectron-transfer. 107

Starting from H_{hvd} , the second protonation of the H-cluster via the catalytic PTP results in the formation of H2 and the regain of Hox. In cycle A, the H-cluster passes through HoxH, an H_{ox}-like state that still holds the non-catalytic proton near the [4Fe-4S]-cluster. Cycles B and C further contain the intermediate species H_{ox}-H₂ and/or H_{hvd}H⁺, which is feasible considering the role of the amine bridge. The H_{hyd}H⁺ state was recently observed in in vivo studies on HydA1 containing E. coli cells and is a plausible intermediate in the $H_{hvd} \rightarrow H_{ox}$ conversion.³⁵

Blanking out the non-catalytic proton in cycle A, cycles B and C can be considered as more detailed mechanisms of cycle A. All used H-cluster states share the Hox-like geometry with a bridging CO, which was supposed to be a prerequisite for fast turn over. However, in recent literature, structures containing a bridging hydride instead of a bridging CO were discussed as well. 80,88 Those structures are thermodynamically more stable and therefore might not justify fast turn-over rates as observed for [FeFe]-hydrogenases, hence they were placed in a secondary cycle (Fig. 20) with exclusively bridging hydrides, which was formerly denoted as "slow cycle".87 We already discussed that both single-electron-reduced states $H_{red'}$ and H_{red} can be formed upon reducing H_{ox} and that pH dictates whether the electron resides at the [4Fe-4S]-cluster, stabilising the Hox-like structure at more alkaline pH, or migrates to the diiron subsite forming $\mathbf{H}_{red} (\mathbf{H}_{red} \mathbf{H}^{\dagger})$ at pH \leq 6. At this point, a problem regarding the nomenclature of the specific states arises. The state which is entered as first species in all cycles, i.e. [4Fe- $4S^{1+}-[Fe_p^{2+}Fe_d^{1+}],$ is called $H_{red}^{76,77,105}$ or $H_{red}^{76,77,88}$ depending on the additional proton near the [4Fe-4S]-cluster. The single-electron-reduced [4Fe-4S]²⁺-[Fe_p¹⁺Fe_d¹⁺] state is called $H_{red}^{80,87,88}$ or $H_{red}H^{+,76,77,102,105}$ depending on the described intermediate and considered mechanism. The same is true for the double reduced state $H_{sred}/H_{sred}H^{\dagger}$.

Unfortunately, no XRD studies that would give hints towards the spatial structure of those single reduced states are present. Therefore, EPR and vibrational spectroscopy in combination with computational techniques are the methods of choice to characterise the molecular structure of each Hcluster state. While EPR spectroscopy can electronically distinguish between $[4Fe-4S]^{1+}-[Fe_p^{2+}Fe_d^{1+}]$ and $[4Fe-4S]^{2+}-[Fe_p^{1+}Fe_d^{1+}]$, the determination of the actual structure requires better suited techniques. Along this line, Lubitz and coworkers proposed the structure of H_{red}H⁺ in 2017 based on the pH dependency of this state (Section 3.6).⁷⁶ Since the secondary amine is the most basic site within the H-cluster and simultaneously the endpoint of the catalytic PTP, a structure with a protonated bridge (NH₂⁺) and a (semi-)bridging CO was justified. Later, this structure was further strengthened by IR und NRVS measurements. There, a bridging CO signal at 1810 and 1803 cm⁻¹ was found within the IR spectrum of H_{red}H⁺ and H_{sred}H⁺, respectively. The absence of a high energy μ-H band in the respective NRVS supported the assumption of a protonated amine bridge for these structures.⁷⁷ As mentioned in Section 3.4, these measurements were, however, conducted at 40-70 K, rendering a direct comparison with spectroscopic results obtained at room temperature cumbersome. IR spectroscopy at room temperature did not show a bridging or semi-bridging CO ligand. 51,72 However, in models without μ-CO ligands, the clear pH-dependent formation of H_{red} was addressed via a bridging hydride for H_{red} and H_{sred} incoming from the catalytic PTP (Fig. 20 left cycle).

Computational simulations on those structures were in overall good agreement with observed spectra. 80 Hox - Hred NRVS difference spectra revealed major differences at the [2Fe-2S]-subsite for H_{red} , whereas the [4Fe-4S]-cluster in H_{red} and Hox is identical. Again, simulations were carried out for a structure with a μ-H⁻ ligand, which agreed well with the found spectra. 80 Interestingly, a high energy band at approx. 750 cm⁻¹ for a bridging hydride was absent in these simulation and present in simulation of Birrell and coworkers.⁷⁷

The current data situation does not allow to favour one of the proposed structures for $H_{red}/H_{red}H^{\dagger}$ and $H_{sred}/H_{sred}H^{\dagger}$ in an unbiased fashion. However, the discussion on these H-cluster states indeed improved the knowledge on the working mechanism of [FeFe]-hydrogenases – at least, if in vitro measurements of isolated enzymes are considered. Recently, in vivo measurements on living E. coli cells that were genetically modified to express the hydrogenase HydA1 and artificially maturated with a synthetic precursor of the diiron site, were performed.³⁵ Here, no $[4Fe-4S]^{1+}$ - $[Fe_p^{2+}Fe_d^{1+}]$ -like state was found, instead H_{red} was enriched under 1% H₂ (99% N₂). Addition of 2 mM NaDT and acidification of the cells led to formation of $H_{ox}H$. Additional treatment with H2 afforded Hhyd. Increasing the NaDT concentration to 100 mM enabled the detection of Hhvd even at pH 8. When the pH was lowered at these conditions, a new species with approx. 15 cm⁻¹ upshifted CO bands compared to previously reported CO signals of H_{hyd} occurred. These signals were attributed to the $H_{hyd}H^{+}$ species, comprising a protonated amine bridge, which was hard to enrich in isolated HydA1 and was therefore not characterised - although postulated (Fig. 20). This state is the missing link between H_{hvd} and Hox (Fig. 21) and provides insight into the H2 formation and cleavage.35

Part B: structural and functional models of [FeFe]-hydrogenases

IV Synthesis of H-cluster models - bridge alterations, metal exchange and ligand substitution

4.1 From enzymes to biomimetic H-cluster models

The pioneering progress in the analysis of natural systems led to the identification of the crucial framework and functional properties of the active site of [FeFe]-hydrogenases. Since the detailed structural characterisation of the natural system at the end of the last century, 28,108 synthetic chemists have constantly devoted efforts towards mimicking the enzyme subsite. These mimics range from small transition metal dinuclear carbonyl clusters to elaborate artificial protein replications.31-33

However, it should herein be mentioned that [2Fe-2S]-mimics were already described in the early 20th century without the knowledge of the hydrogenases' active site but paved the way towards the fundamental synthetic strategies to assemble these cluster mimics. It is furthermore noteworthy that with the exceptional work of Rauchfuss, Darensbourg and Pickett after the elucidation of the enzyme structure, the iron sulphur chemistry underwent a renaissance and became a significant part of bioorganometallic chemistry. The tremendous modification efforts to the catalytic subsite are mainly driven to either achieve ideal catalytic efficiency approaching that of the natural enzyme or to understand the underlaying catalytic mechanism which results in a hitherto unmatched proton reduction activity of the natural system. The diiron subsite can be modified in multiple ways, including e.g. the modification of the bridging S-S linker length, exchanging the bridging sulphur-atoms with other chalcogens or pnictogens, substitution of carbonyl ligands or even the incorporation of other metals. We herein attempt to provide a comprehensive overview on the plentiful modification options.

4.2 H-cluster models with altered dithiolate bridges

4.2.1 Methanedithiolate models. The first synthesis of a diiron methanedithiolate complex was described by Seyferth et al. already in 1981 from in situ generated Li₂Fe₂S₂(CO)₆ (6) and CH2I2. This synthesis afforded the desired compound $Fe_2(\mu\text{-SCH}_2S)(CO)_6$ ((7), Fig. 22) in a yield of 25%. Notably, the reaction of CH2I2 with Fe2(SH)2(CO)6 (8) resulted in significantly improved yields of 84%. 109 In 2012, Liu et al. 110 described a facile novel synthetic route towards a -S2C=CR2 linker. Following this route, Fe₂(μ-S₂C=CHPh)(CO)₆ (9) was obtained by the reaction of Fe₃(CO)₁₂ (10) and 2-phenylethenethione as well as its tautomer (2-phenylethyne-1-thiol). The low yield (4%) of the model yet remains unexplained due

Fig. 22 Structure of exemplary methanedithiolate complexes.

to the extensive and complicated chemistry of thiols with $Fe_3(CO)_{12}$.

In that direction, $Fe_2(C_8H_{12}S_3)(CO)_6$ (11, $C_8H_{12}S_3 = 3,3$ dimethyl-4-(propan-2-ylidene)-thietane-2,2-bis(thiolate)) obtained via the reaction of tetramethyl-1,3-cyclobutanedithione with $Fe_3(CO)_{12}$. A plausible synthetic mechanism for the formation of 11 involves the rearrangement of the dithione to a beta-thiolactone followed by sulphur insertion to yield the respective thiolene, which upon reacting with Fe₃(CO)₁₂ yields 11.

Furthermore, $Fe_2(\mu-S_2C=CHC(O)C_6H_4R)(CO)_6$ (12R, R = F, OMe) with unsaturated functionalities at the thiolate linker were described by Zamora and coworkers. 111,112 While the reaction of $Fe_2(CO)_9$ (13) and $(HS)_2C = CHC(O)C_6H_4R$ in diethyl ether gave the respective saturated products (14R), a 1:1 molar ratio of Fe₃(CO)₁₂ and corresponding ligand in THF yielded the unsaturated product (12R) (Fig. 23). Upon further increasing the ligand amount, a mixture of both products was obtained. Although these models show structural similarities with the active site of the enzyme, they generally lack further investigations on their catalytic activity.

4.2.2 Ethanedithiolate (EDT) models. In the early 1960s, King et al. described the first synthesis of Fe2(edt)(CO)6 (15, Fig. 24) obtained via the reaction of Fe(CO)₅ and 1,2ethanedithiol.113 In a later attempt, the complex was also obtained by reacting Fe₂(CO)₉ with ethanedithiol/dithiocarbonates (C₂H₄S₂CO). 114,115 Subsequently, Huttner et al. reported that complexes of the type $Fe_2(S_2(CH_2)_n)(CO)_6$ (n = 2, 3) can be generally synthesised by the reaction of Fe₃(CO)₁₂ with the respective thiols under reflux conditions. These modified synthetic pathways significantly improved the yields of the respective [2Fe-2S]-complexes from 0.3% to 65% and can thus be regarded as key achievements towards a modern hydrogenase research. 116

With these synthetic possibilities at hand, various modifications have been carried out. Donovan et al. reported the modified analogues Fe₂((SCHR)₂)(CO)₆ (16R, R = CH₃, CH₂OH) wherein methyl and hydroxymethyl groups were introduced to the ethanedithiolate linker. 117 In the absence of any acid, the complexes displayed cathodic shifts in their reduction potential increasing with the number of methyl groups incorporated within the thiolate linker. When studied in the presence of e.g. 4-tert-butylphenol as a proton source, a catalytic peak was observed at approximately $-2.2 \text{ V} \text{ vs. Fc/Fc}^+$ for all complexes. Along this line, Fe₂(SCH₂CH(CH₂OH)S)(CO)₆ (17) providing an

Fig. 23 Influence of the reaction conditions on the product obtained as described by Toledano et al. 111,112

Fig. 24 Exemplary EDT models

additional hydroxyl group at the ethane bridge was synthesised to study the influence of hydrogen bonding within the protonation experiments. It was expected that the hydroxyl group can act as a proton shuttle. While Pickett and coworkers revealed a high degree of hydrogen bonding, which results in selfpolymerised cyclic hexamers, 118 the model was, however, catalytically inactive.117 In addition, the edt complex was modified with a sulfonate moiety to enable catalysis in water. Using ascorbic acid as a proton donor and [Ru(bpy)₃]²⁺ as photosensitizer, the modified edt-complex was reported to generate 88 equivalents of H₂ per catalyst equivalent. 119

Edt-complexes can likewise be obtained by reaction of Fe₂(SH)₂(CO)₆ and diketones – here, reactions e.g. with glyoxal or benzil resulted in the formation of Fe₂((SC(OH)R)₂)(CO)₆ (18R, R = H or Ph, Fig. 24). The rigid unsaturated dithiolate linkers have been broadly known to assist in the facile reduction of the [2Fe-2S]-models via delocalisation of the charge density from the iron centres through $p-\pi$ interaction.¹²⁰

Likewise, the edt-model $Fe_2(SCH_2C(S)=C=CH_2)(CO)_6$ (19, Fig. 24) was synthesised by the reaction of Li₂Fe₂S₂(CO)₆ with excess 1,4-dichloro-2-butyne. Herein, the electron withdrawing nature of the substituent, reduces the electron density at the iron centre thereby resulting in slightly milder potentials for electrochemical reduction $(-1.60 \text{ V vs.} -1.66 \text{ V of Fe}_2(\text{pdt})(\text{CO})_6$ (20)). The unsubstituted diiron buta-2,3-diene-1,2-dithiolato model was found to be capable of proton reduction of CH₃COOH with a low overpotential of −0.65 V in MeCN. ¹²¹

Since the rotated structure of the complexes bearing a bridging carbonyl group is a catalytically relevant key intermediate, studies were also conducted to estimate the influence of steric bulk at the dithiolate linker on stability of the rotated state. Here, model 21 with a rigid norbornane structure was synthesised from norbornyltrithiolane (exo-3,4,5-trithiatricyclo[5.2.1.0]decane) and Fe₂(CO)₉ and was subsequently studied using photoelectron spectroscopy in comparison to Fe₂(pdt)(CO)₆, Fe₂(bdt)(CO)₆ and 2,3-pyridinedithiolato analogues. ¹²² While the reorganization energies of the 1,2-benzenedithiolate, 2,3-pyridinedithiolate, and 1,3-propanedithiolate complexes are comparable, the norbornane model revealed the largest overall reorganization energy. However, the reorganization energies of all models are small compared to the enzymatic active site and further corroborates the importance of the secondary coordination sphere on the proton reduction at unmatched biological rates. 122

4.2.3 Propanedithiolate (PDT) models. The historical unambiguity in the exact nature of the bridgehead atoms led to great efforts in developing models of the active sites.

One example for such a model system that was thought to possess biological relevance is Fe₂(pdt)(CO)₆ (20) and numerous complexes were reported with modifications on the bridgehead carbon (Fig. 30). In this section, Fe₂(pdt)(CO)₆ models are described with focus on modified dithiolate linkers.

The PDT model was originally prepared from Fe(CO)₅ by a reaction with tetrathiacyclophane in 30% yield. Later, this method was modified, and 1,3-dithianes were employed leading to an increased yield of 42%. 124 Nowadays, the synthetic methodology of employing Fe₃(CO)₁₂ as the starting material along with propanedithiol enables excellent yields of up to 92% (Fig. 25).116

In general, Fe₂(pdt)(CO)₆ based analogues can be synthesised from the oxidative addition of cyclic disulphides to Fe₂(CO)₉ or by the reaction of the respective dithiols with Fe₃(CO)₁₂ (Fig. 25). These facile pathways usually lead to the desired complexes in high yields. Alternatively, reaction of Fe₂S₂(CO)₆ with LiEt₃BH, CF₃COOH along with a suitable dihalide compound $(R_1R_2C(CH_2X)_2)$ where X = Cl, Br) can be carried out as shown in Fig. 25. However, usually this method gives lower yields as compared to the first two pathways.

Examples of early modified Fe₂(pdt)(CO)₆ complexes date back to 1982, when Seyferth et al. investigated the reaction of $Fe_2(SH)_2(CO)_6$ with mesityl oxide and α,β -unsaturated ketones in the presence of amines (e.g. triethylamine or piperidine) to afford complex 22.125 Another early report of this class of complexes described the synthesis of a model bearing valeric acid and its C₁-functionalized derivatives. Comparably, complexes 23 to 25 were obtained upon treating Fe₂(CO)₉ with α-lipoic acid or its ester/amide derivatives in THF. 126 Also, the unsymmetrical complex 26 bearing a cyclohepta-4,6-diene unit in the bridge has been reported and was obtained upon treating Fe₂(CO)₉ with the respective trithiolane compound (i.e. 2,3,4-trithiabicyclo[4,3,1]deca-6.8-diene) in THF. 127

Likewise, as the nature of the bridging ligands exerts significant influence on the electrochemical properties of the models, their systematic study was thought to be essential. Hereby, models with longer $(-S(CH_2)_nS-)$ (n = 4-8) dithiolate linkers were reported. Prior to reaction, however, disulphide formation was performed to suppress the formation of polymeric complexes and to favour the formation of the [2Fe-2S]mimics. The required cyclic disulphides were obtained upon reacting the respective dithiols (HS(CH₂)_nSH) (n = 4 to 8) and iodine in a Et₃N solution. Upon refluxing the disulphides with Fe₃(CO)₁₂, the respective sub-site mimics 27 to 31 were obtained. It was shown that oligomerization can hardly be suppressed leading to di-, tetra- as well as hexametallic

$$Fe_{2}(CO)_{9} + S-S$$

$$OC \qquad SSS \qquad CO \qquad reflux$$

$$R_{1}R_{2} \qquad + Fe_{3}(CO)_{12}$$

$$OC \qquad CO \qquad R_{1} = R_{2} = H \quad 20$$

$$10$$

$$1. \text{ LiEt}_{3}BH$$

$$2. \text{ CF}_{3}COOH$$

$$R_{1}R_{2}$$

$$3. \qquad X$$

$$COC \qquad SSS \qquad CO$$

$$OC \qquad Fe \qquad Fe \qquad CO$$

$$OC \qquad Fe \qquad Fe \qquad CO$$

$$OC \qquad CO$$

Fig. 25 Synthetic pathways to PDT and derivatives.

complexes. Likewise, the increased length of the dithiolate linker had negligible influence on the electrochemical properties of the corresponding complexes. 128

4.2.3.1 PDT models with C2-modifications at the dithiolate linker. Most alkyl chain C₂-modifications of the dithiolate linker were - and still are - performed at the C2 position of the pdtligand. Although the general catalytic properties of the resulting model compounds cannot be altered to a significant extent, modified bridges do enable the alteration of physical properties, e.g. solubility, size and adhesive capabilities as well as the introduction of additional functional groups for the linking to surfaces or macromolecules.

Solubility is a key issue in HER-catalyst research and especially in larger scale, solvation in aqueous media is preferred for environmental reasons along with the parallel use of water as solvent and substrate. However, Fe₂(pdt)(CO)₆-like compounds comprise a bad solubility in water due to their non-polar character. Addressing this issue, the solubility of models is usually increased through CO-ligand substitution in favour of phosphines 129,130 or by encapsulating the models in a watersoluble framework (e.g. dextrins¹³¹ or micelles¹³²). The introduction of polar headgroups at the C₂ position of the dithiolate linker is a further possibility to improve the solubility of the [2Fe]_H-subsite models.

Weigand and coworkers showed that upon introducing sugar residues to the C2 position of 20 (Fig. 26) and Fe2(pd-Se)(CO)₆ (32) the biomimetic catalysts became water soluble. ¹³³ The synthesis of those compounds (33 and 34) followed the well-established route via reacting Fe₃(CO)₁₂ with the respective protected dithiol or diselenolane in tetrahydrofuran under reflux and a follow up deprotection of the sugar moiety with sodium methoxide. Interestingly, the selenium version shows an improved stability as well as activity regarding HER in aqueous media, which was explained by the increased electrondonating properties of selenium and therefore stronger

Fig. 26 Structure of PDT with modifications at the C2 position of the dithiolate linker.

 π -backdonation to the CO ligand from the iron centres, resulting in a stronger Fe-C bond. Both models provided good solubility in H₂O:MeCN (5:1) and acted as proton reducing catalysts using acetic acid or water as substrate. 133

Besides sugar residues at the pdt-bridge, more simple models comprising a hydroxy group at the bridge exist as well. This model (35) was synthesised to gain information on the influence of the hydrogen bonding network between the single complexes in solution and to mimic the natural environment of the H-cluster. 134 Using 1,3-disulfanylpropan-2-ol, a binuclear Fe₂(pdt)(CO)₆-like structure (35) was obtained, which is arranged in a helical structure in solid state forming H-bonds between the single hydroxy groups. For the longer butane linker 1,4-dithiothreitol, however, a cyclic tetranuclear complex (36) was formed, in which two dithiolate bridges coordinate two different Fe₂(CO)₆ moieties. In a follow up study, the hydroxy group in 35 was modified via two ways: (a) masked by a methyl group to investigate the influence of the oxygen atom without the hydrogen bonding network to neighbouring complexes (37) or (b) by adding an additional alkyl group to the C2 position of the pdt-bridge (38).135

Notably, such modifications do not alter the catalytic properties of the resulting cluster compared to the hydroxy derivative 35. These results once again show that derivatisation of the pdtlinker does not necessarily influence the catalytic mechanism of [FeFe]-hydrogenase models, unless pK_a and electron density are dramatically changed. A change in reactivity, however, can

be achieved by elongation of the C2-OH distance e.g. by implementing an additional methylene group in the bis-(hydroxymethyl)-functionalised Fe₂((ECH₂)₂C(CH₂OH)₂)(CO)₆ (E = S (39), Se (40)) complexes. 136 The solid state structure of those complexes also revealed a significant intermolecular H-bonding network, forming a rod-shaped cluster with tetrahedrally arranged OH groups. Notably, the reduction potential determined as $E_{44} = -1.53 \text{ V}$ and $E_{45} = -1.49 \text{ V}$ vs. Fc^{+/0}, respectively, which is significantly anodically shifted due to facilitated structural changes upon reduction compared to PDT that exhibits its averaged first reduction at $-1.66~V~\nu s.~Fc^{+/0}$ (see Section VI). Furthermore, as the resulting anionic species in solution is potentially stabilised via intermolecular hydrogen bonding from the hydroxymethyl moiety to either the Fe or the chalcogen atom, respectively. The anion was found to be capable of H₂ production from CH₃COOH via a proposed ECEC mechanism.136

Another way to modify OH-functionalized complexes is the subsequent derivatisation with a carboxylic chloride in presence of Et₃N and offers a wide range of possible alterations. Song et al. described an alternative synthesis of 35 from Li₂Fe₂. S₂(CO)₆ and 1,3-dibromo-2-propanol and further derivatised the OH-modified bridge (41 to 43R) as described in Fig. 27 to explore their influence on the catalytic properties. 137 Herein, model 42 is of special interest, as the pendant phosphine coordinates to one of the iron centres, thereby resembling a closer [2Fe-3S] H-cluster model. 138,139 However, electrocatalytic activity regarding proton reduction was yet solely reported for model 41 bearing the ketone group. 137

The structural features of a headgroup-bound ligand, which can coordinate to the iron centres as shown in complex 42, was also addressed by Pickett and coworkers. 140 A series of models of the type $Fe_2((SCH_2)_2C(CH_3)(CH_2S-p-C_6H_4X)(CO)_5)$ (X = CN (44), NO₂ (45), NH₂ (46)) and Fe₂((SCH₂)₂C(CH₃)(Y))(CO)₅ (Y = 2-pyridine (47), CH₂OH (48), CH₂NH₂ (49), CH₂SMe (50)) was subsequently synthesised. 138-140 It is worth mentioning that models 47 to 50 display a pH dependent CO-binding ("on" or "off"). The labile bridgehead substituent acts either as a chelating ligand or as a base. Upon protonation under COatmosphere the hexacarbonyl compound is generated, which can be reversed with addition of bases wherein ligand to CO displacement occurs. The influence of the coordination of the pendant thioether on the reactivity of 50 is discussed in Section 5.2.

Following organometallic advances, the interest to develop [2Fe]_H analogues displaying better catalytic behaviour or which could be strategically integrated into electrocatalytic systems gained tremendous popularity. 141-143 Thus, models with easily transformable functionalities such as carboxylic acid groups (51) were introduced to Fe₂(pdt)(CO)₆-like models by reacting $Fe_3(CO)_{12}$ with e.g. 1,2-dithiolane-4-carboxylic acid. The carboxyl group enables the functional binding of suitable amines via amide bond formation e.g. with aniline, and therefore allows the covalent attachment to amino-functionalized pyrolytic graphite electrode surfaces - interesting candidates for the design of heterogeneous electrocatalysts. 141,144 The catalytic mechanisms and potentials after attachment of the Fe₂(pdt)(CO)₆-like model to the surface, seem to be unchanged compared to a "free" complex.

4.2.3.2 Chalcogenide and pnictogenide substituted PDT models. As mentioned in the previous section, the selenium version of a sugar-substituted Fe₂(pdt)(CO)₆-like complex shows a higher proton reduction activity and improved stability in aqueous media. To this end, various Se-substituted models (32, 52 to 55) were reported, which can be obtained by refluxing Fe₃(CO)₁₂ with either 1,3-diselenocyanatopropane, a modified diselenolane or 1,3,5-triselanacyclohexane (Fig. 28). 145

In the same manner, models 56 to 58 bearing an oxetane ring were synthesized. 146 The subsequent investigation of these

Fig. 27 Representative modifications of OH-derivatized complexes. 137

Fig. 28 Synthesis of selenium modified H-cluster models.

chalcogenide substituted models regarding their proton reduction capabilities revealed a decreasing activity on moving from S- to Te-analogues owing to an increased Fe-Fe distance and hence disfavouring a bridging ligand, *e.g.* a hydride from direct protonation or a CO ligand from the so called "rotated state". ^{145,146} A further trend that can be observed upon changing sulphur to selenium or even tellurium is a decreasing reorganization energy for the reduction from a Fe^IFe^I to a Fe^IFe^O cluster, which balances the increasing electron density at the iron centres due to the sulphur exchange with stronger electron donors.

Along this line, models with additional methylene groups inside the linker were synthesised via reaction of Fe₃(CO)₁₂ and the respective 1,2-thiaselenane or 1,2-thiatellurane (59 and 60). Using these models, likewise the change of the reorganization energy upon exchange of sulphur by its heavier homologues was studied and lowered energies were observed. ¹⁴⁸

Additionally, to study the influence of the steric bulk on the reduction properties, methyl substituents (61 and 62) were introduced on the bridgehead carbon of the diselenide linker. The altered reduction behaviour along with catalytic abilities of these complexes will be discussed in Section 6.3.

In addition to the above presented synthesis pathways, the reaction of dihalides and $Fe_2E_2(CO)_6$ (E = S, Se, Te) is a further valuable approach (compare with Fig. 25). Following this synthetic scheme, $Fe_2((TeCH_2)_2CH_2)(CO)_6$ (63) was obtained from the reaction of $Fe_2Te_2(CO)_6$ and $Br(CH_2)_3Br.^{149}$

Motivated by a lower acidity of phosphines (R₂PH) as compared to the corresponding thiols, it was likewise postulated that diiron diphosphido models display an enhanced basicity of the iron centres resulting in stable terminal hydrides upon protonation. The control of the diphosphido analogues, the diphosphines (CH₂)_n(PPhH)₂ (n=2, 3) were refluxed in the presence of Fe₃(CO)₁₂ affording complexes 64 and 65. Furthermore, these models were transformed to the diphosphine substituted analogues Fe₂{(CH₂)₃(PPh)₂)}(CO)₄(κ^2 -dppv) (66a) and Fe₂{(CH₂)₃(PPh)₂)}(CO)₄(κ^2 -dppbz) (66b). Due to the increased metal basicity they were predicted to undergo

protonation at the metal centre. However, slow protonation at the metal centre resulting in a bridging hydride state was observed in low temperature experiments $(-90 \, ^{\circ}\text{C})$. For a detailed discussion of the protonation behaviour of H-cluster mimics see Section 5.3.

4.2.3.3 H-cluster models with other group 14 elements in the bridgehead position. The exchange of the C2 carbon for its heavier homologues (Fig. 29) strongly influences the properties of the resulting complexes. A report by Glass et al., for example, described the synthesis of a tin substituted hydrogenase analogue - Fe₂((SCH₂)₂SnMe₂)(CO)₆ (67). The complex was obtained from Fe₂(SH)₂(CO)₆, Me₂Sn(CH₂I)₂ and Et₃N. Later on, studies on a series of silicon modified dithiolato diiron models were described by Apfel et al. 157 The Fe₂((SCH₂)₂SiR₁R₂)(CO)₆ (68 to 70, $R_1 = R_2 = Me$, $(CH_2)_n n = 4, 5$) complexes were obtained by reacting Fe₃(CO)₁₂ and the corresponding bis(mercaptomethyl)silanes. Moreover, the Si-bridged tetranuclear model $(CO)_6Fe_2(SCH_2)_2Si(CH_2S)_2Fe_2(CO)_6$ (71) was synthesised from Si(CH₂SH)₄ and Fe₃(CO)₁₂. Due to C/Si exchange, the basicity of the sulphur centres increased resulting in a higher probability of S protonation. The group of Weigand and coworkers continued the study on such Si-substituted models and further reported a series of Fe₂((SCH₂)₂SiR)(CO)₆ models with bulky Si-bridgehead substituents (R = Si-substituted fluorene (72), xanthene (73) and thioxanthene (74)). 158,159 Furthermore, to investigate the role of bulky dithiolato linkers and their influence on redox properties, models with Ge- and Sn-containing linkers were reported. Adapting the synthetic approach, reaction of $R_2Sn(CH_2I)_2$ (R = Me, Ph) as well as $Me_2Ge(CH_2CI)_2$ and Fe₂(SLi)₂(CO)₆ resulted in the corresponding complexes $Fe_2((SCH_2)_2ER_2)(CO)_6$ (E = Sn (67, 75), R = Me, Ph; Ge (76), R = Me). In case of Sn, the cyclic tetraiiron models 77 and 78 were also obtained alongside in low yields (<9%). Notably, while for $E(CH_2S^-)_2$, (E = S, O, NR', CR₂, Si) the bridge adopts a chair/boat geometry, in case of Ge- and Sn-substituted models, the FeSCECS ring preferentially adopts an almost planar geometry indicating the deformability of these rings due to less

Fig. 29 Literature-known exchanges of the bridgehead carbon atom by heavier analogues.

torsional strain. ¹⁶⁰ These studies were further extended and models with selenium substituted Sn-bridges were reported. ¹⁶¹ $Fe_2((Se_2(CH_2)_nSnMe_2)(CO)_6 \ (n=1\ (79),\ 2\ (80))$ were obtained from $Me_2Sn(CH_2Se)_2$ or $Me_2Sn(CH_2Se)Se$ and $Fe_3(CO)_{12}$ and the desired complexes were obtained in moderate yields (20 to 30%). Herein, also the chalcogenide substitution along with C/Sn exchange causes an increased basicity of the metal core thereby facilitating its protonation.

Notably, the silicon bearing aromatic system possesses light harvesting properties. 162,163 The [FeFe]-hydrogenase model 72 with the covalently attached photosensitizer 1-silafluorene was synthesised $\it via$ the reaction of 1,1′-bis(chloromethyl)-1-silafluorene and Fe₂S₂(CO)₆. Photochemical H₂ evolution experiments were then performed in acetonitrile using trifluoroacetic acid as proton donor and triethylamine as electron donor revealing a turnover number (TON) of 29 and a turnover frequency (TOF) of 2.2 $h^{-1}.^{164}$

4.2.3.4 PDT models with secondary sphere modifications. In Section 4.2.3.1, we already presented mimics that were easily modified by amide bond formation between a carboxylic acid at the C_2 position of the dithiolate bridge and a modified amine. Along this line, [2Fe-2S]-clusters were also incorporated into larger matrices such as e.g. (bio-)polymers. Incorporating the previously highlighted mimics into a larger matrix was shown to enable its protection from undesired influences of foreign substrates and higher complex stability was anticipated. Fe₂(pdt)(CO)₆ models were immobilized on various polymers (e.g. polyacrylic acid^{132,165} and polystyrene-polyethylene glycol¹⁶⁶)

through an amide bond between **81** and amines of the proteins or a redox active group enabling a study of electron transfer processes in such systems. Contrary to those expectations, the PEG environment facilitated degradation of the iron cluster by CO loss and subsequent binding of the ether-oxygen. Likewise, a more unstable behaviour against acidic media was found caused by the surrounding ether moieties in the polymer. ¹⁶⁶

4.2.4 Azadithiolate (ADT) models. Even before the final structural elucidation, it was suggested that a secondary amine in the enzymatic cofactor, in close vicinity to the diiron site, would facilitate a low energetic pathway for proton hydride combination due to its protonation accounting for the unsurpassed catalytic efficiency.^{2,29} The importance of an amine-bridge was further supported by DFT calculations.¹⁶⁷

The synthesis of $Fe_2(adt)(CO)_6$ -like models was first reported in 2001. Here, the dilithium salt of $Fe_2S_2(CO)_6$ was obtained via reaction of $Fe_2S_2(CO)_6$ with $Li[BEt_3H]$ and afforded the N-functionalized models $Fe_2(adt^R)(CO)_6$ (R = Me (82), allyl (83), $(CH_2)_2SMe$ (84)) upon reaction with the respective bis-(chloromethyl)amine precursor bridges. ^{168,169} It was further reported that the bridges can be generated via chloromethylation of various primary amines by a successive reaction with paraformaldehyde in CH_2Cl_2 and the following addition of $SOCl_2$. This method opened up the field for easily accessible N-functionalized models of the [2Fe–2S]-subsite hydrogenases mimics.

Subsequently, Li and Rauchfuss synthesized the cofactor mimic 2 bearing a secondary amine. ¹⁷ Here, condensation of $Fe_2(SH)_2(CO)_6$, which was obtained *via* protonation of $Li_2Fe_2-S_2(CO)_6$, and urotropine ((CH₂)₆N₄) gave the desired complex in

Representative PDT models presented in this section.

Fig. 31 Synthetic pathways towards $Fe_2(adt^R)(CO)_6$ mimics. $(Cp' = MeC_5H_4)$

moderate yields of 24%. The yield could be improved to ca. 40% when using a premixed solution of (NH₄)₂CO₃ and paraformaldehyde instead of urotropine.¹⁷ As shown in Fig. 31, the reaction might proceed via the formation of intermediate Fe₂(SCH₂OH)₂(CO)₆ (85), which feasibility and reactivity was later studied by Stanley et al. 120 Another route to obtain 2 in 28% yield was the reaction of Fe₃S₂(CO)₉ (86) with ammonium carbonate ((NH₄)₂CO₃) and paraformaldehyde.¹⁷

A further method to prepare Fe₂(adt)(CO)₆ was presented by Wang et al. who reported on the synthesis of this complex using organosilicon protecting groups (Fig. 32). The group employed alkylsilylchlorides (¹Pr₃SiCl (87), Et₃SiCl (88), ^tBuMe₂SiCl (89)) along with ammonia and paraformaldehyde, which was reacted with Fe₂(SH)₂(CO)₆ resulting in the highest yield of 36% after deprotection with TFA in case of 87.170

In 2010, Rauchfuss et al. investigated a rather unusual pathway to obtain $Fe_2(adt^R)(CO)_6$ (R = Me (82), Ph (90), Bn (91)). Herein, the organotitanium complex $(MeC_5H_4)_2Ti(adt^R)$ (92R) was used as an azadithiolate transfer agent to [Fe(bda)-(CO)₃] (bda = benzylideneacetone) giving Fe₂(adt^{Ph})(CO)₆ in 42% yield.171

In 2015, a synthesis for isotope-labelled ⁵⁷Fe₂(adt)(CO)₆ starting from ⁵⁷FeBr₂ was reported (Fig. 33).⁷¹ Analogous to the synthesis of **1** reported by Hieber, ¹⁷² ⁵⁷**1** is formed from the reaction of $[H^{57}Fe(CO)_4]^{-}$ (5793) and elemental sulphur. Since the established routes towards the native cofactor are based on iron carbonyl chemistry, and labelling of Fe(CO)₅ with ⁵⁷Fe is challenging on laboratory scale, this new route was developed to avoid Fe(CO)₅ (or derivatives thereof) as starting material. This allowed for explicit spectroscopic investigations by NRVS

Fig. 32 Alternative synthesis of complex 2.

Fig. 33 Synthetic pathway to ⁵⁷Fe₂(adt)(CO)₆.

and Mössbauer of [2⁵⁷Fe]_H first in its H_{ox}-CO state and later in H_{ox} , H_{hvd} and H_{ox} - O_2 as well (see Section III). 64,71

Extending the chalcogenide exchange to the adt models as well, the enzyme cofactor was synthesised with selenium. The straightforward synthetic route comprises of coupling of carbamate protected amine with Li₂Fe₂Se₂(CO)₆ (94) followed by deprotection with BF₃ and Me₂S (Fig. 34).^{32,173} Deprotection of 95 afforded the target product 96 in 20% yield, which was further subjected to ligand exchange resulting in [Fe₂(adSe)- $(CO)_4(CN)_2$ ²⁻ (4). This complex could also be embedded into apo-CrHydA1 and apo-CpI (see Section 2.2).³²

4.2.4.1 N-Alkyl modified ADT models. The conceptually most obvious modification of the amine-linker is a simple alkyl chain, which is, however, not much reported in literature. 17,168,174-176 The shortest version, model 82 bearing a methyl moiety, can be synthesized according Fig. 31 or by saltelimination using Li₂Fe₂S₂(CO)₆ and methylbis(chloromethyl)amine. 17,168 The methyl group resides either in axial or equatorial position, which is dependent on a balance between the anomeric effect (favours an axial position) and steric repulsions between the methyl group and the carbonyl ligands (favours an equatorial positions). 168 Compared to Fe₂(adt)(CO)₆, the methyl substitution changes the catalytic properties in two ways: (1) the inductive effect of the methyl group increases the electron density and therefore the basicity of the amine bridge and (2) the electron density of the iron

centres is increased via hyperconjugation of the N_{1p} and C-S σ^* orbitals. This changed electron density is, however, not visible by a shift of the CO-frequencies within the IR spectra, but results in a shift of the first reduction potential from -1.58 V for NH (2) to -1.72 V for the NMe derivative (82) (vs. Fc/Fc⁺). As a result, the methyl substitution allows for the use of less acidic proton sources during the proton reduction while the potential that has to be applied to reduce the system is more negative. 17,168,170,176

Along this line, $Fe_2(adt^R)(CO)_6$ (97, R = $(CH_2)_2NHTs$) was synthesised but amine deprotection was unsuccessful, thereby restricting its further application. In addition, the tetranuclear model 98 was reported containing two linked adt-units. 177

Lengthening the alkyl chain substituents increases the steric bulk around the metal centres and is advantageous in terms of mimicking specific H-cluster redox states. Modifications with longer alkyl chains range from simple ethyl groups to more complex cyclic alkyls. 17,120,174,176,178–185 The synthetic protocols are similar to those for $Fe_2(adt^R)(CO)_6$ (R = H, Me). Both strategies, condensation of the respective amine with (para)formaldehyde and the following reaction with Fe₂(SH)₂(CO)₆ or the salt-elimination method can be regularly found in literature. The electron donor abilities of the alkyl-moieties increase from ethyl to isopropyl/sec-butyl to tert-butyl, which is reflected by a more cathodic potential, decreasing average CO-frequencies, as well as an increasing Fe-Fe bond distance. 176,182 However, the differences in electronic parameters between

Fig. 34 Synthetic pathway to Fe₂(adSe)(CO)₆ (96).

OC SS CO BH₃THF OC SS CO
$$\frac{1}{100}$$
 CO $\frac{1}{100}$ CO $\frac{1}{100}$ CO $\frac{1}{100}$ R = Me $\frac{1}{100}$ R =

Fig. 35 Binding of BH₃ to the Fe centre

the respective alkyl chain modifications are almost nonsignificant. 179-183,186

To shed light into the protonation behaviour of the active site, the acid base chemistry between the adt moiety and BH₃ was investigated with model system 2 and 82.175 Treating Fe₂(adt^{Me})(CO)₆ with one equivalent of BH₃·THF, wherein the N-coordinating BH₃ group of model 99 binds to the Fe upon decarbonylation afforded complex 100 (Fig. 35). This study served as an illustration for the analogous binding of H2 to the iron centre in the enzyme. 175

4.2.4.2 N-Modification of ADT models by esters and amides. In Section 4.2.3, we reported on modifications of Fe₂(pdt)(CO)₆like complexes with hydroxy- or carboxylic acid-functional groups via esterification reactions or formation of amides.

The same methodology can be applied to Fe₂(adt)(CO)₆-like structures as well. These functional and structural models have been extensively studied to reveal the mechanism of the enzyme.^{187–189}

Song et al. as well as Sun and coworkers reported on the functionalization of hydroxy-modified Fe₂(adt^R)(CO)₆ (101, R = CH₂CH₂OH) to various N-modified complexes (Fig. 36). 186,190 As observed for hydroxy-modified pdt-models, 134 the hydroxy group of 101 forms an intermolecular H-bond network. 186 Via addition of derivatised carboxylic chlorides, aromatic groups (102 to 105) can be added to the bridge. In addition, a terephthalic acid bridged dimer (106) and a thioacetate derivate (107) have been reported. 186,190 However, as also observed for pdt-models, the catalytic properties of these models cannot be altered by changing the substituents at the nitrogen atom. 190

Fig. 36 Towards the syntheses of N-alkanol modified models.

Extending this concept to adSe derivatives, Gao et al. reported a series of double as well as triple cluster cores by linking multiple molecules of $Fe_2(SeCH_2)_2NR(CO)_6$ (R = (CH₂)₂OH) (108). ¹⁷³

Instead of hydroxy alkyl-modified amines, the introduction of carboxylic acids was performed with the same overall aim. Fe₂((SCH₂OH)₂(CO)₆ was reacted with the corresponding amino acids ((2-aminomethyl)benzoic acid, 2- or 4-aminobutyric acid) and afforded the desired hexacarbonyl products 109 to 111. 191

The group of Song reported an additional modification scheme with diverse models bearing N-acyl functionalities (112 to 116) (Fig. 37). 192 For example, model 116 was obtained upon reaction of 2 with 2-chloroacetic acid or chloroacetyl chloride and subsequent treatment with potassium thioacetate. Due the electron withdrawing substituents, these complexes display first reduction potentials in the range of -1.49 V to $-1.54 \text{ V} \text{ vs. Fc}^{+/0}$, which is milder than for unmodified complex 2 (-1.58 V). The Likewise, these models serve as a template for designing systems suitable for photocatalytic studies by modifying the acyl group. 192

4.2.4.3 N-Aryl modified ADT models. Subsequently, an ADT model series of substituted N-phenyl complexes (117 to 121) was established. 193 Crystal structures of the *ortho*-substituted models show sp³-hybridisation of the bridgehead nitrogen, while the para-substituted models 119 and 120 display a rather sp²-behavior at the nitrogen. Due to the steric influence of

the substituents, the lone pair of the nitrogen is unable to delocalise into the aromatic ring and hence these substituted models are capable of conducting proton reduction at near neutral pH 5.5 with low overpotentials in aerobic conditions. On the other hand, ortho unsubstituted models with electron withdrawing substituents (Br and NO₂) at the para position, required harsher acidic conditions for proton reduction due to the diminished basic nature of the nitrogen. 193

In view of developing the ADT models, the nitro functionalized model 120 model was established by treating Fe₂S₂(CO)₆ with N,N-bis(chloromethyl)-4-nitroaniline followed by reduction with Pd/H₂. This procedure yields the corresponding amine derivative $Fe_2((adt^R)(CO)_6 (122, R = p-C_6H_4NH_2).^{194})$ Additionally, condensation of N,N-bis(chloromethyl)-p-methoxyaniline with $\text{Li}_2\text{Fe}_2\text{S}_2(\text{CO})_6$ afforded $\text{Fe}_2((\text{adt}^R)(\text{CO})_6 (123, R = p\text{-}C_6\text{H}_4\text{OMe}).^{195}$ The different electronic effects of the ring substituents are evident in the reductive behaviour of these models. Due to the electron withdrawing nature, the nitro-substituted model 120 is reduced at more positive potential as compared to the amino 122 and the methoxy derivative 123 ($E_{120} = -1.42 \text{ V} \text{ vs. } E_{122} = -1.56 \text{ V}$ and $E_{123} = -1.61 \text{ V}$). ^{194,195} As these previous studies on aromatic substituents suggested a decreased reduction potential, Jiang et al. introduced furan, thiophene and pyridine substituents on the Fe₂(adt^R)(CO)₆ models (124 to 126). Herein, the electronic interactions between the heterocycles and the metal centre via linking C, N, S atoms, influences the redox behaviour.

Fig. 37 Syntheses of N-acyl modified models.

Additionally, bromine was introduced at the thiophene ring in 127 to facilitate functionalization. This bromothiophene model is catalytically active with $HClO_4$ at a potential of $-1.09~V~vs.~Fc/Fc^+$, which is significantly lower as reported for other ADT models. ¹⁹⁶

Extending the study to ADSe mimics, complexes with *N*-aryl diselenide bridges were synthesised (**128** to **130**). Introduction of different substituents (CH₃, NO₂, H) at the *para* positions of the aryl ring aimed at studying the inductive effects. Electrochemical results were in accordance with the trends observed for sulphur bearing models, *i.e.* the nitro substituted complex was more easily reduced than the alkyl substituted complex. Also, crystallographic studies show that sulphur to selenium replacement is responsible for slight elongation of the Fe–Fe bond. ¹⁹⁸

In a more recent approach, the [2Fe–2S]-cluster was attached to a variety of molecules such as nucleosides, redox active fragments (ferrocene and ruthenocene), and luminescent markers (boron-dipyrromethenes–BODIPYs) by introducing an azide functionality (131) and subsequent Cu-catalysed Huisgen cycloaddition (Click-reaction) between terminal alkynes and azides. Advantages of this strategy are high tolerance towards sensitive substrates and a broad range of various functional groups. It was further shown that the resulting triazole rings from the click-reaction can be protonated by strong acids such as $\rm H_2SO_4$ and therefore serve as model for the native adt bridge. 199

4.2.4.4 N/P exchange in ADT models. Even though phosphorus is the heavier analogue of nitrogen, little effort has been conducted to establish active site mimics of [FeFe]-hydrogenases bearing phosphorus in the linker (Fig. 38). A preceding attempt to introduce a tertiary phosphine bridge resulted in ligand substitution in complex 132R (R = Ph, CH₂Fc) due to the high nucleophilicity of the phosphine. Therefore, models 133R wherein phosphorus is incorporated into the bridging position were synthesized from $\text{Li}_2\text{Fe}_2\text{S}_2(\text{CO})_6$ and phosphine oxides O=P(R)(CH₂Cl)₂ (R = Ph, OEt) to avoid interaction of the lone pair phosphorus with the iron centre. Furthermore, this O=P functionality was identified as the protonation site in this model system. Recently, a new strategy to synthesize complexes wherein the phosphorus occupies the bridgehead position

was reported. There, $Fe_3(CO)_{12}$ and the respective dithiols $O=PR(CH_2SH)_2$ (R = OEt, OMe, OPh, OH, Me) were reacted in THF at room temperature to obtain the desired compounds **133R** in moderate yields of approx. 40%.

Another aza-diphosphido model **134** was synthesized *via* deprotonation of a $Fe_2(PPhH)_2(CO)_6$ precursor with MeLi and further incorporation of a $(Cl(CH)_2)_2NR$ (R = CH_2CH_2OMe) linker (Fig. 39). Notably, protonation of these aza-diphosphido analogues occurs exclusively at the amine bridgehead.²⁰³

4.2.5 Oxadithiolate (ODT) models. Based on the first crystal structures of [FeFe]-hydrogenases, $[Fe_2(\text{odt})(\text{CO})_4(\text{CN})_2]^{2-}$ (135) was one of the mimics considered to be the putative active site, since the central oxygen atom has the same number of electrons as the actual active sites head group NH. ^{3,14} Its precursor $Fe_2(\text{odt})(\text{CO})_6$ (136) was originally prepared *via* acidification of $Fe_2(\text{SCH}_2\text{OH})_2(\text{CO})_6$, and later from $\text{Li}_2\text{Fe}_2\text{S}_2(\text{CO})_6$ and $(\text{ClCH}_2)_2\text{O}$ *via* salt-elimination (Fig. 40). ^{17,204} Compared to $Fe_2(\text{adt})(\text{CO})_6$, the average CO frequency of the oxadithiolate derivate is slightly upshifted from 2018 to 2024 cm⁻¹ in hexane indicating the influence of the electron withdrawing oxygen group on the overall electronic structure of the diiron cluster. ¹⁷ However, this electron withdrawing behaviour is not reflected in the respective first reduction potential, which is identical for both complexes at $-1.58 \text{ V vs. Fc/Fc}^+$.

For the oxadithiolate models, sulphur to heavier homologue exchange was carried out and $Fe_2(odSe)(CO)_6$ (137) and $Fe_2(odTe)(CO)_6$ (138) complexes were reported. While 137 was obtained in 45% yield from addition of (HSeCH₂)₂O to $Fe_3(CO)_{12}$,²⁰⁵ 138 was synthesised in 21% yield from $Fe_2Te_2(CO)_6$ and $(ClCH_2)_2O$.²⁰⁶

As already reported for the model counterparts 2 and 20, substitution of CO with more electron donating ligands such as CN⁻, PR₃, NHCs and Cp(CO)₂FeSPh was carried out to further influence the electron density at the iron centres. ^{181,185,207–210} Additionally, tetranuclear models wherein the modified ligand system (dppf, (Ph₂PCH₂)₂NCH₂)₂, (Ph₂PCH₂CH₂OCH₂)₂ and 1,4-(CN)₂C₆H₄) connect two [2Fe–2S]-cores have been synthesised and crystallographically elucidated. ²⁰⁴

4.2.5.1 O/S exchange in ODT models. In contrast to the little explored N-P exchange in adt, incorporation of the heavier

Fig. 38 Synthetic pathways to phosphorus substituted subsite models.

Fig. 39 ADT derived complexes.

Fig. 40 Synthesis routes towards Fe₂(odt)(CO)₆.

homologue sulphur is much more common in odt complexes. According to IR spectroscopy, both models do not show significant electronic differences although sulphur is less electron withdrawing than oxygen. The first reduction potential of $Fe_2(sdt)(CO)_6$ (139), however, is shifted from $E_{146} = -1.58$ V to $E_{150} = -1.51$ V vs. Fc/Fc^+ , showing a more severe change in the electron density at the iron centre. 204,211

The synthesis of **139** was reported initially by Song *et al.* in 2007, using 1,2,4-trithiolane as starting material, which was reacted with $Fe_3(CO)_{12}$ in refluxing THF to yield the desired complex in 42% yield.²¹¹ In parallel, Windhager *et al.* reported various model complexes (**140** to **142**) containing S-substituted bridge structures obtained by reacting $Fe_2(CO)_9$ with different sulphur substituted heterocycles. It was observed that larger

Fig. 41 Syntheses of multi-metallic complexes from sulphur heterocycles

Fig. 42 Synthetic pathways to novel sdt complexes from Fe₂(CO)₉

heterocycles yield trinuclear clusters (141 and 142). The flexibility of the linker strongly affects the structure of the resulting models, e.g. the 7- and 8-membered heterocycles yield triiron clusters wherein the thioether moieties coordinate to one of the Fe centres each. Applying the 9-membered thio compounds affords the diiron complex 140 (Fig. 41).²¹²

The treatment of diiron thiadithiolate with Cp(CO)₂Fe(BF₄), prepared in situ from Cp(CO)₂FeI and AgBF₄, led to cationic model 143a (Fig. 43). This model corroborates the ability of

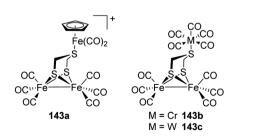


Fig. 43 Trimetallic subsite models with coordination of the bridgehead sulphur to metal centres.

the bridgehead sulphur atom to likewise coordinate to metal centres.211 Taking advantage of this coordinating capability, the parent Fe₂(sdt)(CO)₆ model was reacted with M(CO)₅(THF) (M = Cr, W) (prepared in situ by photolysis of $M(CO)_6$ in THF) and afforded the multi-metallic complexes 143b and 143c.²¹³ These complexes were designed to profit from combined redox properties of the different metals involved; however, they were unstable under electrochemical conditions and hence could not be studied for their catalytic properties.213

Additionally, the introduction of substituted 1,2,4-trithiolanes leads to various di-, tri- and tetranuclear models and is exemplarily shown for the diadamantyl-substituted trithiolane in Fig. 42. The synthesis proceeds through an oxidative addition of the disulphide bond to Fe₂(CO)₉ yielding models 144 to 146.²¹⁴

4.2.5.2 Oxidation of SDT-like models. The Fe₂(sdt)(CO)₆ models were likewise studied for their chemical oxidation response, as it was reasoned that the activity of [FeFe]-hydrogenases is hampered under aerobic conditions due to oxidation of the cofactor. 215 Sulphur-oxidation was achieved through reaction of varying equivalents of dimethyldioxirane (DMD)

Fig. 44 Chemical oxidation of thiadithiolate models-resulting various oxidised complexes

with SDT and derived precursors, yielding S-oxidised models (Fig. 44A). 216 These oxidation investigations were carried out on models 139, 147 and 148 with different substituents on the methylene carbons (R = H, Me, 1/2Cy). An increasing susceptibility for oxidation products with decreasing steric bulk of the substituents was observed. Also, the triiron model 141 was subjected to oxidizing conditions yielding the oxidized thiolate and thioether metal cluster models 1410a and 1410b (Fig. 44B).²¹⁶ Such oxidations were also reported for the Fe₂(edt)(CO)₆ models using *m*-chloroperbenzoic acid.²¹⁷ Chemical oxidation of $Fe_2(pdt)(CO)_{(6-n)}L_n$ (L = CO, PPh₃, PMe₃) also shed light into the site specificity of the oxygenation. DFT calculations suggested that oxidation of the Fe-Fe bond is thermodynamically favoured resulting in μ-oxo species. Contrastingly, the experimental studies revealed oxygenation at the dithiolate sulphur resulting in S-oxygenate products which were crystallographically ellucidated.215 Recently, Berggren, Hammerström and coworkers conducted oxidative degradation studies on Fe2(adt)(CO)6 and Fe₂(pdt)(CO)₆ complexes.²¹⁸ They showed that the interaction of Fe₂(adt)(CO)₆ with molecular oxygen in presence of chemical reductants leads to a transient degenerated state. Although Fe₂(pdt)(CO)₆ showed similar oxygen reactivity as compared with the above-mentioned complexes, the reaction speed was slow. The experiments thus highlight the importance of the secondary sphere on oxidative degradation pathways. The final steps of the oxidative cofactor degradation, however, are still unknown and further experiments are required to pinpoint this important aspect. Likewise, the influence of the cyanides is not understood yet.

4.2.6 Benzenedithiolate (BDT) models. The unsubstituted $Fe_2(bdt)(CO)_6$ (149, bdt = 1,2-benzenedithiolate) model can be synthesized from Fe₂(CO)₉ and 1,2-benzenedithiol. 219,220 Although it was at first not intended as a [FeFe]-hydrogenase mimic, the synthesized complex had significant similarities with the enzyme's active site. The IR spectrum in hexane shows only slightly shifted CO bands of 2079, 2044, 2006, 1967 and 1958 cm⁻¹ compared to 2 which shows its respective bands at 2076, 2036, 2008, 1989 and 1979 cm⁻¹. This shift of the CO-frequencies is indicative for a reduced electron density at the iron core of the 149 and manifests also in the its redox potentials. According to Capon et al., 149 undergoes a 2ereduction at -1.35 V vs. Fc/Fc⁺, which clearly distinguishes this model from all other [FeFe]-hydrogenase models.²²¹ These remarkable redox properties stem from an interaction between the sulphur and benzene p_{π} orbitals that balance the electron count on the iron centres and reduce the energy changes upon reduction of the complex.²²²

These factors stimulated synthetic efforts and led to dedicated research to elucidate the influence of rigid and aromatic bridges on reactivity and electrochemical properties of such [FeFe]-hydrogenase model compounds. Strikingly, **149** was also reported to catalyse the reduction of CO₂ to formate. ²²³

A recent review²²⁴ on [FeFe]-hydrogenase mimics with aryldithiolate ligands covers many aspects of the design and application of these complexes as electro- or photocatalysts. Moreover, also monometallic bdt complexes are discussed, while we strictly focused on complexes with a [2Fe–2S]-core.

The electron withdrawing effect of the benzene moiety in Fe₂(bdt)(CO)₆ was the starting point of several additional studies, aiming at introducing further electron withdrawing groups to shift the reduction potential to more anodic potentials. Hence, the $Fe_2(S_2C_6H_{4-x}Cl_x)(CO)_6$ type complexes 150 to 152 (X = 2 to 4) were investigated in-depth and compared with $Fe_2(bdt^{4Me})(CO)_6$ (153, $bdt^{4Me} = 4$ -methylbenzene-1,2dithiolate). 222,225-227 While the methyl group causes a +I effect, the chlorines possess a strong -I effect.²²⁶ As expected, 152 shows a strongly shifted CO IR pattern (approx. +10 cm⁻¹) compared to the methyl-substituted derivative. This observation reveals a decreased electron density at the diiron centre. Likewise, the reduction potential shifts towards more anodic values from -1.34 V (153) to reach -1.13 V (152). 226 Other substituents at the benzene ring, e.g. hydroxy-moieties, or electron deficient heteroaromatic rings, e.g. pyrazine, were also tested and shown to facilitate the reduction of the diiron centre. 228,229

Models 154R were synthesized via reaction of substituted 1,4-benzoquinones with Fe₂(SH)₂(CO)₆ in the presence of piperidine (Fig. 45).²³⁰ These models were expected to undergo facile reduction, as it was proposed that hydroxyl substituents would aid in the stabilization of the accumulated negative charge on sulphur upon Fe-S bond cleavage. Electrocatalytic investigations of these analogues revealed that these models conduct catalysis at 44 mV lower overpotentials than Fe₂(bdt)(CO)₆. However, these hydroquinone models were also found to be less active for proton reduction from weak acids due to internal hydrogen bonding.²²⁸ The hydroquinone was further functionalised with pyridine carboxylic acid chloride to introduce a basic proton relay²³¹ or with ferrocenoyl chloride to afford 155. This model exhibits three isomers depending on the orientation of the Fc moieties. 232 The ferrocene unit, however, did not possess any interaction with the diiron core.

Another option to alter the properties of the hydroquinones 154R was demonstrated by the oxidation with DDQ (2,3dichloro-5,6-dicyano-1,4-benzoquinone). The respective models 156R possessing non-innocent quinones as ligands were compared to models 157 (1,4-naphthoguinone) and 158 (1,4anthraguinone) with increased π -systems.²³⁰ It was anticipated that the extended π -systems would favour the electron transfer to the [2Fe-2S]-centres and further anodically shift the reduction potential. However, protonation occurs at the quinone oxygen rather than the metal centre and consequently no H₂ evolution was observed.²³⁰

Similarly, functionalization of the carboxylic acid in the cbdt ligand of complex 159 allows for the introduction of a phosphine ligand tethered to the bridging dithiol, exemplarily shown in complex 160 (Fig. 46). This linkage was shown to increase the rotation barrier of the rotation of the respective Fe(L)₃ fragment. Thereby, kinetically labile, terminal hydrides were proposed to be stabilised.²³³

The effect of increased π -systems, as performed in case of 1,4-quionones, was also investigated for N-heterocycles 161 (quinoxaline-6,7-dithiol), 162 (2,3-diphenyl-6,7-quinoxaline dithiol) and 163 (2,1,3-benzothiadiazole-5,6-dithiol) (Fig. 47). These heterocycles withdraw electron density from the metal centres and thereby cause an anodic shift of the reduction potentials ($E_{161} = -1.23 \text{ V}$, $E_{162} = -1.24 \text{ V}$, $E_{163} = -1.25 \text{ V}$ vs. $E_{149} = -1.33 \text{ V vs. Fc/Fc}^+$). Despite related structures, the models 162 and 163 follow different catalytic mechanisms. Due to the electron delocalisation within 161 caused by the additional S-atom, the nitrogen is unable to act as basic site and hence the complex follows an EC mechanism during proton reduction. Due to the absence of such an electron delocalization,

Fig. 46 Complexes 159 and 160

Fig. 45 Functionalized hydroquinone models 154R to 156

Fig. 47 Multi-heteroaryl containing mimics.

Fig. 48 Multimetallic bdt-derived complexes 164 to 166.

protonation at the ring nitrogen of ${\bf 162}$ occurs and leads to a CEEC mechanism. 234

Chen *et al.* designed tetra- and hexametallic clusters (**164** to **166**, Fig. 48) that undergo two or three consecutive two-electron reduction reactions and can act as stable multi-electron relays. The multi-electron reduction steps take place in the range of -1.33 to -1.81 V νs . Fc/Fc⁺. Such consecutive reductions are relevant to natural systems, hence these models were suggested to be valuable templates to design robust catalyst systems. ^{235,236} Further experiments to investigate the potential of such models are, however, still missing.

4.2.6.1 BDT models with secondary sphere modifications. To improve the water solubility and to overcome the oxygen sensitivity and high overpotential the hydroquinone 167 was modified to afford the bis(2-bromo-2-methylpropionate) species 168 (Fig. 49). This complex was then co-polymerized with methyl methacrylate (MMA) and 2-(dimethylamino)ethyl methacrylate (DMAEMA) via atom transfer radical polymerization (ATRP). Here, The flexible –NMe₂ side chains at the DMAEMA monomer,

along with improving the water solubility and stability of the model, also enhance the complexes' catalytic activity and overcome the aforementioned challenges of water solubility and oxygen-sensitivity.²³⁷

Another polymerization technique to produce metallopolymers is the copper-catalysed click-reaction between azides and alkynes (Fig. 50).²³⁸ Depending on the azide used, the general properties of the obtained polymer can be adjusted. Thus, polymer **169Pb** bearing an acetate-substituted backbone, reveals an improved catalytic activity compared to complexes **169Pa** and **169Pc**. The different activity can be explained by enhanced proton shuttle properties of the acetate in **169Pb**.²³⁸ This example shows, that in complex systems not only the catalyst itself is decisive for the overall HER performance, but also the network in which the catalyst is embedded.

4.2.7 Other arenedithiolate models. The different binding motif created by the naphthalene moiety as compared to $Fe_2(bdt)(CO)_6$ derivatives causes an increased S–S distance and a larger S–Fe–S angle (*e.g.* 84.1° in the anthracene derivate νs . 80.7° in bdt). Regarding the electron density at the diiron

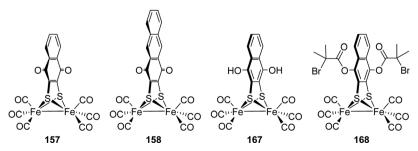


Fig. 49 Representative multiaryl diiron complexes.

Fig. 50 Incorporation of bdt-derived complexes into polymers.

site, the naphthalene moiety in $Fe_2(1,8-S_2C_{10}H_6)(CO)_6$ (170) is less electron-withdrawing compared to its benzene congener. The naphthalene derivative shows CO-frequencies that are shifted by 5 cm⁻¹ to lower wavenumbers as compared to Fe₂(bdt)(CO)₆. ^{220,239} Likewise, the reduction potential of **170** is 40 mV more cathodic. 221,240 Furthermore, naphthalene bridged [2Fe-2S]-mimics exhibit a reversible reduction and stabilise the monoanionic intermediate species obtained upon the first reduction and triggered considerable interest. 239-247 The facile synthetic route, reacting Fe₃(CO)₁₂ and naphthalene 1,8-disulphide, allows for easy manipulation of the naphthalene moiety and a series of modified naphthalene 1,8disulphide models with chloride, 239 alkyl, 239,240 alkoxide, 240 amide and imide substituents were reported. 244,248,249 As already observed for the bdt model, the chloride substituted 171 shows a anodically shifted but irreversible reduction, which was explained with the subsequent loss of chloride. 239 Along this line, electrondonating groups such as OMe and ^tBu shift the reduction potential cathodically. 239,240

Besides iron-sulphur clusters, iron-selenium clusters were reported with a naphthalene bridge as well. Figliola et al. reported an extensive study of such naphthalene and phenanthrene modified dichalcogenide models (170, 172 to 175, Fig. 51).²⁴⁰ The selenium substituted models displayed catalysis at lower overpotentials as compared to their sulphur congeners. 239,250 Likewise, the synthesis and investigation of series comprising naphthalene modified systems with an imide

Fig. 51 1,8-Naphthalenedithiolate bridged derivatives.

functionality in para-position were reported by Weigand and coworkers (176a-c and 177a-c, Fig. 53).244 The modifications increased the stability of the reduced monoanion. Increasing the aromatic system by a further naphthalene moiety resulted in the formation of perylene monoamide-bridged hydrogenase models 178a-c. 247 These analogues show an anodically shifted reduction potential due to the increased π -system. The first reduction potentials (Fe^IFe^I → Fe^IFe⁰) of these species are in the range -0.99 V to -1.04 V and the shift can be attributed to an increased electron withdrawing nature of the perylene linker and an enhanced stability of the reduced species.247

Another increased π -system was reported by Topf et al. who designed the acenaphthylene-based complex 179 (Fig. 52) which acts as redox relay for accelerated electron transfers to

Fig. 52 Synthesis of the BIAN-R model 179.

the diiron site.²⁴¹ By modification of the varying linker substituents, the bis(arylimino)acenaphthene (BIAN-R) models allow for tuneable electronic properties and solubility.

The robust naphthalene derivatives of the diiron dithiolates can be considered a major example of photocatalytically active mimics. Herein, the functionalization of the naphthalene ring with e.g. imides (180) or amines (181) provides a useful method to attach a photosensitizer. $^{248,251-253}$

Instead of increasing the naphthalene π -system, a change to an intrinsically larger aromatic system such as phenanthrene likewise enables the modification of the electronic properties of the [FeFe]-hydrogenase models. Thus, oxidative addition of phenanthro[4,5-cde][1,2]dithiin to Fe₂(CO)₉ gave the phenanthrene-bridged model **182** (Fig. 53). Compared to analogous naphthalene counterparts, the reduction potential was found to be more anodic due to the larger electron withdrawing ability of the dithiolate linker along with better stabilization of the anionic species. 250

4.3 H-cluster models lacking the dithiolate bridge

Already in 1928, Reihlen *et al.* first described the synthesis of $Fe_2(SEt)_2(CO)_6$ (183) without any knowledge on [FeFe]-hydrogenases.²⁵⁴,²⁵⁵ Also, in late 1930s Hieber *et al.* reported on the complex of the type $Fe_2(SR)_2(CO)_6$ (R = Ph (184), Et (183)) by refluxing $Fe_2(CO)_9$ and the respective dithiol.²⁵⁶,²⁵⁷ Further,

the synthetic organometallic chemistry flourished and lead to various diiron($_1$) dithiolate complexes starting from Fe₂S₂(CO)₆ (1).²⁵⁸ As illustrated in Fig. 54, 1 can be modified to obtain a variety of derivatives.

These complexes are commonly low spin, diamagnetic complexes which exist in three isomeric forms *aa*, *ae* and *ee* (Fig. 55) depending on the orientation of the R substituents on sulphur. Contrary to the chemistry displayed in the previous sections, the chelating thiolates lack such steric freedom and hence exist in the *aa* form.

Owing to the significant role of the cysteinyl ligands in both [NiFe]-hydrogenases and [FeFe]-hydrogenases, the introduction of cysteine as a bridging thiolate was pursued. However, unlike most other derivatives, this model could not be obtained through refluxing cysteine (or its methyl ester) with Fe₃(CO)₁₂. Employing the Boc-protected methyl ester of cysteine afforded the target product **185**. When **185** was refluxed in MeOH or in toluene containing CH₃COOH, intramolecular cyclization occurred leading to the EDT-like model **186** (Fig. 56).²⁵⁹

Furthermore, **185** was attached to α -helical peptides resulting in a prototype to replicate the second coordination sphere of the active site. Furthermore, a ferrocene was incorporated and linked the two cysteinyl arms in **187** (Fig. 57). ²⁶¹

Since H-bonding interactions between NH and S are crucial for electron transfer processes in the metalloenzymes such as

Fig. 53 Different naphthalene-derived model complexes.

Fig. 54 Modification possibilities of 1

Isomers of unbridged Fe₂(SR)₂(CO)₆ complexes

Fig. 56 Intramolecular cyclization of 185 resulting in 186.

rubredoxin²⁶² and ferredoxins²⁶³ and also influence their redox behaviour, complexes 188R (R = CH_3 , CF_3 , C_6H_5 , p- C_6H_4F) were synthesised revealing NH···S interactions. This interaction decreased the electron donating capability of the S atom and is visible by an anodic potential shift of 370 and 470 mV as compared to Fe₂(pdt)(CO)₆ and Fe₂(SPh)₂(CO)₆. ²⁶⁴

As described earlier, complexes with aromatic dithiolate linkers conduct proton reduction at more positive potentials. Therefore, the models 189R (R = o-OMe, p-OMe, p-Cl) were synthesized with varying ligand substitution patterns and showing varying electron donating capacities (see Section VI).²⁶⁵

In addition, models 190 to 192 were synthesized comprising pendant pyridine and quinoline groups and were supposed to facilitate an internal proton transfer. Notably, the N-heterocyclic groups were shown to dynamically coordinate to an iron atom. When a solution of these models is subjected to CO, the Fe-N bond is cleaved giving the hexacarbonyl complexes.²⁶⁶

4.4 Metal exchange in H-cluster models

Although numerous synthetic models of the active site have been synthesized and studied, less efforts have been directed towards development of mimics with different transition metals.

Fig. 57 Representative models lacking a dithiolate linker

Indeed, inspiration to realize this goal can be taken from natural systems, wherein a mixed-metal hydrogenase, i.e. the [NiFe]hydrogenase exists. Some synthetic studies described organometallic clusters with structural similarities to the active site of [FeFe]-hydrogenases, which will be discussed in this section.

4.4.1 Incorporation of group 8 elements. Organometallic complexes of heavier analogues of iron i.e. ruthenium and osmium have been reported by Cabeza et al. in 1998.²²⁰ In that work, the synthesis of bimetallic organometallic centres bearing the aromatic benzenedithiolate linker (193 and 194, Fig. 58) is described. These complexes can be synthesised by refluxing the respective metal carbonyls (Ru₃/Os₃(CO)₁₂) with 1,2-benzenedithiol in toluene.192

Subsequently, Ru models with pdt (195) and adt (196) bridges were reported (Fig. 58). Thereby, RuCl₃·nH₂O was carbonylated and in situ reacted with the dithiol as well as zinc. 33,267 In the case of adt, the secondary amine was protected

Fig. 58 Iron substitution by heavier homologues.

with a carbamate (CBz) protecting group (Fig. 59) when introduced as a dithiol (197).33 In 2015, Wu et al.268 reported the synthesis of the diruthenium complex via reaction of 1,3propanedithiol and Ru₃(CO)₁₂. Ru₂(pdt)(CO)₆ was obtained as major product along with the multimetallic side product $\{(\mu-H)Ru_3(CO)_{10}\}_2(pdt)$. The $Ru_2(pdt)(CO)_6$ model along with tri(o-tolyl)phosphine in the presence of formic acid and triethylamine reports photocatalytic H₂ production activity with turnover frequencies of 5500 h⁻¹ and a turnover number over 24 700 h⁻¹. ²⁶⁸

The ligand exchange properties of the ruthenium models are very similar to those of the corresponding iron complexes (see Section 4.6). Phosphines and cyanides are likewise easily incorporated. 220,267-269 Still, it was reported that the cyanation reaction proceeds at faster rates as compared to the [FeFe]analogues due to increased electrophilicity of the Ru-Ru bond.²⁶⁹ In contrast, the protonation chemistry of the heavier homologues is distinctly different (see Section 5.3 for a detailed discussion of the protonation chemistry of the iron complexes). While the Os and Ru models (193 and 194) form a bridging hydride upon treatment with HBF4, the corresponding Fe complex remains unaffected. 192

Along this line, Rauchfuss and coworkers studied the photohydrogenation of Ru₂(pdt)(CO)₄(PCy₃)₂ (198). Notably, a terminal as well as a bridging hydride were observed in the very same complex, $HRu_2(pdt)(\mu-H)(CO)_3(PCy_3)_2$ (198- μ HtH).²⁷⁰ Contrarily, when a solution of 198 was subjected to HOTs, only the bridging hydride $[Ru_2(pdt)(\mu-H)(CO)_3(PCy_3)_2]^+$ ([198- μ H]⁺) was observed.270

Interestingly, the non-inertness of the conjugated acid $[Ru_2(pdt)(\mu-H)(CO)_4(CN)_2]^-$ (199) of the dicyanide model was shown in terms of ligand substitution and the complex readily undergoes decarbonylation when reacted with PMe3. Although showing a labile ligand binding, 199 reveals a hampered H₂ evolution activity as compared to the diiron analogues (-1.6 V vs. $-1.0 \text{ V for Fe}_2(\text{pdt})(\mu\text{-H})(\text{CO})_4(\text{CN})(\text{PMe}_3)$, both values referenced vs. Ag/AgCl).²⁶⁹

The successful synthesis of the [2Ru]_H-precursor [Ru₂(adt)- $(CO)_4(CN)_2]^{2-}$ (5) shed light on the proton reduction mechanistic pathways in the enzyme and was successfully incorporated into

apo-HydA1. Interestingly, the apoenzyme could be cleanly matured also with the bridging hydride. Spectroscopic data reveals that the protein environment affects the structure of these models, as the bridging hydride species, upon maturation, converts to the terminal hydride.

4.4.2 Incorporation of other metals – mixed-metal dithiolates. Inspired by the heterobimetallic centre of [NiFe]-hydrogenases, chemists investigated hetero- as well as homometallic derivatives of the FeFe-, MnFe-, NiFe- and CoFe-type. 271,272 The starting complex Fe(dithiolate)(CO)2(diphos) can be synthesised by reaction of FeCl₂ with a corresponding dithiolate and chelating phosphines (Fig. 60). Treatment of $Fe(pdt)(CO)_2(dppe)$ (200) with [(C₁₂H₁₀)Mn(CO)₃]BF₄ gave two isomeric Fe-Mn bimetallic complexes (unsym-201 and sym-201). The asymmetric isomer slowly converts to the symmetric as is depicted in Fig. 61.²⁷¹

This synthetic scheme was further extended to obtain the analogous ethanedithiolate models as well.271 In the case of edt models, the asymmetric to symmetric isomer interconversion did not occur even at a longer time scale. Upon protonation, $[(CO)_3MnFe(pdt)(CO)_2(\kappa^2-dppe)]^+$ converts to $[(CO)_3MnFe(pdt) (\mu-H)(CO)_2(\kappa^2$ -dppe), whereas it undergoes decarbonylation upon reduction and affords $[(CO)_3MnFe(pdt)(CO)(\kappa^2-dppe)]$.

For the mixed CoFe complex, $Fe(pdt)(CO)(\kappa^2-dppe)$ was reacted with CpCoI₂(CO) to give 202 followed by reduction with CoCp₂ to give the targeted complex 203. This method was found to be more reliable than the direct synthesis using CpCo(CO)₂ in refluxing toluene/THF. Moreover, to generate the protonated complex, the precursors were treated with HBF₄·Et₂O giving rise to the bridging hydride species at room temperature. 271,273

The synthesis of such mixed metal complexes is not restricted to first row transition metals. In 2003 Adam et al. reported on the MoMn complex 204 which was obtained by refluxing Mn₂S₂(CO)₇ with (MoCp(CO)₃)₂. CpMoM(μ-S)₂(CO)₅ reacts with various substrates to give the diverse derivatives 205 to 210 (Fig. 62).^{274,275} Similar to the synthesis routes reported for 201 and 202, the reaction of Mo(bdt)(CO)2-(PMe₃)₂ with Fe(CO)₅ afforded the MoFe complex 211 (Fig. 61).²⁷⁶

4.5 Substitution of carbonyl ligands in H-cluster models

Regarding the native H-cluster, the introduction of two cyanide ligands is the first transformation that comes into mind. However, the generated dianions are unsuitable for modelling the active site in many cases, e.g. because of undesired N-protonation of cyanides 150,153,277,278 or the instability of the oxidised species. 279,280 Therefore, the cyanide ligands are often replaced by phosphines exhibiting similar electron-donating properties without a negative charge. 277,278 Other attempts to

$$\begin{array}{c} \text{Cbz} \\ \text{Cbz} \\ \text{SH} \text{ SH} \end{array} + \text{Ru}_2\text{Cl}_2(\text{CO})_6 \end{array} \xrightarrow{Zn} \begin{array}{c} \text{OC} \\ \text{OC} \\ \text{OC} \\ \text{OC} \end{array} \xrightarrow{Ru} \begin{array}{c} \text{Ru} \\ \text{Ru} \\ \text{CO} \\ \text{CO} \end{array} \xrightarrow{Ru} \begin{array}{c} \text{NH} \\ \text{OC} \\ \text{CO} \\ \text{OC} \end{array} \xrightarrow{Ru} \begin{array}{c} \text{NH} \\ \text{OC} \\ \text{OC} \\ \text{OC} \end{array} \xrightarrow{Ru} \begin{array}{c} \text{NH} \\ \text{OC} \\ \text{OC} \\ \text{OC} \end{array} \xrightarrow{Ru} \begin{array}{c} \text{NH} \\ \text{OC} \\ \text{OC} \\ \text{OC} \end{array}$$

Fig. 59 Synthesis of 196 via the protected amine intermediate.

Synthesis of FeMn and FeCo models

Fig. 61 Isomerisation of 201 and structure of complex 211

modify the mimics involve the usage of N-heterocyclic carbenes (NHCs), isocyanides, or nitrosyls. The reaction pathways summarised in the following section are generally applicable and not limited to the herein described adt or pdt complexes.

4.5.1 Remarks on structural aspects. The ligands are denoted as apical (ap) or basal (ba) with respect to their position in the (idealised) square pyramidal coordination environment of the iron centres (Fig. 63). In some compounds only a single conformation is found, while others exhibit rapid exchange between almost energetically equal conformations at room temperature. This exchange is often referred to as turnstile rotation 281-283 - though there has been a debate, if turnstile rotation differs from a combination of Berry pseudorotation and other isomerisation mechanisms on a molecular level. 284-287 The dithiolate bridge was shown to also influence the position of the ligand beyond steric effects, at least in the solid state. For example, based on molecular structures, Rauchfuss and coworkers reported an apical/basal conformation of the cyanide ligands in $[Fe_2(pdt)(CO)_4(CN)_2]^{2-}$ (212), ²⁷⁹ in contrast to an apical/apical conformation in the corresponding adt and adt^{Me} complexes. 17,168

4.5.2 H-cluster models with cyanide ligands. In 1999, shortly after the crystal structures of [FeFe]-hydrogenases revealed the presence of two cyanide ligands and a bridging dithiolate ligand in the active site, the groups of Darensbourg, 288 Pickett, ²⁸⁹ and Rauchfuss ²⁷⁹ independently described the synthesis of the $[Fe_2(pdt)(CO)_4(CN)_2]^{2-}$ dianion (212, Fig. 64). As observed

in the hexacarbonyl complexes, fast isomerisation of apical and basal positions is observed at room temperature. 283 Following an associative mechanism, 282 complexes of this type are easily accessible in yields over 90% by addition of NEt₄CN to solutions of the corresponding hexacarbonyl complex in MeCN. 279,288,289 The cyanation can also be accomplished using KCN, but this case requires either solubilising crown ethers^{283,288} or elevated temperatures along with extended reaction times.²⁹⁰ The reactions to the dicyanides exclusively yield complexes in which one CO molecule on each iron atom is substituted. As observed for the incorporation of every (monodentate) electron donating ligand, 168,277,278,291 this regioselectivity can be explained by the increased Fe- C_{CO} bond strength due to an increased π -back bonding upon binding of the electron-rich cyanides. The increased bond strength is indicated by a shortened Fe-C_{CO} bond length in the crystal structure of the cyanide complexes²⁷⁹ and a shift of the carbonyl IR modes to higher wavenumbers.^{279,288,289} It is worth mentioning, that for the hexacarbonyl complex with an o-xyloldithiolate (o-xyldt) ligand substantially larger reaction rates for the dicyanation than for the pdt complex were observed. This led to the conclusion that the rotation of the Fe(CO)3 group, facilitated due to steric repulsion in the o-xyldt complex, is the first step of the cyanation in an associative pathway. 282,292 While many hexacarbonyl complexes are sufficiently oxygen-stable to be handled under ambient air, the cyanide complexes are generally less stable in the presence of oxygen - especially in solution. ^{279,289}

Fig. 62 Synthetic pathways towards various MnMo complexes.

Fig. 63 Apical and basal ligands and "turnstile rotation".

Using only one equivalent of NEt₄CN, the monocyanide [Fe₂(pdt)(CO)₅(CN)]⁻ (213) is unintuitively only formed in low yields up to ca. 15%. On the contrary, the dicyanide 212 is observed as the main product.²⁸³ This observation led to the conclusion that the monocyanide complex reacts more rapidly with cyanide ions than the hexacarbonyl complex.282 Indeed, kinetic measurements revealed that the intermediate formed upon mixing the hexacarbonyl complex and cyanides reacts more rapidly with additional cyanides than the hexacarbonyl complex. 282,283 Darensbourg and coworkers calculated a larger rate constant for the substitution reaction for the isolated monocyanide complex than for the hexacarbonyl.²⁸² In contrast, a competition experiment by the Rauchfuss group investigating the reaction of a mixture of hexacarbonyl and monocyanide complex revealed a slower CO/CN⁻ substitution in the monocyanide complex²⁸³ and DFT calculations supported this observation. 292 Moreover, the isolated monocyanide complex, as well as the hexacarbonyl complex 20 were shown to react only slowly with phosphines, while the intermediate rapidly reacts with these nucleophiles. Accordingly, the monocyanide was suggested as an implausible intermediate for the formation of the dicyanide by the authors. 283 As an alternative intermediate, a structure with a (semi-)bridging carbonyl was proposed.^{282,292} To the best of our knowledge, neither theoretical nor spectroscopic studies that further investigate the formation of the dicyanide without assuming the monocyanide as an intermediate were reported in literature. The reaction of the unidentified intermediate with nucleophiles is a useful method for synthesising asymmetric [(CO)2(CN)- $Fe(pdt)Fe(CO)_2(L)]^- (L = e.g. PMe_3 (214)).^{283} In contrast, using$ KCN under reflux conditions, the monocyanide was found to be an isolatable intermediate.290

Still, monocyanide complexes are accessible in good yield (ca. 75%) by converting one of the carbonyl ligands into a cyanide ligand using NaN(SiMe₃)₂. ^{282,288} Following a nucleophilic attack of the amide at the carbonyl carbon atom, the corresponding siloxane (SiMe₃)₂O is released. The increased electron density after the introduction of the cyanide hampers a second nucleophilic attack and accordingly a second CO

Fig. 64 Synthesis routes towards mono- and disubstituted cyanide and phosphine complexes

conversion is inhibited. This holds true even if an excess of NaN(SiMe₃)₂ is used.

The obtained sodium salt Na[Fe₂(pdt)(CO)₅(CN)] was reported to be unstable at room temperature even as a solid under inert atmosphere. 282 The sufficiently stable compound NEt₄[Fe₂(pdt)(CO)₅(CN)] was obtained in similar yields by the use of the decarbonylation agent trimethylamine N-oxide (Me₃NO) and subsequent addition of NEt₄CN to the solution. Widely applied for the selective substitution of one carbonyl ligand, Me₃NO oxidises a single carbonyl ligand to carbon dioxide, followed by dissociation from one iron centre. The vacant coordination site is presumably occupied by a coordinating solvent molecule (or NMe₃) to form a Fe₂(pdt)(CO)₅L (L = MeCN (215), NMe₃) species, that can be observed by IR spectroscopy with a significant lifetime in solution.²⁸³ Further support for this claim was provided by Rudolph, Ishii, Weigand and coworkers who were able to crystallize the corresponding acetonitrile-coordinated intermediate.²⁹³ The decarbonylated species can afterwards be attacked by nucleophiles, in this case cyanide. Similar to the attack of NaN(SiMe₃)₂, a second Me₃NO induced substitution is unfavourable.²⁸³

4.5.3 H-cluster models with phosphine ligands. While the cyanide ligands closely resemble the ligand environment in the native [2Fe]_H, the use of these ligands is accompanied by several drawbacks. First of all, the stability of the complexes – especially towards oxygen – is dramatically decreased. Moreover, protonation studies (*vide infra*) revealed undesired protonation of the cyanide ligands. ^{277,278} This protonation, however, does not occur natively due to encapsulation in the enzyme's binding pocket and H-bonding to the protein. ²⁹⁴ On the contrary, phosphine complexes cannot be protonated at the phosphorous atoms under the typical H₂ formation conditions but at the same time show similar electronic characteristics. The formal exchange of two cyanide ligands by electron-rich trimethyl phosphine ligands shifts the IR bands to higher wavenumbers

by only 15 cm⁻¹ to 20 cm⁻¹ still being shifted the opposite way by *ca.* 100 cm⁻¹ compared to the hexacarbonyl complexes.^{277,278}

The special coordination environment in the enzyme does not only change the reactivity of the system, but also its structure. Phosphines certainly are the most important group of ligands in the field of hydrogenase research due to the same reasons frequently discussed in other fields of homogenous catalysis. 295-297 Due to the strong metal phosphorous bonds, low valent metal complexes are stabilised. In addition, their electronic and steric properties can be easily controlled by their organic substituents and quantified e.g. by the Tolman electronic parameter, 298,299 the ligand cone angle, 300 and the percent buried volume.301 The use of chelating bi- or tridentate phosphine ligands permits additional flexibility in the coordination sphere. Phosphine ligands also allow for a covalent linkage to groups with further functionalities and were used to increase the water solubility, 129,130,302 provide an internal proton relay,303-305 or link the active site mimic to proteins,306 photosensitisers, 307,308 redox-active ferrocenes 204,209,213,309-312 or other redox-active ligands. 313,314

Notably, the reactivity of mono- and bidentate phosphines towards unbridged complexes $Fe_2(SR)_2(CO)_6$ is already known since the 1970s and many results from extensive studies of Haines, Greenwood and coworkers accord with the reports on the dithiolate derivates. ^{315,316}

4.5.3.1 Monodentate phosphines. The synthesis of hydrogenase model complexes substituted with monodentate phosphines is generally straightforward (Fig. 64). Typically, at room temperature the hexacarbonyl complexes undergo direct substitution of one carbonyl ligand by the phosphines more slowly than cyanides.²⁸³ Still, within hours the monosubstituted complexes of less bulky phosphines (e.g. PMe₃, PMe₂Ph) are formed. Disubstituted byproducts are also regularly detected. For the reaction of more bulky phosphines (e.g. PPh₃, P(OEt)₃),

often elevated temperatures are needed. 170,317 By prolonged reaction times and elevated temperatures likewise disubstituted complexes are accessible in good yields for less bulky phosphines, 277,278,318 while for more bulky phosphines the yields range from moderate to low. 129,317,319 As observed for the cyanide complexes, the second substitution always occurs on the remaining Fe(CO)₃-fragment. ^{277,278,317,318} The monosubstituted complexes are also easily available by decarbonylation with Me₃NO in the presence of the desired phosphine. ^{275,283,303} A second substitution using this method only gives low yields, while thermolysis and photolysis are useful tools and afford high product yields (>90%) also for bulky phosphines. 180 In neat PMe₃, photolysis of the edt, pdt and adt hexacarbonyls slowly yields the tetrasubstituted electron-rich complexes $Fe_2(xdt)(CO)_2(PMe_3)_4$ (216xdt) in a yield of ca. 60%. 320

If the model complex is synthesised starting from Fe₂S₂(CO)₆, addition of the phosphine to the intermediate Fe₂(SH)₂(CO)₆, generated at dry ice temperature, also allows for the selective substitution of one carbonyl ligand (217) upon warming to room temperature.321 Fe₂(SCH₂OH)₂(CO)₆ (85) can be substituted analogously and afterwards condensed to the oxa- or azadithiolate complexes Fe₂(xdt)(CO)₅(PR₃) $(e.g. R = Me 218xdt).^{185}$

Mono- and disubstituted phosphine complexes, compared to their cyanide analogues, are typically more stable. The conformations of disubstituted phosphine derivatives are briefly discussed using the example of Fe₂(xdt)(CO)₄(PMe₃)₂ (xdt = edt, pdt; e.g. R = Me 219xdt). In the solid state, a transoid basal-basal arrangement of the phosphine ligands is found for the pdt derivative, while the edt complex features a apical-basal arrangement. 277,278 This arrangement for edt is also found in acetone at -60 °C, while NMR spectra reveal high fluxionality of the Fe(CO)₂(PMe₃) units at room temperature.²⁷⁸ The same interconversion is found for pdt, however, the conformation in solution was found to be highly dependent on the solvent. A comparison of calculated IR spectra with those obtained in different solvents revealed that the transoid basal-basal conformer is almost entirely found in heptane (90%) and hexane, while in acetonitrile and methanol significant amounts of the apical-basal (up to 40%) isomers are found. 322,323 In accordance with the increased amount of the more polar apical-basal isomer in polar solvents is the finding that the apical-basal isomer is predominant in acetone. 278,324,325 The cisoid basal-basal and the apical-apical conformers are not observed.

While monodentate phosphines exclusively give Fe, Fe'-disubstituted complexes, the use of bidentate phosphines makes other coordination modes accessible. By chelation of one iron atom by bidentate phosphines, strongly electron-rich iron centres can be synthesised and an asymmetry between both iron centres can be introduced. However, the different accessible coordination modes render the introduction of bidentate phosphines more complicated. The favourable coordination mode is not only determined by the applied reaction conditions but also by the linker of two phosphines moieties (backbone).

4.5.3.2 Bis(diphenylphosphino)methane (dppm). As κ^2 -chelate complexes with dppm suffer from steric strain of the fourmembered ring, the ligand is known to bridge bimetallic complexes preferably in a κ^1, κ^1, μ -geometry in favour of less strained five-membered cycles. In [FeFe]-hydrogenase mimics this coordination mode is likewise the most prominent (Fig. 65). Complexes $Fe_2(xdt)(CO)_4(\mu-dppm)$ (220xdt, xdt = e.g.edt, pdt, adt^{alkyl}, odt) with two symmetric iron centres are readily prepared (>70% yield) from the hexacarbonyl complexes in refluxing toluene. 183,208,326,327 Contrary, the use of Me₃NO and the phosphine was reported to reduce the yield. 183,326 The chelate complex $Fe_2(pdt)(CO)_4(\kappa^2-dppm)$ (221pdt) was observed as a crystalline side product but could not be isolated in the bulk. 183,326 Similar results were reported for $Fe_2(pdt)(CO)_4(\mu$ -dcpm) (222pdt, dcpm = bis(dicyclohexylphosphino)methane).326 The photolytic introduction of dppm into Fe₂(odt)(CO)₆ was also reported but gave lower yields.²⁰⁸ In contrast to complexes with two monodentate phosphine ligands, the bidenatate phosphine ligands cannot adopt the apical position in these complexes and always adopt a cisoid basal-basal arrangement, which is otherwise sterically unfavourable.

The stable complexes $Fe_2(xdt)(CO)_5(\kappa^1-dppm)$ (223xdt) with a monodentate dppm ligand can also be isolated; in the case of pdt from the reaction of the hexacarbonyl diiron complex and dppm in acetonitrile in the presence of Me₃NO. In case of adt^{n-propyl} the reaction time has to be shortened from 60 min to 30 min. Both, the pdt and adt complexes can also be converted into to the corresponding μ-dppm compounds by thermolysis. 183,326 In an attempt to synthesize the κ^{1} -dcpm complex (224), the oxygen-sensitive ligand was partially oxidised by Me₃NO at both phosphine moieties (Fig. 66).326

In addition, asymmetric κ^2 -chelate complexes can be synthesised if steric bulk is introduced to the backbone carbon atom as displayed by C(Me)₂(PPh₂)₂ i.e. Me₂dppm. The room temperature reaction of the hexacarbonyl complex 20, Me₂dppm, and Me₃NO gives the κ^2 -chelate complex 225 in 63% yield. While in solution the apical/basal and basal/basal isomers were detected in a 2:1 ratio by 31P-NMR, only a dibasal arrangement was observed in the crystalline form. In refluxing toluene, the chelate complex slowly rearranges to the bridged complex with a μ-Me₂dppm ligand.³²⁸

A similar ligation behaviour as observed for Me2dppm was reported for aminodiphosphine-ligands (R₂P)₂NR' (PNP). The amine in the backbone of the ligand was proposed to enable a similar reactivity as the adt-amine and allowing PCET mechanisms. The thermodynamically favoured μ-PNP geometry comprising a five-membered ring is accessible by the direct reaction of the hexacarbonyl complexes with the PNP ligands in high boiling solvents, e.g. toluene or xylene. 130,308,329,330 In an alternative synthesis route, Fe₂(CO)₆(μ-CO)(μ-PNP) (226) can be prepared photochemically from Fe₂(CO)₉, and subsequent reaction with the bridging dithiol.³³¹ By this route, the first dppm complex, Fe₂((μ-S)₂CNR)(CO)₄(μ-dppm) (227), with a bridging dithiocarbamate ligand was synthesised.332

In contrast to dppm, κ²-chelate complexes (228xdt) comprising PNP ligands can be isolated in moderate to high yields

Fig. 65 Different synthetic pathways for the incorporation of dppm and derived ligands as well as PNP ligands into hydrogenase mimics

Fig. 66 Reaction of dcmp with complex 20 in the presence of Me₃NO.

(33% to 90%). The small PNP bite angle ligands favour a dibasal geometry in solution and solid state.330,331 The substitution behaviour of PNP ligands in acetonitrile after decarbonylation with Me₃NO also strongly depends on the substituent R' on the amine. For R' = H, the obtained chelate complex was always formed together with the bridged µ-PNP complex in a 2:1 ratio and could not be isolated in pure form. ³³⁰ PNP ligands with sterically less demanding substituents (e.g. R' = Me, Bu, $(CH_2)_nNMe_2$ (n = 2, 3), etc.) smoothly react to the desired chelate complexes (55% to 90% yield). 130,330,331,333 With increasing steric bulk on nitrogen, the formation rate of the chelate complex slows down and for aryl-substituted amines often very low yields (<10% yield) were reported. Herein, the major product is the monosubstituted complex $Fe_2(pdt)(CO)_5(\kappa^1-PPh_2NHR')$ (229), in which the second P-N bond was hydrolysed. 330,333,334 A more general route towards the chelate complexes is provided by irradiation of the hexacarbonyl precursor with UV light in toluene.333 The PNP chelate complexes also rearrange slowly to the thermodynamically favoured bridged isomers if solutions in toluene are heated to reflux. 130,330,331

4.5.3.3 Bis(diphenylphosphino)ethane (dppe) and bis(diphenylphosphino)ethene (dppv). In contrast to dppm, the chelation of one iron centre is the most common coordination mode for dppe (Fig. 67) and its unsaturated analogue cis-bis(diphenylphosphino)ethene (dppv, Fig. 68). The formed five-membered rings are thermodynamically more favourable than the analogous four-membered rings in the case of dppm. Thereby, asymmetric complexes $Fe_2(xdt)(CO)_4(\kappa^2$ -diphosphine) are accessible. Another effect of the increased bite angle is that the basal-apical geometry of the chelate complexes becomes more favourable.335-337 The chelate complexes 230xdt (dppe) and 231xdt (dppv) form upon decarbonylation of the hexacarbonyl complexes in the presence of the phosphine ligand within hours. 154,210,326,336-338 Because of its rigidity, dppv (Fig. 68) forms the chelate complexes at room temperature, 154,210,326,337 while dppe (Fig. 67) requires elevated temperatures. 336,338 The reported yields vary significantly in the range of 26% to 90% for dppv and 23% to 47% for dppe. The thermal reaction without Me₃NO is also feasible but requires considerably extended reaction times, 326 gives lower yields, 326,339 or demands the use of the more

OC
$$R_2P$$
 PR_2
 $R = Ph$ 232pdt

$$A = Ph$$
 R_2P PR_2
 $R = Ph$ R_2P PR_2
 $R = Ph$ R_2P PR_2
 $R = Ph$ R_2P R_2

Fig. 67 Complexes with dppe ligands comprising different binding modes and their selective synthesis.

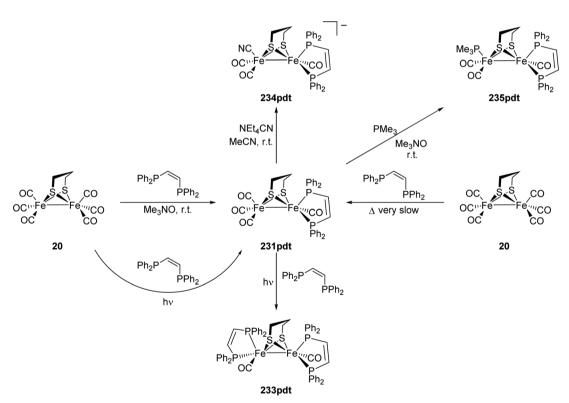


Fig. 68 Complexes bearing dppv ligands comprising different binding modes and their selective synthesis.

electron-rich and sterically less demanding bis(dimethylphosphino)ethane (dmpe) ligand (Fig. 69). 159,340,341 Upon elongated heating the κ^2 -dppe complexes rearrange to afford the thermodynamically even more favourable μ -dppe complexes (232xdt). 342,343 For dppv the light-induced substitution of the hexacarbonyl

complexes was also reported;³⁴⁴ however, this method is more often applied for the introduction of a second dppv ligand on the other iron centre to give the very electron-rich complexes 233xdt of high symmetry. 53,154,210 Interestingly, while Fe,Fe'-disubstituted complexes [L(CO)₂Fe](xdt)[Fe(CO)₂(L)] (L = PR₃, CN⁻) are unreactive

Incorporation of dmpe into complex 237 yielding three different complexes

towards further substitution under various conditions, additional electron donating substituents can be introduced into the complex $Fe_2(CO)_4(edt)(\kappa^2-dppv)$. Cyanide (MeCN, r.t., 234xdt), PMe₃ (Me₃NO, toluene, r.t., 235xdt) and the bulkier phosphines PCy₃ and $P^{i}Pr_{3}$ ($h\nu$) replace a carbonyl ligand on the unfunctionalised iron centre under the same conditions known for the corresponding monosubstituted complexes. The pdt analogue shows similar reactivity but the substitution processes proceeds more slowly (Fig. 68). 337,345

Dppe is also capable of linking two [2Fe-2S]-units giving "intermolecular" bridged complexes of the type [Fe₂(xdt)-(CO)₅]₂(µ-dppe) (236xdt) with half a dppe ligand per [2Fe-2S]unit. This coordination mode is favoured if only one equivalent of dppe is added after decarbonylation with Me₃NO and the reaction mixture is stirred at room temperature. 145,183,339,346 In addition, this coordination mode was also observed for dmpe complexes containing the sterically crowded dithiol (Ph)₂Si(CHPh(SH))₂. From the reaction of the corresponding hexacarbonyl diiron complex 237 with one equivalent of dmpe predominantly the [4Fe-4S]-compound 238 is formed (Fig. 69). If two equivalents of dmpe are used, the chelate complex 239 is only a side product (7% yield) while the bridged complex $Fe_2((\mu\text{-SCHPh})_2SiPh_2)(CO)_4(\mu\text{-dmpe})$ (240) is the main product (35% yield). 159 Interestingly, complex 239 features the unusual rotated geometry, which will be discussed in Section 5.1.

Complexes of the type $Fe_2(xdt)(CO)_4(\kappa^2$ -diphosphine) can also be synthesised starting from iron(II) chloride and Fe(bda)(CO)₃ (bda = benzylideneacetone) (Fig. 70, compare Section 4.4.1). Thereby, the selective labelling with Mössbauer-active ⁵⁷Fe is possible. Here, iron chloride reacts with the diphosphine under one atmosphere of carbon monoxide at room temperature and (the dilithium salt of) the dithiol bridge to give the monometallic precursor. 271,347 The formed Fe(xdt)(CO)₂(κ²-diphosphine) complex then reacts with Fe(bda)(CO)3 as the source for the Fe(CO)3 fragment to give the asymmetric complex $Fe_2(xdt)(CO)_4(\kappa^2$ -diphosphine) (e.g. 231pdt). 271,348 Using 57FeCl₂, the subsequent introduction of a second equivalent of dppv to afford the symmetric complex ⁵⁷FeFe(pdt^{Me})(CO)₂(κ^2 -dppv)₂ (233pdt^{Me}, pdt^{Me} = 2,2-dimethyl-1,3propanedithiol) was enabled.348 Rauchfuss and coworkers were also able to synthesize mixed-metal mimics by exchanging the Fe(CO)₃ source by a Mn(CO)₃ or a Co(Cp) precursor (see Section 4.4.2).271

4.5.3.4 1,1'-Bis(diphenylphosphino)ferrocene (dppf). Redoxactive ferrocene ligands were used to mimic the function of the [4Fe-4S]-cluster providing electrons to the [2Fe-2S]-cluster. While the monodentate, ferrocene substituted phosphines feature the same behaviour as other monodentate phosphines, 213,310,349 dppf shows coordination modes known from dppm and dppe. The connection of two [2Fe-2S]-units is common for the dppf ligand and is accessible via reaction of the hexacarbonyls with 0.5 equiv. of dppf in the presence of Me₃NO at room temperature in varying yields between 24% and 90%. 204,209,213,309,312,343 With toluene-3,4-dithiolate as the bridging ligand, also monodentate dppf in Fe₂(bdt^{Me})(CO)₅(κ^{1} -dppf)

Fig. 70 Synthesis of selectively ⁵⁷Fe-labelled complex **231pdt**.

(241) was observed. If 20 is refluxed in toluene in the presence of dppf for an elongated time, the "intramolecular" bridged complex $Fe_2(pdt)(CO)_4(\mu\text{-dppf})$ (242) is formed.³¹¹

4.5.3.5 Other phosphines. Other bi- or tridentate phosphine ligands follow similar trends in terms of their substitution behaviour. For 1,2-bis(diphenylphosphino)benzene – as observed for dppv – the Me₃NO-induced substitution is more convenient than the thermal reaction in refluxing toluene. The introduction of the electron-poor and thus, less reactive 4,5-bis(diphenylphosphino)-4-cyclopenten-1,3-dione with electron-withdrawing carbonyl-groups requires additional heating under refluxing conditions in toluene. Notably, the related compound 2,3-bis(diphenylphosphino)maleic anhydride is prone to decomposition under these conditions, but can be incorporated into the [2Fe–2S]-framework by photolysis, which is also a known alternative method for the introduction of dppv ligands. 313

In accordance with the stability of five- and six-membered chelate rings, the propane-analogue of dppe, 1,3-bis(diphenylphosphino)propane (dppp) also forms chelate complexes upon CO-substitution. Under the latter conditions the complex slowly rearranges to afford the bridged μ-dppp complex.³⁴² The 2-(n-alkyl)aza-analogues also form the expected chelate complexes as the major product in refluxing toluene together with small amounts of the intramolecular bridged complex and the tetranuclear cluster. 204,350,351 The latter can be synthesised selectively by decarbonylation with Me₃NO. The N-phenyl substituted phosphine gives the "intermolecular" bridged complex either with Me₃NO induced decarbonylation or in refluxing toluene. 351 Decarbonylation and subsequent prolonged heating in toluene allowed for the isolation of the respective chelate complex.352 The additional methylene group in the backbone of 1,3-bis(diphenylphosphino)butane (dppb) renders the formation of the chelate compound unfavourable and the "intramolecular" bridged compound is formed under both conditions. As observed for dppe, the ligands with longer carbon chains in the backbone initially form the linked [4Fe-4S]-complexes. This coordination mode can be exclusively obtained if trans-bis(diphenylphosphino)ethene is used.³⁴²

Similarly, *meta*-substituted pyridyl and pyrimidyl phosphines also adopt this coordination mode if 0.5 equiv. are used. In the presence of one equivalent of the phosphine ligand monosubstituted complexes $Fe_2(xdt)(CO)_5(\kappa^1-L)$ are formed.³⁰⁵ The tridentate phosphine bis(diphenylphosphinoethyl)phenylphosphine can be introduced in refluxing toluene and adopts a μ,κ^1,κ^2 -coordination mode showing both features of the chelate and the bridged complexes. The constraints of this coordination mode distort the square pyramidal coordination environment around the iron centres and lead to complexes with 'rotated state' character. 353,354

4.5.4 H-cluster models with other ligands

4.5.4.1 Isocyanides. Especially in the early days of [FeFe]hydrogenase research, isocyanide ligands gained interest. With a similar Fe-C

N framework, but without a reactive nitrogen centre, isocyanide complexes were shown to give more stable hydrides as well as oxidised species (vide infra) as compared to the cyanides. 355-357 The substitution of the carbonyl ligands (Fig. 71) occurs slowly at room temperature. However, after twenty hours to two days at room temperature the single- or double-substituted complexes can be obtained. 213,355,358 For example, at 70 °C, isocyanomethane (CNMe) gives the doublesubstituted complex Fe₂(pdt)(CO)₄(CNMe)₂ (243) after two hours in acetonitrile.356 Prolonged heating gives the three- and fourtimes substituted complexes (244 and 245) (Fig. 71). 356,357 If Me₃NO is applied, the double-substituted complex is also accessible selectively at room temperature within hours. 213,283 Notably, 1,4-diisocyanobenzene was also highlighted to enable bridging of two [2Fe-2S]-subunits (246).^{204,213}

4.5.4.2 N-heterocyclic carbenes (NHCs). Like phosphines, N-heterocyclic carbenes (NHCs) are strong σ -donor ligands with tuneable steric properties. Their similar donor strength, stability of the corresponding complexes as well as their more convenient handling, e.g. compared to PMe₃, raised the interest in exchanging phosphine ligands by NHCs. ^{359,360} Accordingly, NHCs were also applied for the generation of electron-rich (mono)substituted active site mimics. The substitution of the carbonyl ligands by NHCs in the hexacarbonyl complexes

CNR, Me₃NO MeCN, r.t. slow MeCN/ DCM
$$\Delta$$
 fast

RNC SS CNR OC Fe Fe CO OC RNC CO RNC CO RNC CO R = Me 243

RNC SS CNR OC Fe Fe CO OC RNC CO RNC CO RNC CO R = Me 243

Fig. 71 Substitution of carbonyl ligands by isocyanides.

Fig. 72 Synthesis of NHC-substituted models and further functionalisation

occurs at room or moderately elevated temperatures (up to 60 °C) (Fig. 72). The sterically less demanding ligand 1,3-dimethylimidazol-2-ylidene (IMe) solely forms the monosubstituted complex at room temperature in high yields (83% for pdt),³⁶¹ while at 60 °C the disubstituted complex Fe₂(pdt)(CO)₄(IMe)₂ (247) is preferably formed (50% yield compared to 12% of the monosubstituted complex).291 With sterically more demanding NHCs (e.g. 1,3-dimesitylimidazol-2-ylidene (IMes)) only the monosubstituted complex 248 is observed in high yields (>70%). 181,362-364 Under reflux conditions in toluene, PMe3 can be introduced into the monosubstituted complexes on the Fe(CO)₃ fragment to give very electron-rich asymmetric mimic Fe₂(pdt)(CO)₄(IMes)(PMe₃) (249, Fig. 72). 361,365 The substituents at the nitrogen atoms have also been functionalised with additional donor groups. While thioethers and amines were shown not to coordinate the iron atom even in the presence of Me₃NO, 363 the pyridine nitrogen in 1-methyl-3-(pyridylmethyl)-imidazol-2-ylidene binds to the iron centre immediately (250).364 Interestingly, if the symmetric, N,N'-disubstituted 1,3-bis(pyridylmethyl)-imidazol-2-ylidene ligand is used, only the carbonic carbon atom and none of the pyridines coordinates to the iron centre in 251 at room temperature and even at 60 °C only 20% yield of complex 252 with a κ^2 -NHC ligand was obtained. However, one of the pyridine nitrogen atoms can be coordinated upon addition of Me₃NO to the monodentate complex. By this method the κ^2 -complex is obtained in 90% yield. The coordination of the other pyridine moiety or a ligation of the second iron atom are not observed.³⁶⁴ Similar to bidentate phosphines, bis-carbene ligands with different alkyl linkers were introduced. With these ligands monosubstituted complexes,

chelate complexes (253), intermolecular bridged complexes (254) and mixtures thereof were obtained in low to moderate yields (up to 50%).181,363

Like mono-substituted phosphine complexes, NHC complexes are reasonably stable under ambient conditions even in solution. Carrying an additional phosphine ligand, the stability of the complexes towards air is massively reduced in solution, however the solids can be handled under air. 361

4.5.4.3 Thioethers and sulfoxides. In contrast to cyanides and carbonyls, the third ligand observed in the active site -(metallo-)thioethers - gained considerably less interest as discrete ligands. However, thioethers and sulfoxides can also be introduced as terminal ligands by decarbonylation with Me₃NO and subsequent addition of the ligand at room temperature (255, Fig. 73). 366-368 In an alternative route, thioethers can be incorporated after one of the carbonyl ligands is converted to a Fischer-type carbene (256b) with n-BuLi and [Et₃O]BF₄. The organometallic equivalent of a thioether Cp(CO)2FeSPh or Cp(CO)₂Fe(Cys-κS), mimicking the native cysteine link to the [4Fe-4S]-cluster, can also be incorporated by the former method. 178,204,209 The chemistry of [2Fe-3S]-assemblies is also part of Section V.

4.5.4.4 Amines and pyridines. By heating Fe₂(pdt)(CO)₆ to reflux in propylamine, the complex Fe₂(pdt)(CO)₅(NH₂Pr) was obtained. This complex was shown to be stable in non-coordinating solvents. However, in coordinating solvents, e.g. acetonitrile, the amine ligand is replaced by the solvent molecule.369 In contrast,

Fig. 73 Two methods for the incorporation of thioethers into [2Fe]_H-mimics.

complexes with pyridine donors and derivatives were found to be stable. The substitution with pyridine proceeds via decarbonylation with Me₃NO, ³⁷⁰ while the chelating ligands 2,2'-bipyridine³⁷¹ and 1,10-phenantroline³⁷² were introduced in refluxing toluene. The incorporation of an additional PMe3 substituent was also shown to be feasible with Me₃NO for the latter one. ³⁷³ Adding two additional thiolate functions to 2,2'-bipyridine e.g. in 2,2'-([2,2'-bipyridine]-6,6'-diyl)bis(1,1-diphenylethane-1-thiolate) (= L^{N2S2}) yields the chelate complex 257 (Fig. 74) after adding NiCl₂ in THF at room temperature. Interestingly, changing the metal from nickel to iron, e.g. $Fe(BF_4)_2$, yields $[Fe_2(L^{N2S2})_2H]^+$ (258), a dimeric structure in which one thiolate is protonated.³⁷⁴ Both complexes can be reacted with [CpFe(CO)(MeCN)₂]⁺ yielding the respective NiFe- and FeFecomplexes (259 and 260). 375,376 Thus, the LN2S2 ligand closes the gap between [FeFe]- and [NiFe]-hydrogenase models, enabling both types of metal content.

4.6 Towards structural and functional H cluster models

The rich chemistry of $Fe_2(SR)_2(CO)_{6-n}L_n$ complexes, some of which developed almost a century ago, enabled the preparation of diverse synthetic mimics of the $[2Fe]_{H}$ -cluster. The first and most simple model complexes bearing a dithiolate bridge were synthesised decades before the crystal structure was known. Based on the knowledge gained out of these studies, a variety of more accurate active site mimics were reported after the determination of the crystal structure. At first, the efforts

focused on an exact representation of the ligands on the iron centre and the incorporation of different bridgehead atoms. Aiming at both structural and functional mimics, more complex alterations of the active site were pursued. By now, numerous protocols for the incorporation of different (functionalised) dithiolate bridges as well as many classical organometallic ligands and a combination thereof are available. The Fe(CO)₂ platform enables versatile ligand substitutions by associative or after thermal, photochemical or chemical decarbonylation by dissociative pathways. Both, functionalised ligands and bridges can be used to achieve the desired electronic or structural properties of the mimic and allow for the attachment of photosensitizers, redox active groups, or proteins. Much of the progress, in the development of synthetic protocols was driven by the aim to understand the relationship of structure and function of the [2Fe]_H-cluster. The most prominent example in this regard is the amine in the dithiolate bridge. Even before its presence in the active site was confirmed, the proposed function as a proton relay was confirmed in e.g. protonation experiments and gave strong evidence for its utmost importance. In the next sections, we will present how the synthetic progress gave rise to complexes that were able to mimic almost all relevant features (e.g. rotated structure, CO-binding, protonation behaviour, proton reduction, hydrogen oxidation, etc.) observed in the protein. Although there is no mimic known that is alone capable to display all properties of the natural active site, the variety of model complexes together is able to cover all aspects of [FeFe]-hydrogenases' activity.

V Structural models of the active site

In this section, we will present how by combination of different ligands and dithiolate bridges sophisticated structural or functional models of the active site were designed.

5.1 H-cluster models mimicking the rotated state

The distal iron atom in the native H-cluster features an unusual coordination environment, which is referred to as the 'rotated state'. Namely, the square pyramidal geometry is inverted compared to the hexacarbonyl and dicyanide complexes. One of the formerly terminal carbonyl ligands subsequently adopts a semi-bridging position. This unusual geometry is not found in the hexacarbonyl complexes, in which the carbonyl ligands of the two iron centres are eclipsed and many of the reported, substituted model complexes show the same unrotated geometry. However, due to steric interactions or other constraints deviations

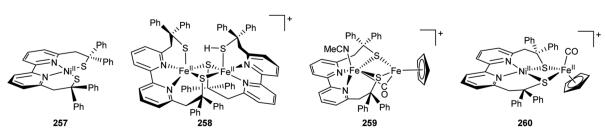


Fig. 74 Two methods for the incorporation of thioethers into [2Fe]_H-mimics.

from the idealised, eclipsed geometry were observed. Aiming at structural models of different enzyme states, more elucidated models featuring the special rotated geometry were developed and will be presented herein.

Already in 2003 Bruschi, Fantucci, and De Gioia³⁷⁷ presented a DFT study on Fe^IFe^I complexes, which proposed stable energetic minima corresponding to a rotated state. In 2006, Tye, Darensbourg, and Hall³⁷⁸ then investigated the influence of various substitution patterns on the stability of rotated structures with DFT calculations and showed that these structures become more favourable when altering the electron donating ligands (e.g. phosphines) from Fe,Fe'- to Fe,Fedisubstituted complexes. In accordance with earlier reports on oxidised and reduced species with bridging carbonyl ligands, 357,379,380 the calculations also affirmed the increased stability of the rotated structures upon oxidation/reduction. In addition, it was proposed that strongly electron donating ligands on one iron atom in combination with strongly electron accepting ligands on the adjacent iron centre is another promising strategy towards "rotated models" - a strategy which is brought to an extreme by the use of NO+ ligands in synthesis.^{381–383} With one exception, in which one carbonyl ligand is forced into a bridging position by binding to strong Lewis acids,⁵³ at least one of these three strategies, is found in every active site mimic featuring a rotated geometry.

5.1.1 Remarks on the rotated geometry. Three criteria are used to describe and quantify the rotation of the iron centre denoted as Fe_d in analogy to the enzyme. First, the coordination environment around the rotated iron centre is evaluated ideally being square pyramidal (Fig. 75). Upon distortion, the coordination is better described as trigonal bipyramidal. Although the Fe_p - Fe_d - C_b angle Ψ , where Fe_p is the unrotated iron centre and Cb is the carbon atom of the semi-bridging carbonyl ligand, is reduced for the formerly basal carbonyl ligand, now in equatorial position, the models are still regarded as unrotated. Contrary, in rotated complexes again a square pyramidal conformation is found. In this case, the coordination around the iron is inverted. The Addison τ parameter is used to quantitatively distinguish between a trigonal bipyramidal and a square pyramidal coordination environment. The value is, according to eqn (1), determined by the difference between the two angles formed by opposing Fe_d-S and Fe_d-C_{CO} bonds divided by 60°.

$$\tau = \frac{\langle S_2 F e_d C_2 - \langle S_1 F e_d C_1 \rangle}{60^{\circ}} \tag{1}$$

In a square pyramidal environment, the four involved bonds span a basal plane. Accordingly, both angles equal 180° giving $\tau = 0$. In reality, for unrotated as well as rotated complexes, the iron atom is distorted out of the basal plane towards the apical ligand, so that both angles adopt values smaller than 180°, e.g. 156° and 160°, in the case of the pdt hexacarbonyl complex.²⁸⁸ Still, in a symmetrical coordination environment τ is almost equal to zero. If the ligands around the iron form a trigonal bipyramid, a carbonyl ligand and a thiolate are in axial positions, while the other two ligands span the trigonal plane together with the third carbonyl ligand. This arrangement corresponds to an angle of 180° between the axial bonds, respectively 120° between the equatorial bonds. Hence, in trigonal bipyramidal environments $\tau = 1$. However, values of $\tau = 1$ are not observed in real structures for two reasons, which will be explained at the example of Fe(pdt^{Et})(CO)₄(PMe₃)₂ (261, pdt^{Et} = 2,2-diethyl-1,3-propanedithiolate, Fig. 76).³⁸² In this complex, the coordination environment of one iron centre can be described as trigonal bipyramidal. The iron, one of the sulphur atoms and both carbonyl atoms span the equatorial plane with only small deviations. Still, the angles between the ligands at the iron centre do not all equal 120°. The two S-Fe-C angles have values of 128° and 129° and the angle between the carbonyl ligand equals 102°, which is at the upper end observed for such complexes. Accordingly, in none of the severely distorted complexes reported in literature the largest angle in this plane $\angle (S_1 - Fe_d - C_1)$ is smaller than 127° . ³⁸¹ In addition, the axial ligands almost never adopt positions, in which the angle △(S₂-Fe_d-C₂) between them is 180°, i.e. because of the constraint S-Fe-S angle. Typical values are 167° to 170°, 169° in the case of 261, with various examples below this range and only two examples 159,382 above a value of 173°, both of which with

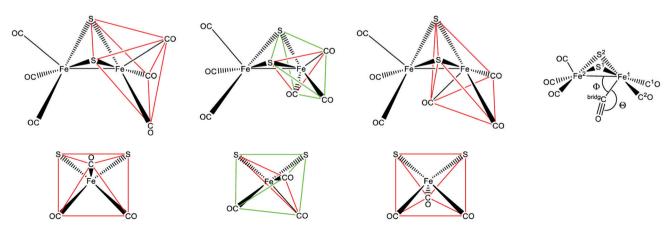


Fig. 75 Left: Two views of the unrotated (left), a distorted, trigonal-bipyramidal (middle) and a rotated (right) geometry from two perspectives. For clarity, bond lengths are exaggerated and all ligands are omitted. Right: Rotated complex, in which the angles Ψ and Θ are indicated.

Fig. 76 Molecular structure of complex **261**. CCDC ID: PEGCOV. 382

Fig. 77 Nitrosyl substituted complexes 262xdt and 263xdt.

smaller τ values (0.48 and 0.17) and sterically crowded dithiolates. Due to these deviations from an idealised trigonal bipyramidal environment Addison's τ -parameter is always considerably smaller than $\tau=1$. For **261**, a value of $\tau=0.67$ is obtained and with the limits of \pm (S_1 – Fe_d – C_1) = 127° and \pm (S_2 – Fe_d – C_2) = 173° described above an estimated maximal value of $\tau=0.77$ can be extrapolated. Considering this behavior, complexes for which τ is about 0.75 can be considered trigonal bipyramidal and represent the transition from an unrotated to a rotated geometry. This upper limit should be considered when using Addisons' τ parameter to quantify the rotation of the iron centre. Moreover, the value is influenced by the position of four out of five ligands and therefore, sensitive to small changes in their position, *e.g.* due to steric effects, which can complicate the comparison of different complexes.

As a second aspect, the torsion angles between the apical ligand on the unrotated iron centre and the ligands on the rotated iron centre can be evaluated. Especially, the largest including the (semi-bridging) carbonyl – and the smallest angle that can be regarded as the distortion of the eclipsed unrotated complex are of interest. Being the most intuitive values, care must be taken as the apical ligand on the unrotated iron can also be distorted from its idealised position (up to 20°) most often due to steric repulsion with the dithiolate bridge. In this context, 261 is again a useful example. Torsion angles of 75° (regarding C¹) and 176° (regarding C_b) are observed indicating a fully rotated complex. However, the apical ligand on the less distorted iron centre does not lie symmetrically between both sulphur atoms. If instead of the apical ligand, the pdt bridgehead carbon, which adopts an almost symmetric position, is used for the calculation, more realistic torsion angles of 56°,

respectively 157° are obtained. Even though the idealised torsion angles are useful quantifiers for the rotation of the iron centre, the described distortion can render these values meaningless, without careful evaluation of the complete complex.

The third aspect is the (semi-)bridging character of the "inverted" carbonyl ligand. The four relevant structural values discussed in the literature to distinguish between terminal, semi-bridging and bridging carbonyls are the Fe_p-Fe_d-C_b angle Ψ , the Fe_d-C_b-O_b angle Θ and both Fe-C_b distances. Crabtree and Lavin³⁸⁴ examined the correlation between these structural parameters in (semi-)bridging carbonyl ligands of different iron carbonyl complexes. In this discussion, structural parameters of different classes of iron carbonyl ligands were used, and even though, all of the complexes herein belong to a single class in different oxidation states, with different ligands, and different steric strain these values and their correlations are astonishingly valid. For terminal carbonyl ligands Ψ is about 100° in hexacarbonyl complexes and substituted complexes without steric bulk, e.g. $Fe_2(pdt)(CO)_4L_2$ (L = CO, CN⁻, PMe₃). ^{277,279,288} If carbonyl ligands adopt positions with a higher bridging character, this value decreases. While for semi-bridging carbonyls this decrease correlates linearly with a decrease of Θ , in the case of terminal carbonyls, uncorrelated values of $\Theta > 170^{\circ}$ corresponding to almost linear carbonyl ligands are observed. Based on the extrapolation of Ψ for which $\Theta = 180^{\circ}$, Crabtree and Lavin suggested $\Psi = 75^{\circ}$ as the frontier between terminal and semi-bridging carbonyl ligands. Even though Θ is not smaller than for terminal carbonyl ligands for Ψ values between 70° and 75°, there is a distinct difference between these ligands. While for values $\Psi < 75^{\circ}$ the oxygen atom is bent away from Fe² corresponding to semi-bridging carbonyls, the orientation of the ligands is governed by steric effects for increased values. For $\Psi = 75^{\circ}$ a distance Fe_p-C_b of 2.69 Å was extrapolated, almost in the middle between the sum of the covalent radii (1.94 Å) and the sum of the van der Waals radii (ca. 3.5 Å). Accordingly, this distance is also a useful criterion for the evaluation of the bridging character. The length of the actual M-CO bond can also be considered. Below an angle Ψ of 70°, the bonds are lengthened due to a diminished π -backbond and correspond to a regular single bond in the symmetrically bridging case. As described above, the π-backbond strengthens and the M-CO bond shortens upon binding of electron rich ligands. The different bond lengths have to be considered, when comparing different complexes, which

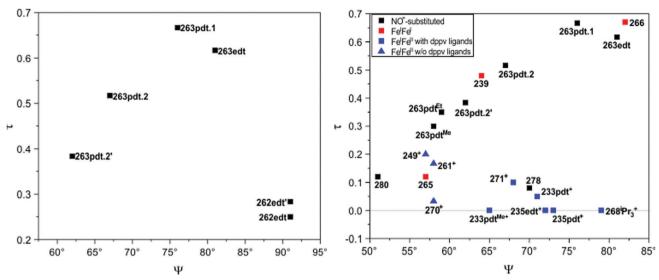


Fig. 78 Left: Addisons' τ parameter against Ψ /° for the nitrosyl complexes **262edt**, **263edt** and **263pdt**. **263pdt**. **1** refers to the ba/ap rotamer and 263pdt.2 to the ba/ba rotamer. The 'denotes different independent molecules in the asymmetric unit. Both 263pdt.2 molecules are weakly disordered, but the angles vary only by 1°. For all molecules of **263pdt** τ was calculated with the semi-bridging carbonyl in apical position. Right: Addisons' τ parameter against $\Psi/^{\circ}$ for different rotated complexes.

makes the other values Ψ , Θ , and Fe_p - C_b more robust in the context of this evaluation.

In Fig. 78, the values Ψ and τ of the nitrosyl substituted complexes $[Fe_2(xdt)(CO)_{4-n}(PMe_3)_{1+n}(NO)]^+$ (xdt = edt, pdt) (Fig. 77) are plotted. The unrotated complex 262edt is characterised by a low τ value and a large angle Ψ . With increasing degree of rotation, Ψ decreases while τ increases initially. The strongly distorted molecule 263edt and the rotamer 263pdt.1 are characterised by high τ values and angles Ψ at the edge of terminal to semi-bridging carbonyls. Both values decrease hereafter to characterise the rotamer 263pdt.2 with a rotated structure.

As an additional aspect in this regard, the carbonyl IR stretching frequencies are useful to evaluate their bridging character especially if no crystal structure is available. Similar to the M-CO bond length, the frequencies are also sensitive to the electron-donating abilities of other ligands, as well as oxidation states of the iron centres and accordingly, have to be assessed carefully. Still, in comparison of similar compounds carbonyl stretching frequencies and Ψ match well. More importantly, the observation of the band of the semi-bridging CO ligand by IR spectroscopy provides an easy, but powerful tool to assess the structure of the complexes in solution, which can differ strongly from the solid state (vide infra). Table 6 summarises various structural and spectroscopic parameters of complexes featuring a rotated or distorted geometry.

5.1.2 Fe^IFe^I models displaying the rotated geometry. Already in 2001, Pickett and coworkers reported a metastable intermediate of the cyanation of Fe₂(κ³-pdt^{MeSMe})(CO)₆ (53, pdt^{MeSMe} = 2-methyl-2-((methylthio)methyl)propane-1,3-dithiolate) bearing a bridging carbonyl ligand. This complex has no open coordination site as the H_{ox} or H_{red} states, but more closely resembles the H_{ox} -CO state (see Section 5.2). ^{138,139,385} Until 2007,

no complex with a bridging carbonyl ligand and an open coordination site was known. Thereafter, Justice, De Gioia and Rauchfuss⁵³ showed that a rotated geometry is indeed accessible outside the protein. They reported on model complexes 264xdt (xdt = edt, pdt), in which the bridging CO ligand is stabilised by binding to strong Lewis acids (AlBr₃, B(C₆F₅)₃) (Fig. 79). Initial experiments with AlBr₃ showed, that strongly Lewis basic carbonyl ligands are required for a sufficient binding strength, accessible by the ligation of the iron centres by electron-donating groups. While $Fe_2(edt)(CO)_4(\kappa^2-dppv)$ and $Fe_2(edt)(CO)_3(\kappa^2-dppv)(PMe_3)$ were shown not to be sufficiently Lewis basic, in cyanide substituted complexes, those ligands would be the preferred binding site for the Lewis acid. Therefore, $Fe_2(xdt)(CO)_2(\kappa^2-dppv)_2$ (233xdt, xdt = edt, pdt) with four electron donating phosphine ligands were chosen for this study. Indeed, IR- and NMR-data are in line with calculated DFT spectra and indicated the formation of the rotated complexes $Fe_2(xdt)(\mu\text{-COAlBr}_3)(\kappa^2\text{-dppv})_2(CO)$ in solution. Unfortunately, no structural data is available for a comparison with other rotated complexes, in which fundamentally different effects cause the rotation. Notably, the binding of the Lewis acids to the pdt derivative is stronger compared to the edt complex. This was explained by the destabilisation of the apical CO ligand by steric interactions with the methylene group. Thus, the Lewis acid-bound conformation with the bridging carbonyl ligand becomes more favourable.

This effect that is caused by steric repulsion between the bridge and the apical ligand is also an integral part of other rotated Fe^IFe^I models. Moreover, all complexes feature, as proposed by Tye, Darensbourg and Hall, an asymmetric ligand environment on the [2Fe-2S]-core, rendering one iron atom strongly electron-rich due to two donating phosphine ligands. The very similar complexes 265³⁴¹ and 266,³⁴⁴ reported in 2013, were the first rotated Fe^IFe^I complexes characterised by X-ray

Table 6 Structural parameters of distorted and rotated active site mimics

		Ox. State	Ligands					Bond lei	ngth			
Complex	Dithiolate	$\mathrm{F}e_d/\mathrm{F}e_p$	Fe_{d}	Fe_p	Ψ / $^{\circ}$	$oldsymbol{arTheta}/^{\circ}$	τ	Fe_d – C_b	$\mathrm{Fe}_{\mathrm{p}}\text{-}\mathrm{C}_{\mathrm{b}}$	μ -CO ν /cm ⁻¹	CCDC ID	Ref.
265	adt ^{Bn}	I/I		dmpe	57	146	0.12	1.815	2.179	1777 ^a	YIJDEC	341
266	pdt ^{Et}	I/I		dppv	65	158	b	1.744	2.499	1818 ^a		344
239	$Ph_2Si(C(Ph)HS^-)_2$	I/I		dmpe	64	156	0.48	1.789	2.379	1801^{a}	KOYKAM	159
235edt ⁺	edt	I/II	dppv	PMe_3	72	170	0.00	1.781	2.628	1883	XIGFEZ	386
235pdt ⁺	pdt	I/II	dppv	PMe_3	73	170	0.00	1.786	2.678	1889	AGEHIE	345
$268P^{i}Pr_{3}^{+}$	pdt	I/II^c	dppv	$P^{i}Pr_{3}$	79	178	0.00	1.785	2.833	1870^{c}	AGEHOK	345
249	pdt	I/II	I_{Mes}	PMe_3	57	152	0.20	1.864	2.196	1861	LIHCAH	365
270^{+d}	pdt ^{Me}	I/II	PMe_3	PMe_3	58	152	0.03	1.813	2.204	1859	NONFEC	387
261 ⁺	pdt ^{Et}	I/II	PMe_3	PMe_3	58	153/146	0.17	1.839	2.222	1874	PEGCUB	382
233pdt ⁺	pdt	I/II	dppv	dppv	71	170	0.05	1.786	2.618	1884	RIXQAS	388
233pdt ^{Me+} e	pdt ^{Me}	I/II	dppv	dppv	65	163	0.00	1.793	2.460	1.854	HELPOG	348
271^{+}	adt ^{Bn}	I/II	dppn		68	167	0.10	1.769	2.516	1896	AVUSAM	389
278	SCR^-	\mathbf{I}/\mathbf{II}^f	NO, PPh_3		70	168	0.08	1.783	2.570	1875 ^a	GICRUF	390
262edt	edt	\mathbf{I}/\mathbf{II}^f	NO, PMe_3		91	177	0.25	1.821	3.170		NOJVUE	381
262edt′	edt	\mathbf{I}/\mathbf{II}^f	NO, PMe_3		91	179	0.28	1.827	3.162		NOJVUE	381
263edt	edt	\mathbf{I}/\mathbf{II}^f	NO, PMe_3	PMe_3	81	173	0.62	1.792	2.872		NOJWEP	381
263pdt.1	pdt	\mathbf{I}/\mathbf{II}^f	NO, PMe_3	PMe_3	76	172	0.67	1.794	2.479		NOJWIT	381
263pdt.2 ^d	pdt	\mathbf{I}/\mathbf{II}^f	NO, PMe ₃	PMe_3	66	160	0.52	1.813	2.472		NOJVOY	381
263pdt.2′ ^d	pdt	\mathbf{I}/\mathbf{II}^f	NO, PMe_3	PMe_3	62	154	0.38	1.780	2.325		NOJVOY	381
280	pdt	$\mathbf{I}/\mathbf{I}\mathbf{I}^f$	NO, I _{Mes}	PMe_3	51	136	0.12	1.747	2.028		UGOGUU	383
263pdt ^{Me}	pdt ^{Me}	$\mathbf{I}/\mathbf{I}\mathbf{I}^f$	NO, PMe ₃	PMe_3	58	148	0.30	1.802	2.182	1877	PEGDEM	382
263pdt ^{Et}	pdt ^{Et}	$\mathbf{I}/\mathbf{I}\mathbf{I}^f$	NO, PMe ₃	PMe_3	59	153	0.35	1.797	2.234	1874	PEGDAI	382
261	pdt ^{Et}	I/I	PMe_3	PMe_3	82	172	0.67	1.747	2.873	1899	PEGCOV	382

' second molecule in asymmetric unit, .1/.2 are rotamers. ^a Only observed in solid state. ^b Not all crystallographic data are reported. ^c Geometric and electronic situation is different in solution. d Disordered, values given for the stronger rotated molecule. e Second molecule in the asymmetric unit with negligible structural deviations. f Regarding the non-innocent NO-ligand other assignments are also reasonable.

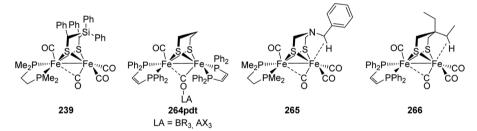


Fig. 79 Fe¹Fe¹ models of the [2Fe-2S]-cluster exhibiting a rotated geometry

crystallography (Fig. 79). Both complexes feature an additional stabilisation of the free coordination site on the rotated iron site through an agostic interaction. This interaction, although being weak, was identified by DFT calculations as crucial feature for obtaining the rotated geometry in the crystalline state. Notably, the asymmetric unit of the crystals of 266 feature two independent molecules one of which is rotated while the other is unrotated. This indicates a small energy difference between both confirmations. Likewise, it suggests that packing effects in the crystal also play an important role for stabilising the rotated conformation.

The coordination geometry of complex 239¹⁵⁹ with a sterically heavily crowded dithiolate ligand shows a distorted, inverted square pyramidal geometry in the molecular structure. Nevertheless, the coordination environment also has a considerable trigonal bipyramidal character ($\tau = 0.48$). The Fe(CO)₃ unit is rotated out of an idealised square pyramidal environment by about 10°-15°. The authors denoted this geometry as

semi-rotated and concluded that the missing agostic interaction, compared to 265 and 239, prevents full rotation. By examination of other geometric parameters, a different conclusion can be proposed. One of the carbonyl ligands has considerable semibridging character (Ψ = 64°, Θ = 156°, Fe_p-C_b = 2.379 Å), less than in 265 but slightly higher than in 266. In addition, the angle $\not \leq$ (S₁-Fe_d-C₁), which is used for the calculation of τ , is unusually large. The value of 175° is at least 5° larger than usual for rotated as well as unrotated complexes leading to an unusual linear arrangement, which could be caused by the high steric strain. Assuming a typical value of 170 $^{\circ}$ for this angle, τ would equal 0.4 in accordance with a less distorted complex. Regarding the positions of C₁O and the phenyl group, that shields the open coordination site, in the crystal (Fig. 80) the distortion may be attributed to steric repulsion between the two groups. Accordingly, it can be proposed that the complex is a fully rotated complex that is distorted due to steric strain. Due to the low number of comparable complexes and the small energy

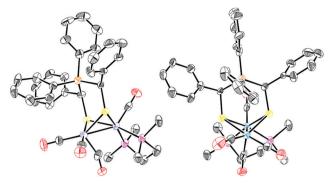


Fig. 80 Molecular structure of complex 239 featuring a rotated geometry. CCDC ID: KOYKAM. 159

differences in the DFT calculations regarding this complex this explanation was not investigated further.

Even though all three complexes 239, 265 and 266 were crystallised in the rotated state and μ-CO bands were detected in the IR spectra of the solids, the solution IR spectra show no bands for bridging carbonyl ligands. Accordingly, for none of the three complexes the rotated geometry is preserved in solution. Rotated complexes in solution were, except from **264xdt**, only observed for oxidised or NO⁺-bound complexes. A very important feature of the three complexes 239, 265 and 266 is that in each complex the Fe(CO)₃ unit is rotated. This is in accordance with the calculations of Tye, Darensbourg and Hall, 378 but no mimic in other oxidation states with a rotated Fe(CO)₃ unit has been crystallised.

These four examples indicate that for Fe^IFe^I complexes a rotated geometry can be obtained by following the following strategies:

- (1) Destabilisation of the apical ligand through steric bulk on the dithiolate.
 - (2) Stabilisation of the bridging carbonyl ligand.
- (3) Stabilisation of the vacant coordination site through weak intramolecular interactions.
- (4) Introduction of an electronic asymmetry on the two iron centres.

5.1.3 H_{ox}-models displaying the rotated geometry

5.1.3.1 $Fe^{I}Fe^{II}$ H_{ox} -models. Since the structure of the active Hox state was identified, dedicated efforts in modelling its key feature - the rotated geometry - were invested. The formation of a bridging CO ligand in Fe^IFe^{II} H_{ox}-models was first reported

for the very same system as for the Fe^IFe^I-complexes by Best, Pickett and their coworkers in 2002. In situ IR measurements of (electro-)chemically oxidized [Fe₂(pdt^{MeSMe})(CO)₄(CN)₂]²⁻ (267) revealed a μ-CO band (see Section 5.2). 379 In 2007, the groups of Rauchfuss³⁸⁶ and Darensbourg³⁶⁵ independently presented the mixed valent Fe^IFe^{II} complexes 235edt⁺ and 249⁺ showing a rotated geometry (Fig. 81). Both complexes were prepared by oxidation of asymmetrically substituted, not rotated model complexes with ferrocenium in non-coordinating solvents and isolated as their [BF₄]-, respectively [PF₆]- salts. A crucial feature of both complexes is a sterically demanding and strongly electron donating ligand on the rotated iron atom. While Rauchfuss and coworkers applied a dppy ligand on the rotated iron centre, Liu and Darensbourg installed an I_{Mes} ligand. In both cases, the unrotated iron atom is substituted with a PMe₃ ligand, which stabilises the oxidised Fe^{II} centre. In these complexes, the rotated geometry is not only present in the solid state, as observed for the rotated Fe^IFe^I models, but also can be detected in solution, as indicated by bands for (semi-)bridging CO ligands in the IR spectra. Complexes with other dithiolates 235xdt⁺ (pdt, adt, adt^{Bn}, odt)³⁹¹ and phosphines $[Fe_2(pdt)(CO)_3(\kappa^2-dppv)(PR_3)]^+$ (268PR₃⁺, PR₃ = PCy₃, PⁱPr₃)³⁴⁵ were also reported by Rauchfuss later on. The exchange of PMe₃ by dppv also afforded rotated complexes 233pdtR+ with pdt^{388,392} and pdt^{Me} bridges.³⁴⁸ By using the redox active phosphine PEt₂Fc* instead of PMe₃, Camara and Rauchfuss presented elaborate, functional model complexes 269adtR+ with adt^R bridges (R = Bn, H). The IR spectra of these complexes also indicate a rotated complex geometry.349 In addition, Darensbourg and coworkers reported that less bulky NHCs lead to less stable complexes that could not be characterised by X-ray diffraction. The bands assigned to μ-CO are strongly blueshifted by 68 cm⁻¹ compared to 249+ in the solution IR spectra of these complexes and indicate a greatly reduced or no semibridging character of the carbonyl ligand. Thus, steric bulk was identified as a key feature of rotated complexes.361 Interestingly, it could be shown that the introduction of steric bulk on the dithiolate bridge (pdtMe and pdtEt) allows for rotated complexes 270⁺ and 261⁺ even with two small PMe₃ ligands. The authors highlighted that, in contrast to 249⁺ and 270⁺, the higher thermal stability of 261⁺ allows for room temperature EPR measurements. 382,387 Rauchfuss and coworkers reported later that the limited thermal stability of their compounds can be avoided by using [BAr^F₄]⁻ instead of [BF₄]⁻ as the counterion.³⁹³

Fig. 81 Structures of selected mixed-valence Fe^lFe^{ll} complexes showing a rotated geometry.

Fig. 82 Rotated, mixed-valence complexes bearing an adjacent amine.

The corresponding complexes showed no decomposition in solution at room temperature for days. The instability of the [BF₄] anion towards electrophilic iron complexes has been reported earlier - interestingly also in the context of complexes for hydrogen activation/generation. 394

Without doubt, all these complexes can be regarded as rotated state mimics (τ , torsion angles, crystal structure). Upon closer inspection of the structural features, some distinct differences are obvious. While the models with the NHC ligand and the bulky bridges all show carbonyl ligands with a high semi-bridging character (Ψ < 60°), in the dppv substituted complexes the carbonyl ligands are on the edge between terminal and semi-bridging ($\Psi = 71^{\circ} - 79^{\circ}$). Only in the complex with the bulky pdt^{Me} bridge, the carbonyl ligand has considerably higher semi-bridging character ($\Psi = 65^{\circ}$) attributed to the steric repulsion between dppv and the dithiolate. Remarkable in this regard is also the structure of complex 268PiPr₃⁺, as the low-energy μ -CO band (1870 cm $^{-1}$) in the solution IR spectrum indicates a higher bridging character of the semi-bridging carbonyl ligand but the opposite is found in the crystal ($\Psi = 79^{\circ}$).

Supported by additional DFT calculations, 382,387,395 in most complexes the iron centres were accordingly attributed the oxidation states $Fe_{p}^{II}Fe_{d}^{I}$. 345,361,382,387 This is especially interesting, as the same assignment is assumed in the active H_{ox} state (see Section 3.2). The EPR measurements of the PiPr₃ and PCy₃ substituted complexes 268PiPr3+ and 268PiCy3+ indicate another assignment despite the similar crystal structure of $268P^{i}Pr_{3}^{+}$ compared to complexes $235edt^{+}$ and $235pdt^{+}.^{345}$ While for all other complexes the spin (S = 1/2) is mainly localised on the rotated iron centre, and in some cases partially on the dithiolate, 382 in the complexes substituted with bulky phosphines ³¹P hyperfine coupling constants indicate that the spin is localised on the other iron centre. Comparing the solution IR spectra to spectra of a rapidly precipitated solid and grown single crystals, in the latter of which no μ-CO band is detected. The authors concluded that in solution the other iron centre Fe(CO)₂(PR₃) is rotated exhibiting a stronger semibridging carbonyl ligand. In this case, the spin resides on the rotated iron centre also in solution.

In order to synthesise a more electrophilic Hox-model for hydrogen oxidation experiments, Camara and Rauchfuss³⁸⁹ synthesised the tetracarbonyl complex $Fe_2(adt^{Bn})(CO)_4(\kappa^2$ dppn) 271 (dppn = 1,8-bis(diphenylphosphino)naphthalene). Its cation 271⁺ is, in contrast to other adt tetracarbonyl cations, stable at room temperature in solution for at least 24 h.

Its structure, the first crystallographically confirmed of a rotated adt complex, is very similar to the dppv complexes reported by Rauchfuss with a higher semi-bridging character ($\Psi = 68^{\circ}$) of the carbonyl ligand on the rotated Fe(μ-CO)(dppn) unit. The complexes $Fe_2(xdt)(CO)_4(\kappa^2-L)^+$ (272xdt⁺, xdt = adt^{Bn}, edt, pdt, bdt^{3Me}; $L = {}^{n}PrN(CH_{2}PPh_{2})_{2}$) also feature the rotated geometry and an enzyme-like proton relay. 396,397 These complexes resemble the only examples of rotated [FeFe]-hydrogenase mimics, in which the unrotated iron centre is unsubstituted. By EPR measurements, the same $Fe_{p}^{II}Fe_{d}^{I}$ electronic structure was identified.

If the oxidation of the electron-rich precursors is conducted in coordinating solvents, e.g. acetonitrile, the coordination of a solvent molecule promotes the oxidation to the diferrous complexes. 337,345,386,398 Likewise, in the presence of phosphines, 398-400 cyanides, 399 isocyanides, 356,357 or even the adt amine^{391,401} oxidation to the Fe^{II}Fe^{II} complexes occurs. A bridging carbonyl ligand is a prominent feature of many of the obtained complexes. Moreover, the diferrous complexes are labile towards further substitution to give highly substituted complexes. An overview over these oxidatively induced ligand substitutions is given in Fig. 83.

5.1.3.2 Hydrogen oxidation by H_{ox} -models. Despite the high activity of [FeFe]-hydrogenases for the dihydrogen oxidation, 402,403 the application of mimics as proton reduction catalysts always dominated research and dihydrogen activation is rare with mimics. Still, H/D exchange reactions catalysed by bridging hydride complexes indicated heterolytic dihydrogen activation. 277,278,355,404 The photolytic oxidative addition of dihydrogen to Ru₂(pdt)(CO)₄(PCy₃)₂ (198) reported in 2004, gave the dihydride complex $HRu_2(pdt)(\mu-H)(CO)_3(PCy_3)_2$. In the presence of coordinating solvents or counterions hydrogen is released upon protonation, while in CH₂Cl₂ with [H(OEt₂)][BAr^F₄] (Ar^F = 3,5-bis(trifluoromethyl)phenyl) a dihydrogen σ -complex is formed.²⁷⁰ For the iron complex 231edt, the photolytic oxidative addition to a similar dihydride complex was also reported. In the presence of B(C₆F₅)₃ heterolytic dihydrogen activation yields a complex bearing a bridging hydride. 405 The ruthenium and the iron complexes are both capable of oxidatively adding other E-H (E = Cl, O, S, Si) bonds.

The first, actual biomimetic dihydrogen activation was reported for the rotated Hox-models 235adtR+ [Fe2(adtR)(CO)3(P- Me_3 (κ^2 -dppv) BAr_4 (R = H, Bn) (Fig. 82). The corresponding pdt and odt complexes are incapable of performing this reaction, again highlighting the importance of the internal

Fig. 83 Top: Oxidation of complex 235xdt in the presence and absence of coordinating solvents. Middle: Substitution of a carbonyl ligand in a diferrous complex. Bottom: Exemplary complexes obtained by oxidatively induced ligand substitution.

proton shuttle for any hydrogenase-like activity. However, the reaction was reported to be slow and required high H₂ pressures (12.4 MPa). In the presence of additional oxidants (e.g. substituted ferrocenium) the reaction rate is strongly increased.³⁸⁹ This observation was explained with the mechanism shown in Fig. 84. Similar to a PCET, the heterolytic cleavage of dihydrogen is proposed to occur simultaneously to the oxidation of the complex by the additional oxidant, which is incapable of oxidising 235adtR+. Without ferrocenium present in solution a second molecule of 235adtR+ is oxidised, which limits the reaction rate. The rate limiting step in the presence of an oxidant is the binding of H2 to the open coordination site. The formed, but not observed doubleprotonated intermediate is then most likely deprotonated, either by an additional base e.g. P(o-tolyl)3 or by reduced 235adt^{R+}, yielding the bridging hydride. At some point during the mechanism, the formed terminal hydride isomerizes to a bridged hydride. Thus, in the absence of additional oxidants and bases only half of the bridging hydride is directly formed from the heterolytic dihydrogen cleavage, while the other half is formed by reduction and subsequent protonation. Contrary, in the presence of excess oxidant and base, more realistic for the situation in the enzyme, the bridging hydride is formed

quantitatively from heterolytic H2 activation. The more electrophilic complex 271⁺ shows an even higher reaction rate. Still, both complexes do not show any catalytic activity, which is attributed to the fact that deprotonation of the bridging hydride is not feasible. Indeed, complexes 272xdt⁺, in which deprotonation of the bridging hydride is easily accessible, shows catalytic hydrogen oxidation activity. 396 For this complex, also the double-protonated intermediate is observed. Other models that show catalytic activity, though at low turnover numbers, are complexes 269adtBn+ and 269adt. Here, the internal oxidant not only increases the rate of H₂ activation compared to 235adt^{R+}, but increases the acidity of the bridging hydride upon oxidation. This allows for the deprotonation required for catalytic H_2 oxidation.

Subsequently, Hogarth, Holt and coworkers reported on the electrochemical H₂-oxidation catalysed by Fe₂(pdt)(CO)₄(dppf) also featuring an internal oxidant in the presence of base. However, the mechanism in the absence of an internal proton shuttle remained unclear and the authors tentatively suggested an intermediary dihydride species.311

5.1.3.3 Nitrosyl substituted Hox-models. Binuclear, nitrosyl substituted iron sulphur complexes are known for more than 150 years. In 1858, Roussin reported on the "red salt"

Fig. 84 Hydrogen oxidation by complexes 235adtR+.

Fig. 85 Exemplary nitrosyl substituted complexes. Note: in the crystalline state 263pdt shows different degrees of rotation.

 $K_2[Fe_2S_2(NO)_4]$ and the corresponding "ester" $Fe_2(SR)_2(NO)_4$. 406,407 Interestingly, the first rotated diiron nitrosyl complex $[Fe_2(\mu,\eta^2\text{-SCR})(CO)_4(NO)(PPh)_3]$ (278) was already reported in 1988 by Behrens and coworkers (Fig. 85). In this complex, the iron centres are bridged by a thioacyl moiety and a semibridging carbonyl ligand, which is not observed in the absence of the phosphine. 390 Nitrosyl substituted hydrogenase mimics were firstly reported in 2008 by the groups of De Gioia and Rauchfuss. 381 Also in this first report, the authors showed that the substitution of carbonyl by nitrosyl ligands can lead to rotated complexes. Since assigning (formal) oxidation states in the presence of nitrosyl ligands can be difficult $(NO^{\dagger} \nu s. NO^{\bullet})$, complexes with nitrosyl ligands are discussed separately.

Generally, NO⁺ used as the [BF₄]⁻ salt replaces a carbonyl ligand within several hours at room temperature or even 0 °C. The reaction is limited by the poor solubility of NOBF4 in CH₂Cl₂, which is used to decrease the rate of decomposition of the products. The diamagnetic nitrosyl complexes are airsensitive and temperature labile, but sufficiently stable at 0 $^{\circ}\text{C.}^{381-383,408}$ Handling and storing the complexes under an atmosphere of CO was also reported to increase the stability of the complexes.³⁸³ In literature, only nitrosyl complexes of electron-rich mimics are known. Not surprisingly, several studies showed that the nitrosyl cation attacks at the more electron-rich iron atom. Further substitution reactions with electron donating ligands (PMe3, CN-) were shown to occur on the nitroso substituted iron centre. 381,383 Apart from the very electron rich $Fe_2(pdt)(CO)_2(\kappa^2-dppv)_2$ (233pdt), only monosubstituted nitrosyl derivatives are reported. Induced by the steric repulsion of the two dppv ligands, $[Fe_2(pdt)(CO)(\kappa^2-dppv)_2(NO)]^+$ (279pdt) is also unusual as the NO+ ligand adopts a basal site, whereas it is usually apical in unrotated complexes. 408

Nitrosyl complexes gained interest after the first report of a rotated complex bearing a nitrosyl ligand. $[Fe_2(pdt)(CO)_3(P-Me_3)_2(NO)]^+$ 263pdt was shown to crystallise as an apical and

a basal rotamer referred to the PMe₃ on the Fe(CO)₂(PMe)₂ unit. In the apical rotamer 263pdt.1 the Fe(CO)(NO)(PMe₃) unit has a high trigonal bipyramidal character ($\tau = 0.67$) and the carbonyl ligand little semi-bridging character ($\Psi = 76^{\circ}$, Fe_p-C_b = 2.479 Å). In contrast, two independent molecules of the basal rotamer **263pdt.2** with higher square pyramidal character ($\tau = 0.52, 0.38$) and higher semi-bridging character of the carbonyl ligand $(\Psi = 67^{\circ}, 62^{\circ})$ were crystallised. This again emphasizes the importance of packing effects on the rotation of crystalline mimics. Also, in the complexes 263pdtR (R = Et, Me) and $[Fe_2(pdt)(CO)_3(NO)(I_{Mes})(PMe_3)]^+$ (280) rotation is induced upon binding of NO^{+, 382,383} Complexes of these type are also rotated if the oxidized state lacks the NO ligand - however, [Fe2(pdt)- $(CO)_2(\kappa^2\text{-dppv})(PMe_3)(NO)]^+$ (281) is in contrast unrotated.⁴⁰⁸ The authors presumed that the nitrosyl ligand is not sufficiently electron-withdrawing to overcome the effect of two donating phosphines and induce electronic asymmetry.

The introduction of a NO $^{^+}$ has effects similar to an oxidation, not only on the molecular but also on the electronic structure as indicated by DFT calculations and Mössbauer measurements. $^{381-383}$ Accordingly, the oxidation states should be assigned to Fe(II){Fe(I)(NO $^{\bullet}$)} with an antiferromagnetic coupling to give diamagnetic complexes, though other assignments are also possible. Indeed, complexes $279pdt,\ 281,\ and\ [Fe_2(pdt)(CO)_3(\kappa^2-dppv)(NO)]^+$ (282pdt) were also prepared by first oxidising the corresponding precursors and a subsequent treatment with NO $^{\bullet}$ under the substitution of a carbonyl ligand. 345,386,408

5.1.3.4 Reactivity of H_{ox} -models towards CO. Within the enzyme the active site of [FeFe]-hydrogenases is reversibly inhibited by carbon monoxide affording the H_{ox} -CO state. A similar reactivity was also observed for some of the rotated complexes with an open coordination site. Though, the eclipsed complexes typically do not exchange carbonyl ligands between both iron centres and do not interact/exchange with

extrinsic CO except after photodissociation of a carbonyl ligand. 282

The reactivity of Hox models towards CO is governed by their stability and basicity. At -78 °C under 1 atm of CO, 249 does not form a stable CO adduct. However, regioselective incorporation of ¹³CO on the rotated iron centre is observed. ^{361,409} The same regioselectivity for 13C-exchange albeit under different conditions (5 °C, $h\nu$) is found in the active site of *D. desulfuricans*. ⁵⁶ At room temperature, ¹³CO is also incorporated into the positions on the unrotated iron centre. 361,409 The corresponding, less stable complex 283+ with an I_{MesMe} ligand instead of an I_{Mes} ligand incorporate ¹³CO in all positions already at -78 °C. Moreover, in addition to the fully labelled complex, a CO adduct is observed. For 249^+ , 283^+ and the I_{Me} substituted complex $283'^+$ CO adducts are formed upon increasing the CO availability by sparging solutions at -78 °C with CO. While the more stable 249⁺ reacts slowly and is in equilibrium with its CO adduct, the CO adduct formation is quantitative and fast for 283⁺ and 283'⁺. In all cases the starting material is recovered upon exposure of the solutions to vacuum or purging with Ar/N₂. Notably, the CO release of 283⁺CO and 283'⁺CO is considerably slower. DFT calculations on 283'+, supported by the EPR spectrum of 283⁺CO, indicated that in both complexes the extrinsic CO does not bind to the NHC substituted iron atom (which exhibits an open coordination site in the solid state of 249⁺) but to the PMe₃ substituted one.³⁶¹

Complexes ${\bf 233edt}^+, {}^{392}{\bf 235edt}^+, {}^{386}{\bf 235pdt}^+,$ and ${\bf 268P}^{\bf i}{\bf Pr_3}^{+345}$ bind CO at −45 °C within seconds/minutes. 235edt⁺ releases the bound CO upon purging with N₂ at 0 °C, while for 233⁺ removal of the CO atmosphere was reported to be sufficient. The adduct unsym-233⁺CO³⁹² formed upon the reaction of 233⁺ with CO shows the same orientation (ba/ba; ap/ba) of the dppv ligands as its precursor. Upon warming a solution of $unsym-233^{+}CO$ to -30 °C or if 233 is oxidised at 0 °C in the presence of CO, the symmetric adduct sym-233⁺CO is formed in which both dppv ligands adopt a basal/apical orientation. This adduct of the cation of the highly basic precursor 233 and CO is significantly more stable and was crystallised from a CO-saturated solution. The crystal structure confirms the results of computational studies on 283'+CO and 233+CO, that predicted a symmetric carbonyl ligand and an unusually elongated Fe-Fe distance (2.70 Å in the crystal). 361,392 The high bridging character of the carbonyl ligand in 233⁺, 235edt⁺, 235pdt⁺ and 268PⁱPr₃⁺ is also reflected in the strongly lowenergy shifted μ-CO bands around 1790 cm⁻¹. DFT calculations and the EPR spectrum of 283+CO are in accordance with a delocalisation of the spin on both iron centres, which is distinctly different from the situation in the H_{ox} models, but again reflects the situation in the enzyme properly.361 Worth mentioning, Silakov, Lubitz, and coworkers also observed spin delocalisation in the H_{ox} -CO model 267^+ . 410

The nitrosyl complexes 263xdt and $[Fe_2(edt)(CO)(\kappa^2-dppv)_2(NO)]^+$ (279edt) bind CO reversible at low temperatures. In contrast to the complexes without a nitrosyl ligand, the bridging position is occupied by a nitrosyl ligand as indicated by IR spectroscopy. ^{381,408} The less basic complexes 262xdt, 282edt, and $[Fe_2(pdt)(CO)_4(I_{Me})(NO)]^+$ (284) are not sufficiently

basic to form stable adducts. 381,383,408 The latter incorporates 13 CO, but presumably via a dissociative pathway. 383

5.1.4 Rotated state models – a highlight in biomimetic modelling. While the rotated geometry of the $[2Fe]_H$ -cluster is not regularly observed in model complexes, the sophisticated design of some model complexes enables the rotated geometry outside the enzyme. This finding emphasises, that the rotation is an intrinsic feature of [2Fe-2S]-clusters and not a unique enzyme-only geometry. In this context, the value of DFT calculations for biomimetic modelling aspects also becomes evident. The relevant electronic features for rotation identified by quantumchemical calculations namely electronic asymmetry, mixed-valency and strongly electron-accepting ligands (NO^+) are all found in synthetic models.

Though rare in Fe^IFe^I complexes, the rotated geometry is accessible in asymmetrically substituted and sterically crowded models. Agostic interactions were identified as a small but crucial contribution to obtain fully rotated complexes. In contrast, mixed-valence complexes obtained by external (Fc⁺) or internal (NO⁺) oxidants more regularly show a rotated geometry. These complexes show limited stability and high reactivity, *e.g.* towards BF₄⁻. Interestingly, some of these complexes do not only mimic the structure but also the reactivity towards exogenous CO and H₂ oxidation in terms of a PCET. Accordingly, these complexes are very powerful models of the H_{ox} state and make this state the most exactly represented by biomimetic modelling.

5.2 [2Fe-3S]-assemblies as H-cluster models

The introduction of a third sulphur ligand allows for more elaborate structural modelling of the active site – e.g. thioether coordination induces significant changes in the molecular structure.

In 2001, models with an additional sulphur donor on the dithiolate bridge were presented by the groups of Pickett¹³⁹ and Rauchfuss (Fig. 86).¹⁶⁹ In case of **50** and **285**, the coordination of the sulphur was observed after the introduction of the dithiol (*via* the thiol route). Contrary, Rauchfuss and coworkers isolated the hexacarbonyl complex **84** (*via* the salt-elimination) and induced sulphur coordination to form complex **286** by decarbonylation with Me_3NO . ^{366–368} A reasonable explanation

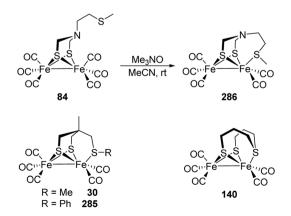


Fig. 86 [2Fe-3S]-assemblies carrying a thioether on the dithiolate bridge.

for the different behaviour of the adt- and pdt-derivatives is the enforced harsher conditions (90 °C (pdt) vs. -78 °C to r.t. (adt)), that could induce the thioether binding. Likewise, the different alkyl linkers between the bridgehead atom and the donor atoms as well as the different bridgehead atoms themselves, led to different binding behaviour due to different strain. Similar effects are also known from complexes comprising the Si/C exchange in tripodal ligands. 411 At last, the additional methyl group on the pdt-derivate renders the coordination more favourable and a comparable trend was observed for adtMeBH3 vs. adtBH3 complexes and attributed to the Thorpe-Ingold effect.175

The additional, hemilabile donor moiety in the bridging ligand has an immense effect on the substitution behaviour of the [2Fe-3S] assemblies (Fig. 87). In contrast to the hexacarbonyl complexes, the monocyanide complex is easily isolable from the reaction of 50 with cyanide. In fact, its formation is several orders of magnitude faster than in the unsubstituted complexes. Again, the thioether moiety is coordinated to an iron centre, but IR studies revealed that the complex $[Fe_2(\kappa^3-pdt^{MeSMe})-$ (CO)₅(CN)]⁻ (287) is formed initially. Similar results were reported for the introduction of P(OMe)₃ into complex 140.⁴¹² Herein, the pentacarbonyl intermediate is sufficiently stable to allow for its crystallisation. For the equilibrium between the two monocyanide complexes 287 and 288, as well as for the initial attack of 50 by a cyanide, a transition state with a bridging CO was proposed. The iron centre bearing the thioether is attacked by additional evanide to form a metastable species 289 with a bridging carbonyl ligand. Being stable at 0 °C, the complex slowly converts to complex 267 at room temperature. With large excess of cyanide present in solution, complex 267 is directly formed from 50.138,139,385,413

Complex 299 closely resembles the Hox-CO state in terms of the ligand environment of the iron centre but not in their oxidation states. Through transient IR and EPR spectroscopy of the product obtained upon chemical and electrochemical

electron oxidation of 267, formation of $[Fe_2(pdt^{MeSMe})(\mu-CO)-$ (CO)₃(CN)₂] was proposed. 379 With a comparable first coordination sphere, the same spin state and very similar IR bands with respect to the Hox-CO state, this complex was the first close resemblance of an actual state of the [FeFe]-hydrogenases. These spectroscopic similarities affirmed the assignment of the (unusual) oxidation states (Fe^IFe^{II}) in the active site. The spectroscopically and theoretically studies on [Fe₂(adt^{SMe})- $(CO)_4(L)_2^{+/-}$ (adt^{SMe} = (methylthio)ethylbis(sulfidomethyl)amine, L = PMe3, CN-) further supported the suggested structure - however, comprising a bridgehead nitrogen atom as an additional detail.^{280,414,415}

Notably, Tard et al. reported on complex 291, representing the hitherto only complete iron-sulphur framework of the H-cluster (Fig. 88). 416 Contrary to the natural H-cluster, the [2Fe-2S]- and the [4Fe-4S]-clusters are interconnected by an organic thioether moiety instead of a cysteine residue, completing the H-cluster framework. The formal exchange of the methyl group of 50 by a [4Fe-4S]-cluster results in a downshift of the IR bands of about 15 cm⁻¹. Electrochemical studies revealed that the [4Fe-4S]²⁺-cluster is reduced at milder potentials, than the [2Fe-2S]-cluster (compare H_{red} vs. $H_{red'}$, though the oxidation states of [2Fe]_H vary). While complex 291 revealed HER activity, despite its remarkable structural resemblance with the H-cluster, further in-depth studies on this model are quite limited due to the inherent instability of the H-cluster mimic. 416-418

In general, the electronic influence of the connection of the thioether to the dithiolate bridge is apparently small. From pdtand adt-hexacarbonyls to the thioether substituted models 50 and 286 the highest-energy IR frequency shifts by 26 cm⁻¹, respectively 21 cm⁻¹ to lower wavenumbers. 138,139,169,288 For the diethyl sulphide substituted pdt-complex a shift of 27 cm⁻¹ is observed.366 For discrete and linked metallothioethers shifts of 30 to 38 cm⁻¹ are reported. 178,204,209,416,419

If the unprotected trithiol CH₃C(CH₂SH)₃⁴²⁰ and its silasubstituted derivative¹⁵⁷ are reacted with triiron dodecacarbonyl,

Fig. 87 Proposed mechanism for the dicyanation of **50**. 138,139,385,413

Fig. 88 Synthesis of the H-cluster model 291

Fig. 89 Synthesis of complex 292E

the only isolable products are the tetra iron clusters 292E, in which two [2Fe-2S]-units are bridged by the two thiolate arms (Fig. 89). $^{420-422}$

All in all, as shown for **50**, an additional thioether strongly influences the reactivity of model complexes and enables the formation of the H_{ox} -CO mimic 267^+ with the complete first coordination sphere of the iron atoms in the enzyme.

5.3 Protonated H-cluster models

As a hydrogen forming catalyst, protonated states of the active site of [FeFe]-hydrogenases are an integral part of the catalytic cycle. Model complexes with bridging hydrides are readily formed upon protonation; 277,278,423,424 however, their relevance for the catalytic cycle was questioned and is still under discussion (Section 3.10). Terminal hydrides, though rare in mimics, are more appealing as key intermediates for rapid $\rm H_2$ formation due to their lower reduction potential, 154 their higher hydridic character, 270 and the proximity to the amine proton shuttle by the adt-ligand. 30

The reactivity of hydrogenase mimics towards other nucleophiles than protons and nitrosyls was also extensively studied. While giving bridged complexes for a variety of electrophiles, 425-435 terminal intermediates were reported as well. 433,434 In addition, other electrophiles, especially alkylation agents, also showed cyanide, 153,290 thiolate, 404,436 or carbonyl 53 centred reactivity as well.

5.3.1 Remarks on terminal and bridging hydrides. In terms of metal centred protonation reactions, mainly two different binding modes are observed, (apical) terminal hydrides and thermodynamically favourable bridging hydrides (Fe(μ-H)Fe). Notably, basal terminal hydrides also have been proposed as intermediates in isomerisation processes. 437 The two main binding modes are easily distinguishable by the hydride resonances in the ¹H-NMR spectra. Terminal hydrides cause a signal at ca. -5 ppm, ⁴³⁸ whereas bridging hydrides show resonances in a range of -12 ppm to -20 ppm. 423,424 These signals feature a characteristic coupling to the 31P nuclei of phosphine ligands. In the case of terminal hydrides, strong coupling ($J_{PH} \approx 75 \text{ Hz}$ chelating phosphines, 50 to 100 Hz monodentate) to phosphines ligated to the same iron centre is detected, 154,210,320,336,438 while bridging hydrides couple to phosphines in basal positions on both iron centres ($I_{PH} \approx 25$ Hz) and only weakly (5 Hz) to the apical phosphines. 277,278,423,424,438 Protonation of the iron core to give hydrides is accompanied by an average upshift of the IR bands by 60–110 cm⁻¹. 140,150,153,278,363

It is worth to mention that in the case of terminal hydrides, a μ -CO band is detected. The spectroscopic properties of many protonated complexes have been summarised by Tschierlei *et al.* and we would like to direct the reader to this review for more detailed information.

The structure of the bridging hydride complexes is very similar to the unprotonated complexes. The metal–metal distance is only slightly elongated (0.02 Å to 0.05 Å) upon protonation despite the loss of the Fe–Fe bond. In the face-sharing octahedrons the iron atoms are less displaced from the equatorial $S_2(CO)L$ planes compared to the displacement form the basal plane in the neutral complexes. 153,277,278,440,441 Importantly, the high fluxionality of the FeL3 units is lost upon protonation consistent with higher site exchange barriers in the octahedral coordination environment. 153,278

5.3.2 Iron-centred protonation of H-cluster models. The hexacarbonyl complexes are not basic enough to be protonated

by strong acids, e.g. HBF₄·Et₂O, but require harsh, super-acidic conditions (e.g. $[SiEt_3][B(C_6F_5)_4] + HCl_{(g)}$). 277,278,442,443 Contrary, the electron-donating abilities of a thioether in 50 are sufficient to allow for its protonation by HBF4·Et2O in dichloromethane. 140,444 In addition, protonation of dicyanide complexes gives several complexes with resonances corresponding to bridging hydrides in the ¹H-NMR spectrum, but cyanide ligand protonation and their intrinsic instability render these complexes unsuitable for protonation studies.²⁷⁷⁻²⁷⁹ However, the pdt dicyanide complex 212 reduced protons from strong acids under decomposition to sub-stoichiometric amounts of dihydrogen.²⁷⁹ Using only a small excess of acid, Pickett and coworkers were able to reveal the protonation process and suggested an initial protonation of the cyanide ligand with a subsequent rearrangement to afford a bridging hydride. 324,325 Notably, protection of the cyanides by boranes allows for the formation of stable protonated species bearing a bridging hydride.²⁹⁰ The formal exchange of one cyanide ligand by PMe₃ to give the complex $[Fe_2(pdt)(CO)_4(CN)(PMe_3)]^-$ (214) also allows for the isolation of a stable bridging hydride, while the corresponding P(OMe)₃ complex and the unsubstituted monocyanide are protonated at the cyanide. For the phosphine and the phosphite complex double protonation is observed with very strong acids. 150,153

Stable diphosphine complexes with bridging hydrides are known since the 1970s from early studies by Poilblanc, Mathieu and coworkers. 423,424,440 The protonation of Fe2(pdt)(CO)4-(PMe₃)₂ (219pdt) gave the first and representative example for a bridging hydride $[219pdt-\mu H]^+$ with a dithiolate bridge (Fig. 90). 277,278 Subsequently, stable bridging hydrides were obtained by protonation of a variety of electron-rich cofactor mimics. 270,278,326,342,353-355,445,446 The protonation behaviour of diphosphine bridged complexes Fe₂(pdt)(CO)₄(μ-diphosphine) shows remarkable dependence on the nature of the diphosphine. While the complexes with electron-poor, small-bite-angle phosphines dppm, dppe and (Ph2P)2NR are sluggishly protonated to give unstable bridging hydrides, complexes with the electron-rich

dcpm and the more flexible chelating phosphines form stable bridging hydrides. 326,342,353,354

Bridging hydride complexes are either not, or only slowly deprotonated by amine bases. 278,283,342 However, the less basic but smaller chloride ion deprotonates bridging hydrides and can also increase the deprotonation rate by amine bases when added sub-stoichiometrically (Fig. 93). 278,283,447

Protonated disubstituted phosphine and isocyanide complexes enable H/D exchange reactions between D2, D2O, alkenes, and Fe(μ-H)Fe under photolytic conditions. The reaction was proposed to proceed after dissociation of a carbonyl ligand, or a hydride shift to a single iron centre to provide a binding site for the substrate. This assumption is supported by the inhibition of the scrambling reactions by CO, acetonitrile, and in the case of D₂/D₂O by alkenes. ^{277,278,355,404}

After the first detection in the electrocatalytic proton reduction with a [2Fe-2P]-complex, 448 the first terminal hydride [216edt-tH] observed in a [2Fe-2S]-complex was synthesised in 2005 by Rauchfuss and coworkers. 438 In contrast to earlier studies, the complex was prepared by addition of a hydride source to the diferrous complex [Fe2(edt)(CO)2(PMe3)4(MeCN)]- $(PF_6)_2$ ([274edt]²⁺, Fig. 90). At low temperatures (-25 °C), this reaction yields a terminal hydride that was studied by NMR and IR spectroscopy and structurally characterised by X-ray diffraction. The isomerisation to the favourable bridging hydride [216edt-µH]⁺ occurs slowly at these temperatures (as well as in the solid state) but proceeds rapidly at room temperature. Notably, the corresponding complex with only three phosphine ligands showed no terminal hydride complex. Complexes [216xdt-tH]+ are also accessible by protonation of the corresponding complexes 216xdt. 320 The protonation of the edt and pdt analogues yields a mixture of bridging and terminal hydrides, while the protonation of 216adt only yields the terminal hydride. Though the terminal hydrides [216xdt-tH] isomerise to the bridging hydrides [216xdt-µH]⁺ at room temperature, terminal hydrides are no intermediates in the initial formation of the bridging hydride in the case of the edt

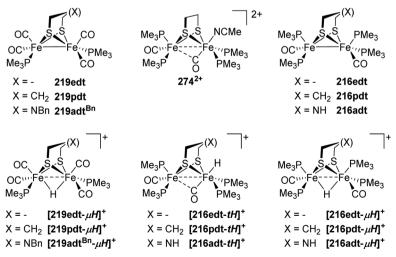


Fig. 90 Schematic structure of typical hydride complexes and their precursors

and pdt complexes. For the initial formation of the bridging hydride an intermolecular reaction from an S-protonated intermediate was proposed. Interestingly, $[216edt\text{-}tH]^+$ releases hydrogen upon treatment with strong acids, which is not always observed for terminal hydrides, e.g. $[216adt\text{-}tH]^+$ and $[\text{HFe}_2(\text{pdt})(\text{CO})_2(\kappa^2\text{-}d\text{ppv})_2]^+$ ($[233pdt\text{-}tH]^+)$ and generally not observed for bridging hydrides. 320,438,449

In contrast to the symmetric complexes 216xdt and 219xdt (xdt = edt, pdt), 320,324,325 terminal hydrides were proposed to be intermediates in the protonation of the asymmetric complexes $Fe_2(xdt)(CO)_4(\kappa^2-L)$ (κ^2-L = dppe, dmpe, dppv, phen, bis(NHC), NHC-PPh₂; xdt = edt, pdt), Fe₂(pdt)(CO)₃(PMe₃)(κ^2 -dppv), as well as in the sterically demanding complexes $Fe_2(xdt)(CO)_2(\kappa^2-dppv)_2$ (xdt = edt, pdt, adt, odt). 154,210,336,340,363,372,437,450 In contrast, the protonation of Fe₂(xdt)(CO)₄(κ^2 -dppp)³⁴² at -70 °C mainly yields bridging hydrides and for $Fe_2(edt)(CO)_{4-x}(PMe_3)_x(\kappa^2-dppv)$ (x = 0, 1)⁴³⁷ no terminal hydride is observed. Not only the dithiolate bridge but also the strength of the acid were reported to influence the occurrence of terminal hydride intermediates. 373,451 Admittedly, especially in cases where bridging hydrides isomerise quickly, distinguishing between a terminal hydride as a necessary intermediate and a terminal hydride as a (side) product that isomerises quickly is not trivial and requires elaborated experiments.

While the asymmetric terminal hydride complexes isomerise quickly at low temperatures ($-30~^{\circ}\text{C}$ to $-90~^{\circ}\text{C}),^{336,340,363,372,437}$ the sterically crowded terminal hydrides [HFe₂(xdt)(CO)₂(κ^2 -dppv)₂]⁺ [233xdt-tH]⁺ are reasonably stable at $-20~^{\circ}\text{C}.^{154,210,437}$ Unintuitively, protonation of the less electron-rich Fe(CO)₃ unit in the asymmetric complexes is regularly observed under these conditions. 336,340,363,437 A stable terminal hydride was reported for (Cp*)Fe(pdt)(μ -CO)Fe(κ^2 -dppe)H (293-tH), which only isomerises upon oxidation. Interestingly, the corresponding reduced bridging hydride partially isomerises to the terminal hydride – a process which is not observed for any other bridging hydride in context of hydrogenase mimics (Fig. 91). 452

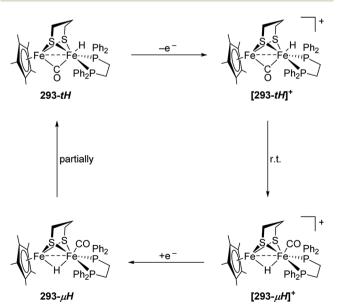


Fig. 91 Redox-Isomerisation of the terminal hydride [293-tH].

5.3.3 Influence of adjacent amines on the protonation of H-cluster models. The formal exchange of the bridgehead atom from pdt- to the basic adt-complex has, in contrast to the small influence on electronic and structural properties, an enormous influence on the protonation behaviour of hydrogenase mimics. Experimental observations highlight the advantages of the unique adt-ligand for the reversible proton reduction and, in this context, the relevance of nature's choice for the adjacent amine becomes obvious.

The adt^R amine is protonated by strong acids (e.g. HOTf, HBF₄/Et₂O), ^{151,168,177,183,196,203,346,354,447,453-459} while weaker acids (e.g. CF₃COOH, HOAc) are not sufficiently acidic to protonate the hexacarbonyl complexes in organic media. 195,343,460 Electron donating ligands on the iron core were reported to increase the basicity of the amine and allow for protonation with weaker acids, 210,320,451,459 while some hexacarbonyl complexes are only partially protonated by triflic acid or deprotonated in neutral solution. 454,459 Amine-functionalised phosphines show a similar protonation behaviour as the adt-bridge. 152,302,303,350-352,461 In the IR spectrum of the N-protonated complexes, the bands are shifted by ca. 15 cm⁻¹ under retention of the band structure. 459 The reversibility of the N-protonation was demonstrated with amine bases. 151,183,350,453-455 In electron-rich complexes bearing an adjacent amine, metal centred protonation is likewise observed. Importantly, in some cases the corresponding complexes lacking the amine are very slowly protonated or are only protonated by stronger acids. 210,320,351 In accordance with similar basicities, this effect is attributed to reduced kinetic barriers and is crucial for rapid H₂ formation. ^{154,210,350,351} Due to two basic sites, the protonation behaviour of these mimics can be complex and is affected by several parameters. While in some cases the amine only facilitates the protonation of the iron centre by decreasing the kinetic barrier and itself remains unprotonated, 154,462 mixtures of N- and Fe-protonated species 152,350,351 as well as solely N-protonated complexes were obtained. 151,447,455,459 The latter is observed if the amine is not in proximity to the site of metal protonation and can represent a metastable intermediate that slowly coverts to the bridging hydride. 151,455,459 This tautomerization can be accelerated by chloride. 455,459 Still, in some cases metal protonation is not observed at all. 447,463 Both amine and iron can be the more basic site and accordingly be the thermodynamically favourable site for protonation. If the basicities of both sites are similar, the equilibrium between the ammonium and the hydride tautomer is influenced by the solvent, 152,154,351,451 the used acid/counterion, 351,451 and the basicity of the amine.225,352

In general, more polar solvents stabilise the ammonium tautomer as well as the ability of the counterion to form hydrogen bonds. With strong acids double-protonated species are accessible, where the hydride can occupy a terminal or a bridging position. ^{151,154,320,447,451,455} It was shown that decreasing the basicity of the amine in Fe₂(pdt)(CO)₄((Ph₂PCH₂)NR) (294R) from 294Me to 294Ph is sufficient to prevent double-protonation by triflic acid (Fig. 92). This behaviour exemplifies the high acidity of the double-protonated complexes, which are sensitive to weak bases *e.g.* methanol, water or even

Fig. 92 Protonation and deprotonation pathways of the complexes 294R.

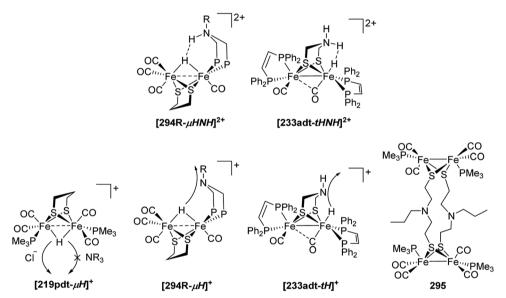


Fig. 93 Top: H-H interaction in double-protonated complexes. Bottom: Deprotonation of hydride complexes and structure of complex 295 (Note: P = PPh2 in 294R).

acetonitrile. 151,154,320,454,455 By NMR spectroscopy and singlecrystal XRD hydrogen-hydrogen interactions were suggested in these complexes (Fig. 93). 351,451

The deprotonation of the ammonium proton is typically fast, while the deprotonation rate of the hydride is dependent on the structure of the complex and the base used. In the complexes [294R-µH]⁺ and [233adt-tH]⁺ where the amine is in close proximity to the hydride, the deprotonation of the hydride can be accomplished with amine or phosphine bases (Fig. 93). The formal exchange of the NH groups by CH₂ groups prohibits any deprotonation, 210,351,464,465 while an oxygen atom showed a decreased, yet existent proton relay ability.210 In the case of 295 (Fig. 93), the iron protonated tautomer is metastable and rearranges to the N-protonated tautomer. If the hydride and the amine are spatially separated, the amine does not function

as a proton relay and deprotonation by amine bases is prohibited. 151,455 However, selective deprotonation by chloride was reported to be efficient.447

To emphasize the importance of the adjacent amine, we briefly summarize the different reactivities of the complexes shown in Fig. 94. The initial product of the protonation of 219adt is the ammonium salt and not the bridging hydride as for 219pdt^{151,277,278,324,325,455} and in contrast to 294R, 294C is only protonated by H(Et₂O)BF₄ if used in large excess. 342,351 While the protonation of 216adt and 233adt is feasible with medium strength acids as ammonium or phosphonium salts, the protonation of their pdt analogues requires strong acids.210,320 Accordingly, in the cases where the protonation site is in close proximity to the amine (216adt, 233adt, 294R), the protonation rate of the iron centre is accelerated. The same

Fig. 94 Exemplary complexes, whose reactivity towards protons is strongly altered by the adjacent amine

was observed for the deprotonation of the terminal hydride in [233adt-tH]⁺ and the bridging hydride in [294Me-µH]⁺, but not for [219adt-μH]⁺, where the bridging hydride and the amine are spatially separated. 210,351,455 At last, all amine containing complexes except from 294Ph allow for double-protonation to give a complex bearing an ammonium proton and a hydride.

5.3.4 Significance of the adjacent amine revealed by protonation studies. Protonation of electron-rich [2Fe]_H-mimics efficiently yields bridging hydrides accompanied by very small changes in the geometry of the [2Fe-2S]-core. However, their reactivity along with spectroscopic features on the enzyme questioned their relevance for rapid H2 formation. The more relevant terminal hydrides are formed as intermediates or metastable products both converting to the thermodynamically more stable bridging hydrides. In protonation experiments, the adjacent amine proved crucial for rapid proton transfer to and from the metal centre as well as for protonation with weak acids. These findings highlight the importance of the unprecedent adt-bridging ligand for the activity of the enzyme and fully explain the loss of activity upon incorporation of bridgeheadaltered [2Fe]_H-models.

VI Electrochemistry of H-cluster models - redox and catalytic properties

6.1 Redox chemistry of H-cluster models

Due to the mild potentials at which the native enzyme operates the hydrogen conversion/formation, models of its active site were extensively studied as noble-metal-free catalysts for the hydrogen evolution reaction. In order to evaluate potential catalysts, the redox behaviour of numerous complexes described in the previous sections was intensively investigated. In general, the interplay of the reduction potentials and the basicity of these clusters heavily influences the proton-coupled electron transfer processes that are crucial for HER.

Before describing the electrocatalytic capabilities of the active site models to serve as potent HER catalysts and the specific mechanisms involved, a short discussion on the noncatalytic redox behaviours of the diiron subsite models is advisable. Subsite models are generally in the Fe^IFe^I resting state and undergo reversible or quasi-reversible (stepwise)

reductions. Therefore, the complexes act as precatalysts and form the actual catalyst upon reduction. In the anodic scan, the mimics commonly display a one-electron oxidation resulting in a Fe^{II}Fe^I state followed by another electron oxidation resulting in a Fe^{II}Fe^{II} state. Herein the four most discussed complexes i.e. $Fe_2(pdt)(CO)_6$, $Fe_2(adt)(CO)_6$, $Fe_2(odt)(CO)_6$ and $Fe_2(sdt)(CO)_6$ will be considered. However, the oxidation only plays a minor role in the chemistry of hydrogenases and we will thus mainly focus on their reduction properties. In the following, all given potentials are referenced versus the ferrocene/ferrocenium couple if not otherwise specified.

Fe₂(pdt)(CO)₆ (20) displays a quasi-reversible single-electron reduction at $E_1 = -1.74 \text{ V} (-1.34 \text{ V } \nu \text{s. NHE})$ in MeCN and a second irreversible reduction is observed at a more negative potential of -2.35 V (-1.95 V vs. NHE). Notably, the reduction potential (E_1) for **20** has been reported with an averaged value of $-1.66 \text{ V.}^{283,289,302,317,329,367}$ The azadithiolate model Fe₂(adt)(CO)₆ (2) is reduced from a Fe^IFe^I to a Fe^IFe⁰ state at -1.58 V.^{170} This value is slightly less negative than the reduction potential of 20, which is reasonable due to the higher electronegativity of nitrogen compared to carbon and therefore a decreased electron density at the diiron core. Likewise, Fe₂(odt)(CO)₆ (136) undergoes a single-electron quasireversible reduction at -1.59 V and a further irreversible reduction at -2.1 V.²⁰⁴ Furthermore, the sulphur analogue $Fe_2(sdt)(CO)_6$ (139) is reduced to the Fe^IFe^0 stage at -1.51 V followed by reduction at -1.94 V to the Fe⁰Fe⁰ state.²¹¹ Thus, the bridgehead atom exerts only a small influence on the noncatalytic reduction behaviour of these models. Comparatively, models with aromatic thiolate linkers, especially Fe2(bdt)(CO)6 (149), are reduced at milder potentials (about -1.3 V). 221,467,468 The reduction comprises two overlapping single-electron processes. It is worth mentioning that the reduction of 149 proceeds with structural changes wherein one of the Fe-S bonds is cleaved, and a CO ligand transforms from a terminal to a bridging position. This geometric transformation causes a potential inversion, making the second reduction more feasible than the first one (Fig. 95). 468-471

6.2 Influence of modified thiolate bridges on the redox behaviour of H-cluster models

A common method to alter the redox-properties of metal centres is manipulation of their ligand environment and hence

Fig. 95 Scheme depicting the reduction of Fe₂(bdt)(CO)₆ based on spectroscopic data and DFT studies. 470-472

their electron density, which influences thermodynamics and kinetics of the redox-event. For C2-substituted Fe2(pdt)(CO)6like models, the inductive effect of the substituent is therefore decisive for the potential shift. Alkyl chains, that exhibit a +I effect, are expected to negatively shift the reduction potential due to the increasing electron density. However, models $Fe_2((SCH_2)_2CR_1R_2)(CO)_6$ (R₁ = R₂ = Me (296); R₁ = R₂ = Et (297); $R_1 = \text{Et}$, $R_2 = \text{Bu}$ (298)), bearing methyl-, ethyl- or butylgroups at the bridging position, undergo the first reduction at -1.61, -1.67 and -1.64 V, respectively.⁴⁷³ These values are within the reported range for the reduction of 20. Thus, inductive effects of alkyl-substituents at this specific position can be neglected. Contrary, substituents bearing an electron withdrawing group, should direct the reduction potential to more positive values due to the decreased electron density. Here, especially ketons, ¹³⁷ carboxylates ^{133,137,141} and alcohols ^{134–136,474} are present in literature. For example, complexes 35 and 38 are functionalized by a hydroxy group in C2 position and can be reduced at -1.61 and -1.60 V, respectively. Compared to the alkyl-substituted PDT-models and unsubstituted 20, the reduction potential is barely shifted to more positive values, which again shows the limited influence of inductive effects at the C2-position. 134,135 The same is true for ester bearing models 38, 43c and 43d that show a reduction potential of approx. -1.59 V. 133,137 The strongest shift in reduction potential (-1.53 V) is observed for model 39, which however, has kinetic reasons since the reduced state of this model is stabilized by an intermolecular H-bond between the hydroxy group and the reduced iron centre. 136 These few examples show that the influences of C2-substituents on the reduction potential of Fe₂(pdt)(CO)₆-like models is generally low and highlights the electronic remoteness of this position within the mimics.

Along this line, substituted azadithiolate models likewise possess reduction potentials between -1.49 V to -1.59 V (Section 6.5.2). Contrary to the pdt-models, more pronounced shifts in the reduction potential are observed upon modifying the mimics at the nitrogen atom. For example, alkyl substituted models are generally reduced at more cathodic potential (-1.63 V to -1.68 V). ^{182,186,190} Furthermore, strongly electron withdrawing systems attached to a phenyl-ring such as in $Fe_2(adt^R)(CO)_6$ (R = p-C₆H₄NO₂, **120**) cause a reduction potential shift to -1.42 V. 194

In contrast to Fe₂(pdt)(CO)₆ and Fe₂(adt)(CO)₆ models, the aryl-substituents of Fe₂(bdt)(CO)₆ have a severe influence on the redox potentials of the hydrogenase mimics (Section 6.5.6). 222,228,232,239,240,242 Notably, bdt-models comprising

ligands with +I effect are rare in literature and show only slightly increased electron density at the diiron centre. Complex 153 is a good example for such a system and possesses a single methyl group in m/p-position. The complex shows a reduction potential of -1.37 V compared to -1.36 V for 149. Attachment of electron withdrawing groups, especially chlorides, reduce the electron density at the diiron core to a significant extent. The most prominent effect can be observed in 152 with a per-chlorinated benzene ring and exhibits a reduction potential of -1.13 V in MeCN. 226 Moreover, models with N-heterocyclic substituents (e.g. 2,3-quinoxalinedithiolate) are more conveniently reduced owing to the decreased electron density on the diiron centre due to the electron withdrawing N-substituted aryl rings (Section 6.5.6). 222,228,229

In contrast to Fe₂(bdt)(CO)₆ and its analogues, 1,8naphthalenedithiolate models mainly undergo two singleelectron reductions (Section 6.5.7). 239,242,244,247,251 The first reduction usually occurs at -1.52 V, which is approx. 200 mV more negative compared to Fe₂(bdt)(CO)₆.²³⁹ Electron density manipulating groups alter the redox potential in the same way as described above for bdt-derivatives but to a somewhat lesser extent.

The biphenyl⁴⁷⁵ as well as the o-carborane⁴⁷⁶ modified Fe₂S₂ complex also undergoes two successive single step electron reductions. Moreover, due to the delocalisation of the negative charge on the aromatic ring and rigidity of the naphthalenelinker these models prove to generate more stable reduced states. In addition, the phenanthrene-4,5-dithiolate-bridged compound 182 is reduced at more positive potentials due to the enhanced electron delocalization of the phenanthrene system.240

Despite of the increased electron density at the diiron centre in chalcogenide-substituted complexes, anodically shifted reduction potentials upon sulphur to chalcogenide exchange were observed (e.g. 39 vs. 40). 133,136,145,146,204,205,211,239,240,251,477 This positive shift is overall < 50 mV and can be rationalised by the improved stabilization of the reduced species. Furthermore, Weigand and coworkers further reported decreasing reorganization energies for the reduction to a Fe^IFe⁰-species within the S, Se and Te series that partially counteracts the trend of increasing electron density at the diiron centres. 146

In contrast, Fe₂((ECH₂)N-p-C₆H₄R)(CO)₆ derivatives show almost no shift in the reduction potentials for the respective selenium containing models (R = H, E_{128} = -1.57 V vs. E_{128} = -1.58 V; R = NO₂ E_{130} = -1.48 V vs. E_{120} = -1.49 V). ¹⁹⁸ Along this line, it is worth to mention that the reduction of 2 (adt) and 96

(adSe) occurs at almost same potentials of approx. $-1.2 \text{ V} \nu s$. SHE. ³² Thus, the experimentally observed shifts are depending on the interplay of the electron density modulating properties of the headgroup and the chalcogenide as well as the counteracting change of reorganisation energies for the reduction.

6.3 Influence of CO-ligand substitution on the redox behaviour of H-cluster models

While the carbonyl to cyanide replacement aims at replicating systems that resemble the natural subsite more closely, the complicated electronic behaviour of the cyanide restricts their investigations; in this regard many studies adopted phosphines and carbenes as non-native ligand systems. Herein, it is necessary to highlight that the redox behaviour of the metal centres are considerably affected by the nature of the binding ligand as the LUMO has major contributions of metal–metal and metal–ligand anti-bonding orbitals. 138

6.3.1 Cyanide and isocyanide substituted models. The first irreversible reduction of the dianionic dicyanide model $[Fe_2(pdt)(CO)_4(CN)_2]^{2-}$ (212) takes place at -2.73 V and the irreversible oxidation takes place at -0.51 V. 289 Contrarily, $[Fe_2(pdt)(CO)_5(CN)]^-$ (213) displays its first irreversible reduction at -2.17 V and the oxidation takes place at +0.13 V. 283 Therefore, compared to $Fe_2(pdt)(CO)_6$ (20), successive CO to CN $^-$ exchange causes a 0.51 V to 0.59 V cathodic shift of E_{pc} due to the increasing electron density at the metal centres owing to the strong electron donating nature of the CN $^-$ ligand and thereby making them harder to reduce. 138,283,289,478 Replacement of CN $^-$ ligands with weaker electron donating ligands such as methyl isocyanide results a moderate ΔE_{pc} of -0.15 V upon single CO-replacement. Along this line, the second substitution proceeds with a further shift of -0.27 V. 283

6.3.2 Phosphine or phosphite substituted models. As a general trend, the replacement of one CO ligand by a phosphine or phosphite results in cathodic shifts of the reduction potential $E_{\rm pc}$ of about 0.12 to 0.3 V. Herein, the electron density at the metal centre is severely increased due to σ-donation and decreased back donation of the phosphine or phosphite. ^{121,170,195,215,222,229,317,473,479–481} Comparing PPh₃ and PMe₃ substituted mimics, the former are reduced at less cathodic potentials (Table 7), which can be attributed to the weaker electron donating capability of the PPh₃ ligand. ^{170,302,317} Notably, when a second CO ligand is replaced, $E_{\rm pc}$ further shifts by approx. 0.2 V to 0.46 V to more negative

potentials. 170,182,191,195,317,454,479 Thus, the second ligand exchange results in in case of two PMe $_3$ ligands (219pdt) in a reduction potential of -2.31 V, which is notably 0.4 V less negative as for 212. In case of monosubstituted phosphine models of 149, the cathodic scan shows a stepwise two-electron reduction at more cathodic potentials, *e.g.* approx. 0.2 V for a single PPh $_3$ ligand in acetonitrile and contradicts the concerted two-electron transfer found in the hexacarbonyl complexes. This behaviour was reasoned by slower electron transfer kinetics of the monoanion in the substituted models. 229,480,482

The alteration of $E_{\rm pc}$ for chelating phosphine ligands is comparable to the shifts observed for mimics possessing two monodentate phosphines. The reduction potential of Fe₂(pdt)CO₄(κ^2 -dppe) (232pdt) is with -2.33 V similar to that of Fe₂(pdt)CO₄(PMePh₂)₂ (299) with -2.30 V. Likewise, analogues with bridging phosphine ligands reveal comparable reduction potentials as their chelating counterparts. The reduction potentials are the reduction counterparts.

6.3.3 NHC substituted models. Much like phosphine substituted complexes, carbene substituted models show similar trends, i.e. cathodic shifts of the reduction potential. However, the influence of the carbene ligand is much stronger than reported for phosphines. 197,291,361,362,364,473 Herein, upon each CO to NHC exchange, potential shifts of up to 0.44 V are observed. For example, the E_{pc} of $Fe_2(pdt)(CO)_{6-n}(IMe)_n$ (n = 1(300), 2 (247)) is observed at -2.06 V and -2.47 V, respectively, and is thus more cathodic as compared to the phosphine (PMe₃) analogues (218pdt & 219pdt) that exhibit the respective reduction signals at -1.94 V and -2.31 V.291,302 Aromatic N-substituents at the imidazole result in less negatively shifted reduction potentials of the respective mono-substituted complexes, e.g. -2.01 V in the IMes ligand. Interestingly, the potential change caused by the IMe and IMes ligand, is smaller than the difference observed for PPh3 vs. PMe3 substituted compounds. This difference shows that the effect of the additional substituents of imidazole based NHCs is not as significant as in phosphines (Table 8). As found for IMes substituted complexes, furan and pyridine modified disubstituted carbene models are reduced at potentials 0.95 V more cathodic than the hexacarbonyl models. 197

6.3.4 Models substituted with other ligands. The natural [2Fe–2S]-cofactor is not only coordinated by two cyanide and four CO ligands, but also linked to the [4Fe–4S]-cluster *via* a cysteine that forms a thioether with the cubic cluster. Therefore,

Table 7 $E_{pc}(Fe^{l}Fe^{l}/Fe^{l}Fe^{0})$ of phosphine/phosphite-substituted models

Complex	Ligand	$E_{\text{pc,mono}}$ [V] $n = 1$	$E_{\text{pc,di}}$ [V] $n=2$	ΔE vs. CO [V]	Ref.
Fe ₂ (pdt)(CO) ₆		-1.66			215, 283, 289, 302, 317, 329 and 367
$Fe_2(pdt)(CO)_{6-n}(L)_n$	PMe_3	-1.94	-2.31	0.28, 0.37	221, 467 and 468
$Fe_2(pdt)(CO)_{6-n}(L)_n$	PMe ₂ Ph	-1.90	-2.30	0.24, 0.40	283
$\operatorname{Fe_2(pdt)(CO)}_{6-n}(L)_n$	PPh_3	-1.84		0.18	291
$Fe_2(pdt)(CO)_{6-n}(L)_n$	$P(OEt)_3$	-1.81	-2.27	0.15, 0.46	473
$Fe_2(pdt)(CO)_{6-n}(L)_n$	$P(OMe)_3$	-1.98	-2.30	0.32, 0.32	473
$Fe_2(pdt)(CO)_{6-n}(L)_n$	PTA	-1.94	-2.14	0.28, 0.20	473
$Fe_2[(SCH_2)_2(NH)](CO)_6$		-1.58			473
$Fe_2(adt)(CO)_{6-n}(L)_n$	PMe_3	-1.88		0.30	170
$Fe_2(adt)(CO)_{6-n}(L)_n$	PPh_3	-1.70		0.12	170

Table 8 $E_{pc}(Fe^{l}Fe^{l}/Fe^{l}Fe^{0})$ of carbene, cyanide and isocyanide-substituted models

Complex	Ligand	$E_{\text{pc,mono}}[V] n = 1$	$E_{\mathrm{pc,di}}\left[\mathbf{V}\right]n=2$	ΔE vs. CO [V]	Ref.
$Fe_2(pdt)(CO)_{6-n}(L)_n$		-1.66			215, 283, 289, 302, 317, 329 and 367
$[Fe_2(pdt)(CO)_{6-n}(L)_n]^{n-}$	CN^-	-2.17	-2.72	0.51, approx. 0.55	221, 467 and 468
$Fe_2(pdt)(CO)_{6-n}(L)_n$	CNMe	-1.81	-2.08	0.15, 0.27	283
$Fe_2(pdt)(CO)_{6-n}(L)_n$	IMe	-2.06	-2.47	0.40, 0.41	291
Fe ₂ (pdt ^{Me})(CO) ₆		-1.61			473
$\operatorname{Fe_2(pdt^{Me})(CO)_{6-n}(L)_n}$	IMes	-2.01		0.40	473
$Fe_2(pdt^{Et})(CO)_6$		-1.67			473
$\operatorname{Fe_2(pdt^{Et})(CO)_{6-n}(L)_n}$	IMes	-2.02		0.35	473

Table 9 $E_{pc}(Fe^{l}Fe^{l}/Fe^{l}Fe^{0})$ and $E_{pa}(Fe^{l}Fe^{l}/Fe^{l}Fe^{l})$ of **50** and **285** and their respective cyanides¹³⁸

Complex	$E_{ m pc}$ vs. SCE (vs. Fc ^{+/0}) [V]	E_{pa} vs. SCE (vs. Fc ^{+/0}) [V]
$\begin{split} & Fe_2(pdt^{MeSMe})(CO)_5 \\ & Fe_2(pdt^{MeSBn})(CO)_5 \\ & [Fe_2(pdt^{MeSMe})(CN)(CO)_4]^- \\ & [Fe_2(pdt^{MeSBn})(CN)(CO)_4]^- \\ & [Fe_2(\mu\text{-}CO)(pdt^{MeSMe})(CN)_2(CO)_3]^{2-} \\ & [Fe_2(pdt^{MeSMe})(CN)_2(CO)_4]^{2-} \\ & [Fe_2(pdt^{MeSMe})(CN)_2(CO)_4]^{2-} \\ & [Fe_2(pdt^{MeSBn})(CN)_2(CO)_4]^{2-} \end{split}$	-1.38 (-1.78) -1.36 (-1.76) -1.83 (-2.23) -1.83 (-2.23) -2.40 (-2.83) -2.40 (-2.83) -2.40 (-2.83)	+0.67 (+0.27) +0.77 (+0.37) +0.17 (-0.23) +0.12 (-0.28) -0.10 (-0.03) -0.25 (-0.65) -0.26 (-0.66)

thioether and sulphoxide substituted models were synthesized to mimic the properties of the native cysteine. 138,139,367 Thioethers generally were shown to cause cathodic shifts of up to 100 mV for the reduction of Fe^IFe^I (Table 9). Contrary, sulphoxides induced no noteworthy shift of the respective potential due to the competing inductive effects of the oxide and alkyl chains.³⁶⁷ Along this line, the incorporation of a strong electron accepting ligand, such as NO^+ , lead to a 1 V cathodic shift of E_{pc} as compared to the related hexacarbonyl complexes. 150,153,381

6.4 Remarks on the electrochemical oxidation of H-cluster models

As described above, the electron donating capabilities of the ligands effect the complexes reduction potentials. Analogously, the oxidation processes are likewise affected by ligand exchanges. Thus, strong donor ligands such as PMe3 and CN^- e.g. cause a cathodic shift of E_{pa} (Table 10). Importantly, the influence of the ligand substitution is greater on the oxidation potentials than on the reduction potentials, a trend

that can be explained by different HOMO-LUMO participation. As oxidation involves removal of electrons from the HOMO, which usually has a strong contribution of the Fe-Fe bond, attaching strong donor ligands to the iron atoms directly eases this process. However, in case of any reduction, the less metalcentred LUMO with a significant Fe-S bond character is involved and renders this process comparatively less sensitive to ligand exchange.138

The first oxidation of [FeFe]-hydrogenase mimics leads to bivalent Fe^IFe^{II} species, which is related to the H_{ox} state of the native active site. The effect of the dithiolate bridge on the stability of the oxidised state was reported by Justice et al. investigating Fe₂(xdt)(CO)₃(PMe₃)(dppv) (235xdt, xdt = edt, pdt, adt).337 Upon chemical oxidation with Cp2FePF6 in MeCN, rotated states mimics with bridging CO and NCMe coordination at the "open site" resulting in [Fe₂(xdt)(CO)₂(μ-CO)(PMe₃)- $(dppv)(NCMe)^{2+}$ (235xdt(MeCN)²⁺) was observed. Notably, the dicationic state of edt was found to be stable at room temperature, while the pdt bridged model decomposed within 30 minutes at -40 °C. Remarkably, no dicationic species was yet detected for any adt analogue. Based on these observations, it was concluded that bulky ligands, e.g. PMe3 are not accommodated by pdt and adt linkers in the rotated state, due to steric limitations which are not present in edt-bridged derivative, thereby explaining the preference of nature to apply sterically less demanding CN- and CO as ancillary ligands. Adding substituents to the amine bridgehead allows for the facile oxidation of such complexes as seen in e.g. Fe₂(adt^{Bn})-(CO)₄(dmpe) (265). The described reason for the stabilization of the rotated Fe^IFe^{II}-oxidation state is an anagostic interaction between the methyl group of the Bn substituent and the

Table 10 $E_{pa}(Fe^{\dagger}Fe^{\dagger}/Fe^{\dagger}Fe^{\dagger})$ and shift of the peak potentials compared to the corresponding hexacarbonyl ΔE vs. (CO)₆ of various substituted models

Complex	Ligand	$E_{ m pa}\left[{ m V} ight]$	ΔE vs. $(CO)_6$ $[V]$	Ref.
Fe ₂ (pdt)(CO) ₆		+0.80		
[Fe ₂ (pdt)(CO) ₅ (CN)] ⁻	CN^-	+0.13	0.67	283
$[Fe_2(pdt)(CO)_4(CN)_2]^{2-}$	CN^- , CN^-	-0.52	0.66	289
Fe ₂ (pdt)(CO) ₅ (CNMe)	CNMe	+0.57	0.17	283
$Fe_2(pdt)(CO)_4(CNMe)_2$	CNMe, CNMe	+0.21	0.42	283
Fe ₂ (pdt)(CO) ₅ (PMe ₃)	PMe_3	+0.23	0.57	317
$Fe_2(pdt)(CO)_4(PMe_3)_2$	PMe ₃ , PMe ₃	-0.20, -0.14	0.43, 0.37	153 and 215
$Fe_2(pdt)(CO)_5(P(OEt)_3)$	$P(OEt)_3$	+0.44	0.36	317
$Fe_2(pdt)(CO)_4(P(OEt)_3)_2$	$P(OEt)_3, P(OEt)_3$	0.00	0.44	317
$Fe_2(adt)(CO)_6$		+0.59		170
$\operatorname{Fe}_{2}[(\operatorname{adt})(\operatorname{CO})_{5}(\operatorname{CN})]^{-}$	CN^-	-0.043	0.63	478
Fe ₂ (adt)(CO) ₅ (PPh ₃)	PPh_3	+0.19	0.40	170

Fig. 96 Structural reorganisation of 265 upon oxidation.

iron centre (Fe···H-C), which also allows its facile second oxidation. It was also observed that these weak interactions are lost under CO atmosphere. In this case, the [Fe^{II}Fe^I(adt^{Bn})-(CO)₄(dmpe)]⁺ is stabilized by binding of an additional carbonyl ligand to an iron centre, rather than an anagostic stabilisation (Fig. 96).

6.5 Redox potentials of selected H-cluster models

While the above-mentioned trends are explained on distinct examples, comparable trends can be observed for numerous synthetic mimics. The large number of mimics would certainly distract the reader from the actual message and trends observed. However, to present a full picture and allow the reader to obtain the formation without a detailed literature research, we subsequently present the redox properties of numerous complexes in tabular form (Tables 11-23).

6.5.1 PDT and derivatives thereof

The tabular section can be summarized as follows:

- Fe₂S₂ models undergo electrochemical reductions resulting in a Fe⁰Fe⁰ state which might be achieved in concerted twoelectron transfer steps or require two single-electron potentialseparated reductions.
- Functionalisation of the dithiolate bridge has only minimal influence on the redox potentials.
- Upon CO-replacement, electron donating ligands shift both, the reduction and oxidation potential, towards more cathodic values. The opposite is true for electron withdrawing ligands. However, the oxidation potential is more affected due to the nature of the HOMO and LUMO orbitals.

These investigations of the diiron models in the absence of acids give a fair idea about their respective redox properties, which can be mainly correlated to the basicity of the Fe-Fe bond. This basicity is an important factor for the catalytic mechanism of hydrogen evolution, which will be discussed in Section 6.6.

Table 11 Mono-substituted complexes

L_1	Solvent	$E_{\rm pc}({\rm Fe^IFe^I/Fe^IFe^0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^{II}})^a/{\rm V}$	Ref.
CO CO	MeCN THF	-1.65 to -1.74 -1.25^b	-2.32	0.74 to +0.82	215, 283, 289, 302, 317, 329 and 367 283

Table 11 (continued)

			L ₁ CO		
L_1	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm II})^a/{\rm V}$	Ref.
PTA	MeCN	-1.94^{c}		+0.34	302
$P(NC_4H_8O)_3$	MeCN	-1.86		$+0.27^d$	129
(-4 8 -)3	MeCN/H ₂ O 10:1				
	$MeCN/H_2O 3:1$	-1.80			
	MeCN/H ₂ O 2:1	-1.78			
$(PPh_2)NH((CH_2)_2NMe_2)$. 2	-1.87		+0.348	303
(PPh ₂)NH(o-C ₆ H ₄ NH ₂)		-1.86		+0.259	303
PPh_2 -(C ₆ H ₄)-2CH ₂ N(Me) ₂		-1.85			303
Val-Epa-Leu	MeCN	-1.93		+0.22	306
-	$MeCN/H_2O 3:1$	ND			
	$MeCN/H_2O3:2$	ND			
Val-Ipa-Leu	MeCN	-1.90		+0.26	306
-	$MeCN/H_2O3:1$	-1.78		+0.22	
	$MeCN/H_2O 3:2$	-1.74		+0.22	
Val-Ppa-Leu	MeCN	-1.82		+0.31	306
•	$MeCN/H_2O3:1$	-1.77		+0.29	
	$MeCN/H_2O 3:2$	-1.72		+0.32	
$PPh_2(S(o-C_6H_4NH_2))$	MeCN	-1.79	-2.06		307
$PPh_2(p-C_6H_4NH_2)$	MeCN	-1.85	-2.10		307
$PPh_2(S(o-C_6H_4NH_2-L_a^e))$	MeCN	-1.76		+0.34	307
$PPh_2(p-C_6H_4NH_2-L_a^e)$	MeCN	-1.83		+0.36	307
PMe ₃	MeCN	-1.94^{f}		+0.31, +0.65	317
PMe ₂ Ph	MeCN	-1.90^{f}		+0.25, +0.66	317
PPh ₃	MeCN	-1.84^{f}		+0.26, +0.62	317
3		-1.87	-2.24	+ 0.27, +0.66	215
$P(OEt)_3$	MeCN	-1.81^{f}		+0.44, +0.63	317
PPh ₂ Py		-1.74		+0.46, +0.68	319
PPy ₃		-1.66		+0.65	319
$PPh_2NH(p-C_6H_4Br)$	MeCN	-1.80			334
$PPh_2NH(p-C_6H_4Me)$	MeCN	-1.82			334
$PPh_2(R_1)^g$	DCM	-2.05			305
$PPh_2(R_2)^g$	DCM	-2.02	-2.17		305
$PPh_2(R_3)^g$	DCM/DMF 1:4	-1.83	-2.07		305
IMe	MeCN	-2.06/-2.01		+0.11	291 and 361
IMes	MeCN	-2.10		+0.11	361
IMes		-2.10^{c}		+0.11, +0.72	362
IMeMes	MeCN	-2.12		+0.23	361
IPic^h	MeCN	-2.07^{f}		+0.10	364
SEt ₂	MeCN	-1.72^{i}			367
S(CH ₃ CH ₂)((CH ₂) ₂ Cl)	MeCN	-1.76^{i}			367
$S(CH_2CH_3)(C_6H_5)$	MeCN	-1.77^{i}			367
SO(CH ₂ CH ₂ CH ₃) ₂	MeCN	-1.65^{i}			367
$SO(CH_3)_2$	MeCN	-1.68^{i}			367
MeCN	MeCN	-1.68			369
NH ₂ n-Pr	MeCN	-1.80			369
PySEt ^j	MeCN	-1.65^{i}	-2.22		370
P(piperidyl) ₃	MeCN	-1.98		+0.18	484
$P(OMe)_3$	MeCN	-1.98	-2.29	+0.37	485
DAPTA	$MeCN^{l}$	-1.83^{c}	-		486
CNMe	MeCN	-1.81^{i}		+0.57	283
CN-	MeCN	-2.17^{i}		+0.13	283
		•			

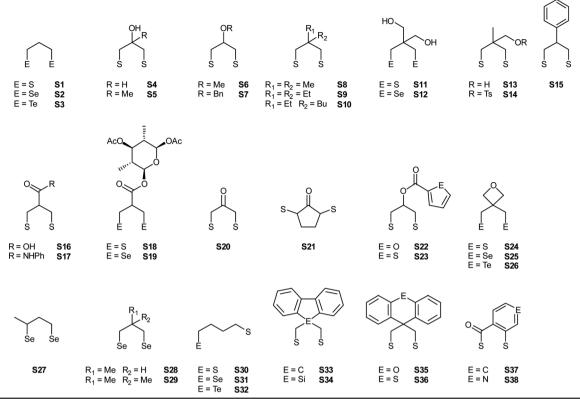
 $[^]a$ Second oxidation corresponds to Fe^IFe^{II}/Fe^{II}Fe^{II}. b Reported vs. Ag/AgCl. c Reported vs. NHE, converted. d Electron count not defined. e L_a = CO(bpy)(ppy)₂Ir. f Reported vs. 0.01 M Ag/Ag(NO₃), converted. g R₁ = 6-(diphenylphosphaneyl)pyridin-2-yl, R₂ = 6-(diphenylphosphaneyl)pyrimidin-4-yl, R₃ = 6-(tert-butyl)pyrimidin-4-yl. h 1,3-Bis(2-picolyl)imidazol-2-ylidene. i Reported vs. Ag/AgCl, converted. j 4-(Ethylthio)pyridine. k DAPTA = 3,7-diacetyl-1,3,7-triaza-5-phosphabicyclo[3.3.1]nonane. l Reference for redox potentials in different H₂O:MeCN ratios.

Table 12 Multi-substituted complexes

In the set of the set o		L ₁ ,L ₂ //L ₃	κ²-geometry	μ-geometry		
$\begin{split} & \text{PinC4}_{+}(p_0)_{j}/ \text{PinC4}_{+}(p_0)_{j} \\ & \text{MeCN}^{\dagger} & -2.08 \\ & \text{MeCN}^{\dagger} & -2.10 \\ & \text{MeCN}^{\dagger} & -2.10 \\ & \text{MeCN}^{\dagger} & -2.09 \\ & \text{MeCN}^{\dagger} & -2.10 \\ & \text{MeCN}^{\dagger} & -2.20 \\ & \text{MeCN}^{\dagger} & -2.27 \\ & \text{MeCN}^{\dagger} & -2.28 \\ & \text{MeCN}^{\dagger} & -2.23 \\ & \text{MeCN}^{\dagger} & -2.28 \\ & \text{MeCN}^{\dagger} & -2.28 \\ & \text{MeCN}^{\dagger} & -2.21 \\ & \text{MeCN}^{\dagger} & -2.23 \\ & \text{MeCN}^{\dagger}$	Ligands: Fe ₁ //Fe ₂	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm O}/{\rm Fe}^{\rm O}{\rm Fe}^{\rm O})/{\rm V}$	$E_{\rm pa}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm II})^{a}/{\rm V}$	Ref.
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	PTA//PTA	MeCN	-2.18^{b}		+0.00	302
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	$P(NC_4H_8O)_3//P(NC_4H_8O)_3$		-2.08		+0.01 ^c	129
μ-(PPh_2) ₂ N-(CH ₂) ₂ NMe ₂ BzBr McCN	κ^2 -(PPh ₂) ₂ N-(CH ₂) ₂ NMe ₂ BzBr	$MeCN^d$	-2.10		+0.06	130
McON		$MeCN: H_2O(3:2)$	-1.92		+0.06	
McON	μ-(PPh ₂) ₂ N-(CH ₂) ₂ NMe ₂ BzBr	$MeCN^d$	-2.09		+0.40	130
μ-dppf MeCN	. (-)- (-)		-1.93		+0.42	
$ P(OEb_{)}/P(OEb_{)} P(PoEb_{)}) P(Ph_{2}Py_{)}/P(Ph_{2}Py_{)} P(Ph_{2}Py_{)}/P(Ph_{2}Py_{)} P(Ph_{2}Py_{)} P(P$	μ -dppf ^e		-2.10	-2.19	+0.05	311
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	PMe ₂ Ph//PMe ₂ Ph	MeCN	-2.30^{f}		-0.14, +0.20	317
$\begin{split} \text{pip}_{j} $	$P(OEt)_3//P(OEt)_3$	MeCN	-2.27^{f}		+0.00	317
$\begin{array}{c} \mu \text{dppm} \\ \mu \text{dppm} \\ \text{MeCN} \\ -2.28^f \\ \text{MeAgdppm} \\ \text{MeCN} \\ -2.16^g \\ -2.23 \\ \text{Ho.} 19, +0.04 \\ 328 \\ \text{Ho.} 24, +0.60 \\ \text{Ho.} 13 \\ 328 \\ \text{Ho.} 19, +0.14 \\ 328 \\ \text{Ho.} 19, +0.14$	$P(Ph_2Py)_3//P(Ph_2Py)_3$		-1.92		+0.32, +0.72	319
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$P(Py)_3//P(Py)_3$		-1.70		+0.62	319
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	μ-dppm	MeCN	-2.28^{f}		+0.22, +0.60	183
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	κ²-Me ₂ dppm	MeCN	-2.16^{g}	-2.23	-0.19, +0.04	328
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	μ-Me ₂ dppm	MeCN	-2.50		+0.74	328
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	μ-(PPh ₂) ₂ NPr	MeCN	-2.17	-2.46	+0.33	329
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	κ^2 -(PPh ₂) ₂ N(allyl)	MeCN	-2.19		-0.11	330
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$		DCM	-2.23		+0.07	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	μ-(PPh ₂) ₂ N(allyl)	MeCN	-2.15		+0.31	330
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$		DCM	-2.23		+0.65	
$ \begin{array}{c} \Gamma \\ \mu \text{-dppe} \\ \mu \text{-dppe} \\ \text{HeCN} \\ \text{THF} \\ -2.23 \\ \text{THF} \\ -2.37 \\ \\ \text{IMes//PMe_3} \\ \text{MeCN} \\ -2.47 \\ -2.36 \\ -2.36 \\ -0.47 \\ 361 \\ \text{IMes/PMe_3} \\ \text{MeCN} \\ -2.52 \\ -0.33 \\ 361 \\ \text{IMe/Mes/PMe_3} \\ \text{MeCN} \\ -2.52 \\ -0.33 \\ 361 \\ \text{IMe/Mes/PMe_3} \\ \text{MeCN} \\ -2.52 \\ -0.33 \\ 361 \\ \text{IMe/PMe_1} \\ \text{MeCN} \\ -2.53 \\ -0.24 \\ 361 \\ \text{Meron PMe_2 MeCN} \\ -2.42^h \\ -0.41 \\ 363 \\ \text{κ}^2 \text{-NHCMePy} \\ \text{MeCN} \\ -2.42^h \\ \text{-0.16} \\ \text{-0.16} \\ \text{-0.25} \\ \text{-0.25} \\ \text{-0.27} \\ \text{-0.27} \\ \text{-0.27} \\ \text{-0.29} \\ \text{-0.20} \\ -0.$	κ^2 -(PPh ₂) ₂ N(p-C ₆ H ₄ Me)	MeCN	-2.21			334
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	κ²-dppe	MeCN	-2.07^{g}			338
THF		THF	-2.12			
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	μ-dppe	MeCN	-2.23			338
$\begin{array}{cccccccccccccccccccccccccccccccccccc$		THF	-2.37			
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	IMe//IMe	MeCN	-2.47			291
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	IMes//PMe ₃	MeCN	-2.36		-0.47	361
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	IMeMes//PMe ₃	MeCN	-2.52		-0.33	361
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	IMe//PMe ₃	MeCN	-2.53		-0.24	361
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$		MeCN	-2.42^{h}		-0.41	363
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$				-0.16		
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	κ²-bpy		-2.06			371
$\begin{array}{cccccccccccccccccccccccccccccccccccc$		DCM		-0.25		
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$P(OMe)_3//P(OMe)_3$	MeCN	-2.30			
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	PMe ₃ //PMe ₃	MeCN			-0.20, -0.14	153 and 215
$\begin{array}{cccccccccccccccccccccccccccccccccccc$						
$\begin{array}{cccccccccccccccccccccccccccccccccccc$				-0.98		381
$\begin{array}{cccccccccccccccccccccccccccccccccccc$						152
$\begin{array}{cccccccccccccccccccccccccccccccccccc$						487
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	PMe ₂ Ph//PPh ₃				+0.03	487
$\begin{array}{cccccccccccccccccccccccccccccccccccc$						
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$PMe_3//P(OEt)_3$				+0.03	487
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$PPh_3//P(OEt)_3$					487
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$PCy_3//P(OEt)_3$					487
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	0 0					
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	5 55				-0.02	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$			-2.06^{p}			
CNMe//CNMe MeCN -2.08^k $+0.21$ 283 CN^- //CN $^-$ MeCN -2.72^m -0.50 289			-2.14^{b}_{L}			486
$\text{CN}^-//\text{CN}^-$ MeCN -2.72^m -0.50 289	-		-2.58^{κ}_{L}			
$\text{CN}^-//\text{CN}^-$ MeCN -2.72^m -0.50 289	CNMe//CNMe	MeCN	-2.08^{κ}		+0.21	283
$CN^{-}//CN^{-}$ MeCN -2.72^{κ} -0.52 283			-2.72^{m}			
	CN ⁻ //CN ⁻	MeCN	-2.72^{κ}		-0.52	283

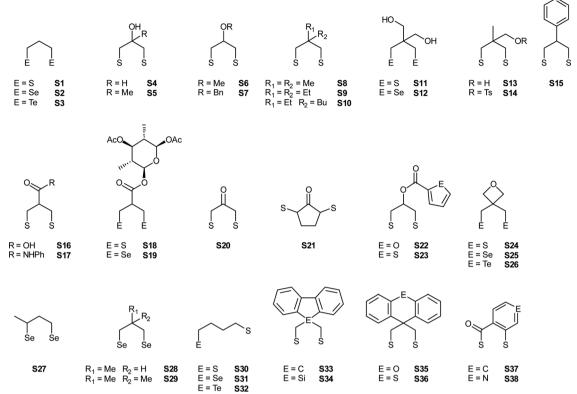
^a Second oxidation corresponds to- $Fe^{I}Fe^{II}/Fe^{II}Fe^{II}$. ^b Reported vs. NHE, converted. ^c Electron count not defined. ^d Reference for redox potentials in different $H_2O:$ MeCN ratios. ^e dppf = 1,10-bis(diphenylphosphino)ferrocene. ^f Reported vs. 0.01 M Ag/Ag(NO₃), converted. ^g Reported vs. Ag/AgCl. ^h $Fe^{I}Fe^{I}/Fe^{I}Fe^{I}$ 1-Methyl-3-(2-pyridyl)imidazol-2-ylidene. ^f $Fe^{I}Fe^{II}Fe^{II}$ Reported vs. Ag/AgCl, converted. ^l Under CO. ^m Reported vs. SCE, converted.

Table 13 Alternative dithiolate bridges



Dithiolate	Ligands: Fe ₁ //Fe ₂	Solvent	$E_{\rm pc}({\rm Fe^IFe^IFe^IFe^0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^I/Fe^IFe^{II}})^a/{\rm V}$	Ref.
S16		MeCN	-1.64^{b}	-2.40		141 and 144
S16		$MeCN^c$	-1.64	-2.51		
S16	PMe ₃ //PMe ₃	MeCN	-2.42			141
S17		$MeCN^c$	-1.67	-2.60		141
S18		MeCN	-1.58^{d}	-1.97	+1.01	133
S19		MeCN	-1.54^{d}	-2.10	+0.89	133
S4		MeCN	-1.61^{a}	-2.23	+0.77	134
S5		MeCN	-1.60^{a}	-2.26	+0.70	135
S6		MeCN	-1.60^{d}	-2.17	+0.79	135
S15		MeCN	-1.61	-2.17	+0.75	479
S15	PPh_3	MeCN	-1.78		+0.39	479
S15	κ¹-dppm	MeCN	-1.80	-2.16	+0.19	479
S15	μ-dppm	MeCN	-2.22			479
S 7		MeCN	-1.61	-2.12	+0.70	339
S 7	PPh_3	MeCN	-1.63	-2.41	+0.40	339
S 7	PPh ₃ //PPh ₃	MeCN	-1.96	-2.26	+0.07	339
S21	PMe ₃ //PMe ₃	DCM			+0.04, +0.29	474
S11		MeCN	-1.53		+0 . 67 ^e	136
S12		MeCN	-1.49		$+0.58^{e}$	136
S13	μ-(PPh ₂) ₂ NCH ₂ -pyridin-2-yl	THF^c	-2.25	-2.34		488
	,		-2.27	-2.38		
S13	μ-(PPh ₂) ₂ NBn	THF^c	-2.27	-2.36		488
	. (=/-	THF^c	-2.21	-2.31		
S14	μ-(PPh ₂) ₂ NCH ₂ -pyridin-2-yl		-2.19	-2.41^{f}		488
	. (=,= = =10		-2.12	-2.29^{f}		
S22		$MeCN^c$	-1.58	-2.21	+0.85	137
S23		$MeCN^c$	-1.59	-2.23	+0.84	137
S20		$MeCN^c$	-1.52	-2.39	+1.01	137
S9		$MeCN^c$	-1.67	-2.27	+0.82	473
S10		$MeCN^c$	-1.64	-2.27	+0.78	473
S8		$MeCN^c$	-1.61	-2.24	+0.73	473
S8	IMes	$MeCN^c$	-2.01		+0.76, +0.05	473
S8	PPh_3	$MeCN^c$	-1.79	-2.29	+0.69, +0.35	473
S9	IMes	$MeCN^c$	-2.02		+0.73, +0.16	473

Table 13 (continued)



			332			
Dithiolate	Ligands: Fe ₁ //Fe ₂	Solvent	$E_{\rm pc}({\rm Fe^IFe^IFe^IFe^0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^I/Fe^IFe^{II}})^a/{\rm V}$	Ref.
S24		MeCN	-1.60		+0.81	146
S25		MeCN	-1.55		+0.78	146
S26		MeCN	-1.54		+0.71	146
S2		MeCN	-1.61	-2.15	$+0.73^{e}$	477
S2		MeCN	-1.90^{bg}		$+0.95^{e}$	145
S27		MeCN	-2.08^{b}		$+0.32^{g}$	145
S28		MeCN	-1.61			148
S29		MeCN	-1.55^{h}			148
S30		DCM	-1.66^{i}		+0.76	128 and 147
S31		DCM	-1.63^{h}		+0.72	128
S32		DCM	-1.64^{h}		+0.71	128
S3		MeCN	-1.58	-2.09	$+0.64^{e}$	149
S3	μ -(PPh ₂) ₂ N- n Pr	MeCN	-2.06	-2.45	+0.14	149
S33	/-	DCM	-1.48^{h}			158
S35		DCM	-1.56^{h}			158
S36		DCM	-1.56^{h}			158
S34	PPh_3	DCM	-1.81			158
S38	-		-1.18			489
S37			-1.28			489
S37	PPh ₃		-1.47			489
S37	PPh_3		-1.47			

 $[^]a$ Second oxidation corresponds to- Fe^IFe^{II}/Fe^{II}Fe^{II}. b Reported νs . Ag/AgCl, converted. c Under CO. d Reported νs . 0.01 M Ag/Ag(NO₃), converted. e Fe^IFe^I/Fe^{II}Fe^{II}Fe^{II} Fe^{II}F

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6.5.2 ADT and derivatives thereof

Table 14 ADT and derivatives thereof

R	Ligands: Fe ₁ //Fe ₂	Е	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm II})^a/{\rm V}$	Ref.
S39		S	MeCN	-1.58		+0.59	170
S39		S	MeCN	-1.20^{b}			32
S39		Se		-1.20			32
S39	PPh_3	S	MeCN	-1.70		+0.19	170
S39	PMe_3	S	MeCN	-1.88		+0.51	170
S39	$P(p-tol)_3^c$	S	MeCN	-1.81		+0.26	321
S39	$P(m-tol)_3$	S	MeCN	-1.83		+0.33	321
S39	$P(p-C_6H_4F)_3$	S	MeCN	-1.77		+0.34	321
S39	$P(m-C_6H_4F)_3$	S	MeCN	-1.71		+0.39	321
S39	$P(C_4H_3O)_3$	S	MeCN	-1.80		+0.45	321
S39	PPh ₂ (OEt)	S	MeCN	-1.84		+0.40	321
S39	CN-	S	MeCN	-2.23		-0.043	478
S39	μ,κ^1,κ^2 -triphos ^d	S	DCM			-0.45	354
S60		S	MeCN	-1.42^{e}	-1.79	+0.70	194
S61		S	MeCN	-1.56^{e}		+0.72	194
S49		S	MeCN	-1.57^{e}		+0.56	177
S47		S	MeCN	-1.66		+0.59	186
S57		S	MeCN	-1.65		+0.61	186
S74		S	MeCN	-1.49	-1.96	+0.86	192
S73		S	MeCN	-1.51	-2.00	+0.87	192
S71		S	MeCN	-1.54	-1.99	+0.81	192
S72		S	MeCN	-1.52	-1.97	+0.82	192
S62		S	MeCN	-1.54	-2.01	+0.55, +0.87	460
S62	PPh_3	S	MeCN	-1.67		+0.34, +0.61	343
S50		S	MeCN ^f	-1.63	-2.33	+0.59	190
S51		S	$MeCN_{f}^{f}$	-1.62	-2.29	+0.56	190
S55		S	$MeCN^f$	-1.63	-2.10	+0.53	190

Table 14 (continued)

R	Ligands: Fe ₁ //Fe ₂	E	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^{II}})^a/{\rm V}$	Ref.
S56		S	$MeCN^f$	-1.64	-2.09	+0.56	190
S76		S	MeCN	-1.61		+0.62	191
S76	PMe ₃ //PMe ₃	S	MeCN	-2.01		-0.23	191
S53		S	MeCN	-1.60		+0.62	191
S53	PMe ₃ //PMe ₃	S	MeCN	-2.05		-0.27	191
S54		S	MeCN	-1.63		+0.64	191
S54	PMe ₃ //PMe ₃	S	MeCN	-1.94		-0.12	191
S63		S	MeCN	-1.61	-2.10	+0.48, +0.81	195
S63	$PHPh_2$	S	MeCN	-1.78	-2.22	+0.26, +0.49	195
S66		S	MeCN	-1.55			193
S43		S	MeCN	-1.66	-2.22	+0.55, +0.88	182
S43	PMe ₃ //PMe ₃	S	MeCN	-2.00	-2.32	-0.21, -0.09	182
S44		S	MeCN	-1.66	-2.24	+0.59, +0.92	182
S44	PMe ₃ //PMe ₃	S	MeCN	-1.99	-2.29	-0.20, -0.08	182
S45		S	MeCN	-1.68	-2.21	+0.58, +0.90	182
S45	$PMe_3//PMe_3$	S	MeCN	-2.00	-2.35	-0.22, -0.078	182
S82		S	MeCN	-1.55		+0.71, +0.73	196 and 457
S82	IMe//IMe	S	MeCN	-2.53			197
S83		S	MeCN	-1.64		+0.65	196
S84		S	MeCN	-1.54		+0.72	196
S85							
S85	IMe//IMe	S	MeCN	-2.49			197
S60		Se	MeCN	-1.48^{e}	-1.79	+0.58	198
S58		Se	MeCN	-1.57^{e}	-2.10	+0.50, +0.78	198
S59		Se	MeCN	-1.58^{e}	-2.08	+0.49, +0.81	198
S69		Se	$MeCN_{f}^{f}$	-1.50	-1.97	+0.72 ^g	173
S70		Se	$MeCN^f$	-1.48	-1.95	$+0.73^{g}$	173
S 77	$PMe_3//PMe_3$	S	DCM			-0.24, -0.02	318
S67		S	MeCN	-1.53	-2.02	+0.89	481
S67	PPh_3	S	MeCN	-1.74	-2.22	+0.44	481
S64		S	MeCN	-1.54	-2.02	+0.61	481

Table 14 (continued)

R	Ligands: Fe ₁ //Fe ₂	E	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^{II}})^a/{\rm V}$	Ref.
S64	PPh ₃	S	MeCN	-1.73	-2.09	+0.41	481
S78		S	MeCN	-1.56^{e}	-2.05	+0.61	454
S78	PMe ₃ //PMe ₃	S	MeCN	-2.18^{e}		-0.13	454
S79		S	MeCN	-1.56^{e}	-2.07	+0.67	454
S80		S	MeCN	-1.56^{e}	-2.06	+0.67	454
S59		S	MeCN	-1.55		+0.55	358
S59	$CN-pC_6H_4I//CN-pC_6H_4I$	S	MeCN	-1.70		+0.13	358
S65		S	MeCN	-1.67		+0.44	490
S65	$P(p\text{-tol})_3$	S	MeCN	-1.79		+0.30	321
S65	$P(m-tol)_3$	S	MeCN	-1.75		+0.34	321
S65	$P(p-C_6H_4F)_3$	S	MeCN	-1.72		+0.38	321
S65	$P(m-C_6H_4F)_3$	S	MeCN	-1.67		+0.39	321
S65	$P(C_4H_3O)_3$	S	MeCN	-1.72		+0.33	321
S46	1	_					
S46	PPh ₃	S	DCM	-2.04		+0.26	185
S46	$PPh_2(o-py)^h$	S	DCM	-2.08		+0.18	185
S46 S52	$P(p\text{-tol})_3$	S	DCM	-2.10		+0.22	185
S52	PPh_3	S	DCM	-2.05		+0.25	185
S52	$PPh_2(o-py)$	S	DCM	-2.06		+0.17	185
S52 S41	$P(p\text{-tol})_3$	S	DCM	-2.10		+0.23	185
S41 S42	μ-dppm	S	MeCN	-2.25^{i}		+0.08, +0.42	183
S42	κ^2 -dppe	S	MeCN THF	$-2.01 \\ -2.22$			338
S42	u dnno	S	MeCN	-2.22 -2.12			338
S42 S42	μ-dppe κ²-1,10-phenantroline	S	MeCN				
S42 S48	•			-2.22			463
S48	κ²-dppe	S	MeCN	-1.98			338
			THF	-2.16			

Table 14 (continued)

R	Ligands: Fe ₁ //Fe ₂	E	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\mathrm{pa}}(\mathrm{Fe}^{\mathrm{I}}\mathrm{Fe}^{\mathrm{I}}/\mathrm{Fe}^{\mathrm{I}}\mathrm{Fe}^{\mathrm{II}})^{a}/\mathrm{V}$	Ref.
S48	μ-dppe	S	MeCN	-2.10			338
			THF	-2.36			
S81		S	MeCN	-1.56			456
S91		S	MeCN	-1.56		+0.55	490
S91	PPh_3	S	MeCN	-1.67		+0.52	490
S87		S	MeCN	-1.56	-1.98	+0.61	490
S92		S	MeCN	-1.58	-2.06	+0.46	490
S88		S	DMF	-1.59	-2.49	+0.64	490
S89		S	MeCN	-1.56	-2.08	+0.60	490
S94		S	MeCN	-1.56	-2.08	+0.57	490
S86		S	MeCN	-1.54	-1.99	+0.55	490
S90		S	MeCN	-1.56	-2.03	+0.55	490
S93		S	MeCN	-1.55	-2.02	+0.50	490
S58	κ^2 -(PPh ₂) ₂ N(CH ₂ CHMe ₂)	S	MeCN	-2.10			491

 $[^]a$ Second oxidation corresponds to- Fe^IFe^{II}/Fe^{II}Fe^{II}. b Reported vs. SHE. c tol = methylphenyl. d Phenyl bis(diphenylphosphinoethyl)phosphine. e Reported vs. 0.01 M Ag/Ag(NO₃), converted. f Under CO. g Fe^IFe^{II}/Fe^{II} by = pyridyl. i Reported vs. 0.001 M AgNO₃, converted.

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6.5.3 EDT and derivatives thereof

Table 15 EDT and derivatives thereof

Dithiolate	Ligands: Fe ₁ //Fe ₂	Solvent	$E_{\rm pc}({\rm Fe^IFe^IFe^IFe^0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^{II}})/{\rm V}$	Ref.
S95		MeCN	-1.63^{a}			117
S95	PMe ₃ , NO ⁺	DCM	-0.45^{b}	-0.95		381
S95	PMe_3 , $NO^+//PMe_3$	DCM	-0.67^{c}	-0.98		381
S95	κ^2 -(PPh ₂) ₂ N(CH ₂ CHMe ₂)	MeCN	-2.23			491
S98		MeCN	-1.66			117
S96		MeCN	-1.68			117
S99		MeCN	-1.63			117
S99			-1.67^{d}		+0.91	138
S99	CN ⁻ //CN ⁻	MeCN	-2.75^{d}		-0.47	138
S100		MeCN	-1.64^{e}	-2.02		119
S100		H_2O	-1.07^{f}			
S101		MeCN	-1.60	-2.11	+0.96	121
S101	PPh_3	MeCN	-1.74		+0.57	121
S101	dppm	MeCN	-1.75		+0.46	121
S97		DCM	-1.11	-1.25		492

^a At fast scan rate (0.1 V s⁻¹). ^b Reported vs. Ag/AgCl at 0 °C. ^c Reported vs. Ag/AgCl. ^d Reported vs. SCE, converted. ^e Reported vs. NHE, converted. f Reported vs. NHE.

6.5.4 ODT and derivatives thereof

Table 16 ODT and derivatives thereof



X	E	Ligand L	Solvent	$E_{\rm pc}({\rm Fe^IFe^I/Fe^IFe^0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\mathrm{pa}}(\mathrm{Fe^{I}Fe^{I}/Fe^{I}Fe^{II}})/\mathrm{V}$	Ref.
S	S	CO	MeCN	-1.51	-1.94	+0.79	211
S	S	CN-tBu	MeCN	-1.76	-1.98	+0.49	213
S	Se	CO	MeCN	-1.49	-1.97	$+0.64^{a}$	205
O	S	CO	MeCN	-1.59	-2.10	$+0.81^{a}$	204
O	Se	CO	MeCN	-1.55	-2.06	+0.71 ^a	205

^a Fe^IFe^I/Fe^{II}Fe^{II}.

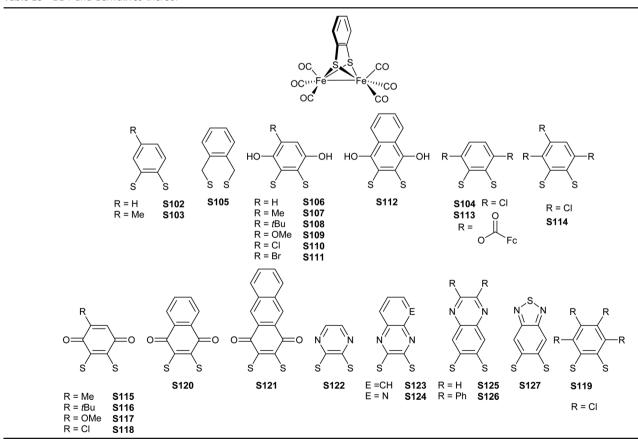
6.5.5 XDT and derivatives thereof

Table 17 XDT and derivatives thereof

X	G//G	Е	Ligands: Fe ₁ //Fe ₂	Solvent	$E_{\rm pc}({\rm Fe^IFe^IFe^IFe^0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^{II}})/{\rm V}$	Ref.
SiMe ₂	CH ₂ //CH ₂	S		DCM	-1.71	-1.84	+0.77	160
$GeMe_2$	$CH_2//CH_2$	S		DCM	-1.72	-1.85	+0.74	160
$SnMe_2$	$CH_2//CH_2$	S		DCM	-1.68	-2.20	+0.70	160
$SnMe_2$	CH ₂ //—	Se		DCM	-1.75^{a}			161
$SnMe_2$	$CH_2//CH_2$	Se		DCM	-1.63^{a}			161
$SiPh_2$	CHPh//CHPh	S		DCM	-1.57			159
				MeCN	-1.43			
$SiPh_2$	CHPh//CHPh	S	κ²-dmpe	DCM	-2.21			159
			-	MeCN	-2.00	-2.30		
$SiPh_2$	CHPh//CHPh	S	μ-dmpe	DCM	-2.32			159
				MeCN	-2.14			
P(O)Me	$CH_2//CH_2$	S		MeCN	-1.41			202
P(O)(OEt)	$\mathrm{CH_2/\!/CH_2}$	S		MeCN	-1.42			202
^a Fe ^I Fe ^I /Fe ⁰	Fe ⁰ .							

6.5.6 BDT and derivatives thereof

Table 18 BDT and derivatives thereof



Bridge	Ligands: Fe ₁ //Fe ₂	Solvent	$E_{\rm pc}({\rm Fe^IFe^I/Fe^IFe^0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^{II}})/{\rm V}$	Ref.
S102		MeCN ^a MeCN MeCN	$-1.35, ^{b} -1.36$ -1.36^{b} -1.31			222 and 232 232 226

Table 18 (continued)

Bridge	Ligands: Fe ₁ //Fe ₂	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^{II}})/{\rm V}$	Ref.
		MeCN	-1.32^{b}			468
			-1.27^{b}			467
S102	PMe ₃ //PMe ₃	MeCN	-2.09			222
S102	$PR_3^{\ c}$	DCM	-1.44^{d}	-1.64		493
S102	$P(OMe)_3$		-1.59		+0.64	482
S102	PPh_3	DCM	-1.82		+0.58	480
		MeCN	-1.57	-1.80		
S102	PPh_2Me	DCM	-1.89		+0.60	480
		MeCN	-1.59	-1.88		
S102	PPh_2H	DCM	-1.80		+0.60	480
		MeCN	-1.54	-1.76		
S102	PPh ₃ //PPh ₃	DCM	-2.17		+0.13	480
		MeCN	-1.83			
S102	PPh ₂ Me//PPh ₂ Me	DCM	-2.21			480
		MeCN	-1.89			
S102	$PPh_2H//PPh_2H$	DCM	-2.02		+0.21	480
		MeCN	-1.74			
S102	Val-Ppa-Leu	MeCN	-1.62		+0.47	306
		$MeCN/H_2O 3:1$	-1.51		+0.48	
		$MeCN/H_2O 3:2$	-1.47		+0.46	
S106		MeCN	-1.28			228^e
S109		MeCN	-1.32			228
S107		MeCN	-1.28			228
S108		MeCN	-1.27			228
S110		MeCN	-1.22			228
S111		MeCN	-1.22			228
S112		DCM	-1.34			228
S106		$MeCN^a$	-1.34^{b}			232
		MeCN	-1.34^{b}			
S113		$MeCN^a$	-1.28^{b}			232
			-1.28^{b}			
S103		MeCN	-1.37			222
			-1.34^{f}			226
S103	$PMe_3//PMe_3$	MeCN	-2.08			222
S104		MeCN	-1.23			222
			-1.20			226

Table 18 (continued)

Bridge	Ligands: Fe ₁ //Fe ₂	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	E _{pa} (Fe ^I Fe ^I /Fe ^I Fe ^{II})/V	Ref.
S104	PMe ₃ //PMe ₃		-1.91			222
S115		DCM	-1.23^{g}			230
S116		DCM	-1.20^{g}			230
S117		DCM	-1.32^{g}			230
S118		DCM	-1.24^{g}			230
S120		DCM	-1.29^{g}			230
S120	$PMe_3//PMe_3$	DCM	-1.71^{g}			230
S121		DCM	-1.25^{g}			230
S122		MeCN	-1.17^{b}			229
S122	PPh_3	MeCN	-1.38	-1.65		229
S124		MeCN	-1.23^{b}			229
S124	PPh_3	MeCN	-1.42	-1.70		229
S123		MeCN	-1.22			222
S123	PPh_3	MeCN	-1.41	-1.70		229
S123	PMe ₃ //PMe ₃		-1.88			222
S125		MeCN	-1.34^{b}			234
S125	PPh_3	MeCN	-1.55	-1.85		494
S125	$P(OEt)_3$	MeCN	-1.58			494
S125	$P(OEt)_3//P(OEt)_3$	MeCN	-1.9			494
S126		MeCN	-1.34^{b}			234
S126	PPh_3	MeCN	-1.56	-1.80		494
S126	$P(OEt)_3$	MeCN	-1.57			494
S126	$P(OEt)_3//P(OEt)_3$	MeCN	-1.84			494
S127		MeCN	-1.35^{b}			234
S127	PPh_3	MeCN	-1.56	-1.83		494
S127	$P(OEt)_3$	MeCN	-1.60			494
S127	$P(OEt)_3//P(OEt)_3$	MeCN	-1.86			494
S105		DCM	-1.68^{b}_{c}			221
S114		MeCN	-1.15^{I}			226
S119		MeCN	-1.13^{f}			226

^a Under CO. ^b Fe^IFe^I/Fe⁰Fe⁰. ^c 2,2'-(2-Phenyl-4,5,6,7-tetrahydro-2*H*-isophosphindole-1,3-diyl)dipyridine. ^d Reduction associated with the ligand at -1.76 V. ^e For this reference, reported are E_{ov}° values, *i.e.* standard potential for overall 2e⁻ reduction defined as $E_1^{\circ} + E_2^{\circ}/2$. ^f Electron count not defined. ^g Reduction associated with quinone.

6.5.7 1,8-Disulfidonaphtalene complexes and derivatives thereof

Table 19 1,8-Disulfidonaphtalene complexes and derivatives thereof

$$R = H = 13128 \quad R = CH_0 CH \\ R = H = 13128 \quad R = CH_0 CH \\ R = (R + H + 13128) \quad R =$$

			K - CI 3152		
E	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm II})^a/{\rm V}$	Ref.
S	DCM	-1.76	-2.00		239
	MeCN	-1.52	-1.96	+0.87	
S	DCM	-1.60		+1.00	239
S	DCM	-1.84	-2.12		239
	MeCN	-1.59	-2.05	+0.78	
S	DCM	-1.51		+0.86	242
S	MeCN	-1.46	-1.60	+0.75	242
S	DMF	-1.36	-1.63		242
S	DCM	-1.54		+0.69	242
S	MeCN	-1.45	-1.55	+0.77	242
S	DMF	-1.37	-1.58		242
S	DCM	-1.62	-1.88	+0.82	242
S	MeCN	-1.53	-1.88	+0.79	242
S	DCM	-1.65	-1.87	+0.82	242
S	MeCN	-1.54	-1.87	+0.81	242
S	DCM	-1.60	-1.93	+0.89	242
S	MeCN	-1.51	-1.87	+0.85	242
S	DCM	-1.63	-1.95	+0.90	242
S	MeCN	-1.51	-1.89	+0.84	242
S	DCM	-1.61	-1.93	+0.90	242
S	MeCN	-1.51	-1.88	+0.82	242
S	DCM	-1.63	-1.96	+0.89	242
S	MeCN	-1.51	-1.90	+0.82	242
	S S S S S S S S S S S S S S S S S S S	S DCM	S DCM -1.76 MeCN -1.52 S DCM -1.60 S DCM -1.84 MeCN -1.59 S DCM -1.51 S MeCN -1.46 S DMF -1.36 S DCM -1.54 S MeCN -1.45 S MeCN -1.45 S MeCN -1.62 S MeCN -1.62 S MeCN -1.63 S DCM -1.65 S DCM -1.65 S MeCN -1.54 S DCM -1.65 S MeCN -1.54 S DCM -1.65 S MeCN -1.54 S DCM -1.65 S MeCN -1.51 S DCM -1.63 S MeCN -1.51 S DCM -1.63 S MeCN -1.51 S DCM -1.61 S MeCN -1.51 S DCM -1.61 S MeCN -1.51 S DCM -1.63	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$

Table 19 (continued)

$$R = H \quad S128 \quad R = CHO \quad S131 \quad R = CHO \quad S132 \quad R = CH_0 CH \quad S132 \quad R = CH_0 CH \quad S133 \quad$$

Bridge	E	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe^IFe^0/Fe^0Fe^0})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^IF})^a/{\rm V}$	Ref.
S139	S	DCM	-1.62	-1.92	+0.85	242 and 243
S144	S	DCM	-1.34	-1.68	+0.93	244
S144	Se	DCM	-1.39	-1.63	+0.89	244
S145	S	DCM	-1.12	-1.54	+1.11	244
S145	Se	DCM	-1.17	-1.55	+0.98	244
S129	S	MeCN	-1.65		+0.99, +1.20	240
S129	Se	MeCN	-1.64		+0.88, +1.13	240
S128	Se	MeCN	-1.54		+1.00	240
S128	S, Se	MeCN	-1.60		+0.68, +1.12	240
S130	Se	MeCN	-1.34		+0.55, +0.90	240
S130	S, Se	MeCN	-1.61		+1.07	240
S143	S	MeCN	-1.64		+0.87, +1.14	240
S143	Se	MeCN	-1.52		+0.98, +1.31	240
S140	S	MeCN	-1.51	-1.74	+1.00	251
S140	Se	MeCN	-1.45	-1.86	+0.96	251
S141	S	MeCN	-1.51	-1.85	+0.46, +0.76	251
S141	Se	MeCN	-1.51	-1.90	+0.52, +0.68	251
S148	S	DCM	-0.99	-1.23	+0.98	247
S147	S	DCM	-1.01	-1.24	+0.97	247
S149	S	DCM	-1.04	-1.25	+0.99	247
S150	S	DCM	-1.58			241
S146	S	MeCN	-1.1318	-1.49		248
S153	S	MeCN	-1.279		+0.42	250
S151	S	MeCN	-1.09	-1.30		475
S152	S	DCM	-1.05	-1.40		492

^a Second oxidation corresponds to Fe^IFe^{II}/Fe^{II}Fe^I.

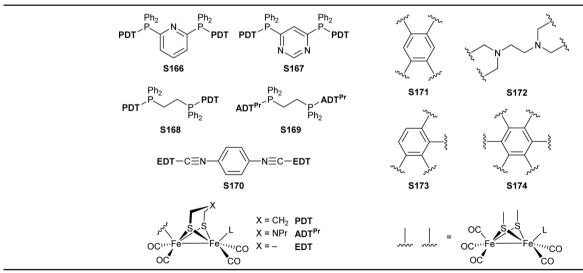
6.5.8 Other models

Table 20 Other models with monodentate thiolates

Thiolate	Ligands: Fe ₁ //Fe ₂	Solvent	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm I}/{\rm Fe}^{\rm I}{\rm Fe}^{\rm 0})/{\rm V}$	$E_{\rm pc}({\rm Fe}^{\rm I}{\rm Fe}^{\rm o}/{\rm Fe}^{\rm o}{\rm Fe}^{\rm o})/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^I})^a/{\rm V}$	Ref.
S161		MeCN	-1.29	-1.73	+0.74	264
S162		MeCN	-1.22	-1.68	+0.88	264
S163		MeCN	-1.24	-1.71	+0.75	264
S164		MeCN	-1.19	-1.66	+0.77	264
S157		MeCN	-1.44	-2.26	+0.81	265
S158		MeCN	-1.51	-2.42	+0.93, +1.06	265
S159		MeCN	-1.35	-2.11	+0.79	265
S160		MeCN	-1.55	-2.29	+0.73, +1.08	265
S160	$PMe_3//PMe_3$	MeCN	-2.08	-2.45	-0.23, +1.03	265
S165		MeCN	-1.33		+0.61	495
S165	PMe_3	MeCN	-1.49		+0.31	495
S165	$P(p-C_6H_4OMe)_3$	MeCN	-1.66	-2.37	+0.29, +1.00	496
S156	2 2 1 70	$MeCN^b$	-1.73	-2.37	•	141
S154		DMF	-1.25^{c}	-1.64		497
S155		DMF	-1.20^{cd}	-1.60		497

^a Second oxidation corresponds to- Fe^IFe^{II}/Fe^{II}Fe^{II}. ^b Under CO. ^c Reported vs. SCE. ^d Average for syn and anti-isomer.

Table 21 Other models with two or more [2Fe-2S] units



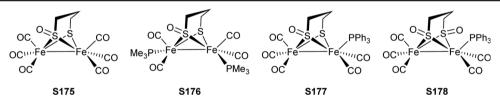
Unit	Ligands L	Solvent	$E_{\rm pc}({\rm Fe^IFe^IFe^IFe^0})^a/{\rm V}$	$E_{\mathrm{pa}}(\mathrm{Fe^{I}Fe^{I}/Fe^{I}Fe^{II}})^{e}/\mathrm{V}$	Ref.
S166	CO, CO	DCM	-2.06^{b}		305
S167	CO, CO	DCM	$-2.04, -2.25^{b}$		305
S168	CO, CO	DCM	-1.94^{c}	+0.67	183
S172	CO, CO	MeCN	-1.59^{b}	+0.58	177
S169	CO, CO	DCM	-1.93^{c}	+0.62	183

Review Article Table 21 (continued)

Unit	Ligands L	Solvent	$E_{\rm pc}({\rm Fe^IFe^IFe^IFe^0})^a/{\rm V}$	$E_{\rm pa}({\rm Fe^IFe^IFe^IFe^{II}})^e/{\rm V}$	Ref.
S170	CO, CO		-1.65, -1.82	+0.50, +0.85	204
S171	CO, CO	DCM	$-1.38,^d$ -1.66^d		235
S171	CO, PPy ₃	DCM	$-1.42,^d$ -1.70^d		235
S171	PPy_3, PPy_3	DCM	$-1.47, ^d -1.79^d$		235
S173	CO, CO	DCM	$-1.40,^d$ -1.66^d		235
S174	CO, CO, CO	DCM	-1.33 , d , -1.56 , d , -1.81		236

 $[^]a$ More than one Fe^IFe^I/Fe^IFe⁰ steps possible. b Involving ligand reduction. c Reported vs.~0.001 M Ag/AgNO₃. d All values involving 2e⁻. e Electron count not defined.

Table 22 Oxidised models



Complex	Solvent	$E_{\rm pc}({\rm Fe^IFe^I/Fe^IFe^0})/{\rm V}$	$E_{ m pc}{}^a/{ m V}$	$E_{\rm pa}({\rm Fe^IFe^I/Fe^IFe^{II}})/{\rm V}$	Ref.
S175	MeCN	-1.58	-1.70, -2.25	+0.95	215
S176	MeCN	-2.21	-2.38	+0.03	215
S177	MeCN	-1.77	-1.97, -2.3	+0.48	215
S178	MeCN	-1.72	-1.98	+0.65	215

^a Further undefined reductions.

 Table 23
 Photocatalytic efficiencies of the diiron subsite models

Catalytic site	Sensitizer	Electron/proton source/solvent	TON	TOF	Ref.
105	ZnTPP	2-Mercaptobenzoic acid, CF ₃ COOH in CH ₂ Cl ₂	0.16		528
91	Re(imidazo[4,5-f]-1,10-phenanthroline) (4-(phenylethynyl)pyridine) (CO) ₃ -Fc	Ascorbic acid CH ₂ CN	0.35		527
170	ZnPn ₃ PhP	<i>p</i> -Anisidine, CF₃COOH toluene	0.5		248
170	Modified ZnTPP	2-Mercaptobenzoic acid, CF ₃ COOH CH ₂ Cl ₂	0.56		252
150	$[Ru(bpy)_3]^{2+}$	Ascorbic acid DMF/H ₂ O	200	$2.7 h^{-1}$	537
91	$[Ru(bpy)_3]^{2+}$	Ascorbic acid CH ₃ CN/H ₂ O	4.3		536

Chem Soc Rev

Electrocatalytic proton reduction by H-cluster models

Besides the basic redox properties, the function of numerous diiron various complexes in the presence of a proton source was analysed. Although water is considered to be an ideal proton source and electrolyte, aqueous conditions are mostly not attainable for the catalysis tests due to the insolubility and instability of the mimics (Section 4.6). As a result, artificial Fe₂S₂ systems are commonly studied in various organic solvents (MeCN, DCM, THF, DMF) in presence of acids of varying strengths, e.g. organic acids (CH3COOH, CF3COOH, HOTs, Pivalic acid) or HBF4·OEt2. While native [FeFe]-hydrogenases catalyse the proton reduction starting from the Fe^{II}Fe^I H_{ox} state (Section 3.1), the active site mimics mostly possesses an inactive Fe^IFe^I resting state and reveal HER activity only upon their reduction (Section 4.7). Studying the proton reduction mechanisms of the subsite models is thus important to develop an in-depth understanding of the underlying principles and to design an "ideal", optimized artificial catalytic system.

Although the specific mechanism varies between different complexes and depends upon various factors, the general pathways for the proton reduction by subsite analogues are present and highlighted in Fig. 97. Their combination depends on the applied complex as well as acid utilized. Also, it is to be noted that the L_nFe₂^{x,x}H state does not specifically represent protonation at the metal centre, rather it represents unspecific protonation of the subsite model.

As discussed in the previous sections, hexacarbonyl mimics in their Fe^IFe^I state rarely undergo direct protonation. However, if the carbonyl ligands are substituted by electron donating ligands (phosphines (PMe3, P(OEt)3, dppe, dppm, dppv, carbenes (IMe, IMes), cyanide - Section 5.3.2), the Fe centre(s) becomes basic enough to undergo a direct protonation 150,151,154,454 resulting in $[L_nFe_2^{II,I}H]$ or $[L_nFe_2^{II,II}HH]$. Elsewise, models

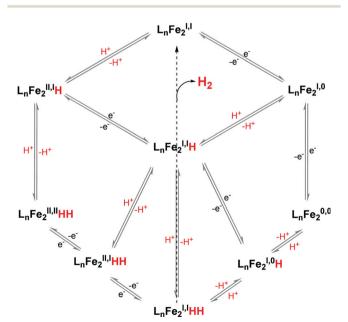


Fig. 97 Commonly observed electron and proton transfer pathways facilitated by subsite models.

undergo one- or two-electron reduction steps affording the $L_n Fe_2^{I,0}$ or $L_n Fe_2^{0,0}$ state. These states show greater affinity for protons due to their increased electron density/basicity at the metal centres and hence are readily converted to the hydride intermediates $L_n Fe_2^{I,I}H$ or $L_n Fe_2^{I,0}H$ even in the presence of weaker acids such as HOAc. ^{135,151,186,239,242,466} Further protonation of the hydride intermediates then results in the formation of dihydride intermediates $L_n Fe_2^{II,I}HH$ or $L_n Fe_2^{I,I}HH$ and hydrogen might then be released from a two-electron reduced double-protonated intermediate involving either a dihydrogen or dihydride species. Cleavage of H₂ then reforms the starting complex $L_n Fe_2^{I,I}$.

Notably, the presence of an additional basic site (e.g. adjacent amines) within the subsite analogues was shown to affect the overall catalytic mechanism. In such cases, ligand protonation can be kinetically favoured and then dominantly occur. Usually, such a protonation is then followed by an intramolecular proton transfer to the metal centre and eases the catalytic progress through facilitated formation of the hydride state.

A detailed analysis of the proton reduction mechanism can be only derived from a combination of electrochemical, spectroscopic (IR, UV-vis), and spectroelectrochemical data. Notably, the proposed mechanisms were often supported or even forecasted by theoretical calculations that are therefore an anchor stone in these detailed analyses.

6.6.1 Mechanisms of proton reduction by unsubstituted H-cluster models. The electrocatalytic ability of 20 was studied in detail in the presence of HOTs utilizing electrochemical as well as spectroelectrochemical methods. 380,498 DFT calculations assisted in describing the plausible proton reduction. 499 As 20 in the Fe^IFe^I state is itself not protonable even by strong acids, the catalytic cycle herein was shown to begin with a 1e⁻-reduction step resulting in a monoanionic [Fe^IFe⁰] state. This intermediate is sufficiently basic to be protonated by HOTs and enables the generation of a bridging hydride intermediate (Fe^I–μH–Fe⁰) (Fig. 98). Subsequently, this bridging hydride species undergoes a second reduction at potentials milder than $-1.59 \text{ V } (-1.2 \text{ V } \text{vs. SCE}) \text{ yielding } [\text{Fe}^0 - \mu \text{H} - \text{Fe}^0]^-$. This double-reduced state can then undergo further protonation either at the metal centre resulting in breaking of the metalmetal bond and leads to formation of a dihydridic Fe^{II}Fe^{II} state. Likewise, the formation of an intermediate possessing a bridging hydride as well as carbonyl with concomitant Fe-S bond cleavage and protonated sulphur was suggested. 380,498,499 This thiol carrying intermediate, on further reduction at E = -1.84 V (-1.45 V vs. SCE) generates an anionic species, which releases H2 and transforms back to the singly reduced [FeIFe0] state $(K = 1.7 \times 10^8)$.

On the other hand, H2 release from the dihydride intermediate of Fe₂(pdt)(CO)₆ is slow ($K = 10^4 \text{ vs. } 1.7 \times 10^8$). 380,498,499 Notably, when Fe₂(pdt)(CO)₆ was assessed in the presence of weaker acids such as AcOH, the catalysis progressed via a $Fe^{0}Fe^{0}$ state formed at more negative potentials -2.35 V.^{466}

Contrary, as the azadithiolate models possesses a basic amine, the catalytic cycle can start with an initial protonation step of the bridgehead followed by a single-electron reduction.

Fig. 98 Catalytic pathway for proton reduction by Fe₂(pdt)(CO)₆.

However, the protonation behaviour of azadithiolate modified models strongly depends on the basicity of the bridgehead amine (and with it the substitution thereon) and strength of the employed acid. The interplay between these factors greatly affects the mechanism of the proton reduction. For instance, 2 is protonated with moderately strong acids such as HOTs (or Cl₃CCOOH) at the bridgehead nitrogen leading to intermediate [2-NH]+ which undergoes reduction at -1.27 V. Hereafter, the Fe site is rendered more basic than the amine $(\Delta p K_a 3.3)$. Thus, internal H⁺ transfer takes place leading to a metal protonated-terminal hydride state [2-NtH] (Fig. 99 scheme A). 500 Contrastingly, for N-protonated intermediates of Fe₂(adt^R)(CO)₆ (R = iPr, CH₂CH₂OCH₃), no such tautomerisation is observed. The high electron density at the bridgehead nitrogen by the electron donating substituents was suggested to restrict the H⁺ transfer. Furthermore, as depicted in Fig. 99 scheme B, the reduced [2-NRH] intermediate takes up a second proton and subsequently yields the bridging hydride state [2-NRHµH] which after additional 1e reduction regenerates the Fe^IFe^I state. Moreover, with stronger acids such as HBF₄· OEt₂ further protonation of the $[2-N^R(H)_2]$ gives an $[2-N^R(H)_3]^+$ state and regenerates the singly protonated reduced species [2-NRH] upon reductive H₂ elimination (Fig. 99 scheme B'). 458,501,502 In the case of 2, after the terminal hydride state is achieved, the catalytic cycle proceeds with a second protonation step resulting in double-protonated state [2-NHtH]+. Further reduction of this species gives [2-NHtH], which releases H₂ and achieves the Fe^IFe^I state (Fig. 99 scheme A).

Notably, when utilizing weaker acids (*e.g.* CH₃COOH), Fe₂(adt^R)(CO)₆ models adopt a pathway similar to the PDT derivatives (Fig. 99 scheme C). Here the model must take up electrons before being protonated and the adjacent amine is not involved in the overall hydrogen formation mechanism. *E.g.*, Fe₂(adt)(CO)₆ is protonated by ClCH₂COOH (p $K_a = 15.3$) at the metal centre (p $K_a = 17.1$) only upon undergoing 1e⁻ reduction at -1.60 V. There is also a possibility to establish a concerted proton electron transfer pathway resulting in [2-NttH] from [2-NtH]⁺ (Fig. 99 scheme C'). However, this pathway has not been explored in detail and thus will not be further discussed.⁵⁰⁰

Underlining the control of acidic strength on the catalytic pathway, $Fe_2(adt^R)(CO)_6$ (R = p-C₆H₄COOMe, 301)⁴⁶⁰ was shown to adopt two different pathways for the proton reduction depending on the strength of the utilized acid. In case of CF₃COOH, the complexes follow an ECCE mechanism as the reduced state can be protonated twice. Subsequent reduction generates H2. Contrarily, with HOAc an EECC mechanism is observed. Herein, only the dianionic complex of the models is basic enough to be protonated by weak acids and consequently H₂ is only released at very negative potentials. 491 The catalytic efficiencies with CF₃COOH (TON = 10.6) were shown to be larger than with HOAc (TON = 4).460 Similarly, in case of a missing suitable protonation site, it is justified that the double-reduced species is protonated generating H2 via an EECC mechanism with weak acids. Hereby it can be ascertained that the weaker acid (HOAc) requires the Fe⁰Fe⁰ state Chem Soc Rev Review Article

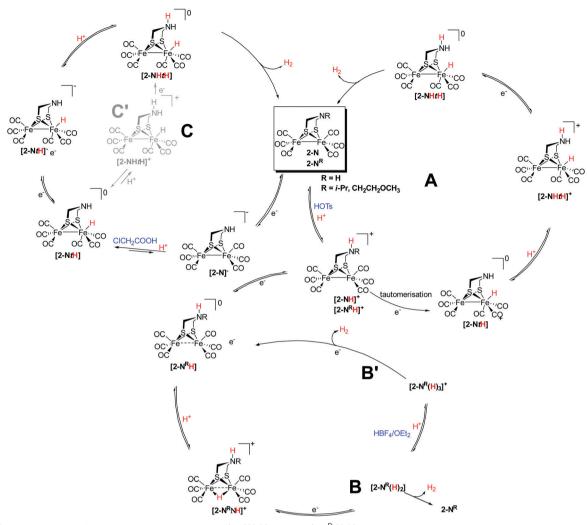


Fig. 99 Catalytic pathways for the proton reduction by $Fe_2(adt)(CO)_6$ and $Fe_2(adt^R)(CO)_6$ analogues with acids of varying strength.

for catalytic activity. 134,190,192,195,466,481 Contrastingly, the presence of a stronger acid such as HOTf, HClO $_4$ or HBF $_4$ ·OEt $_2$ leads to a CEEC mechanism. 177,194,196,456,481

As discussed in Section 4.3, BDT models possess a metal site that is sufficiently electron deficient and hence readily undergo a $2e^-$ -reduction yielding the Fe^0Fe^0 state. This state participates in the catalytic cycle, generating H_2 from HOAc upon double-protonation following an EECC mechanism. However, with HOTs the monoanionic mono-protonated state formed undergoes another reduction resulting in dianionic state, which only then releases H_2 and reforms the Fe^1Fe^0 state (Fig. 100). 467,468

A similar observation was made for phosphorus-substituted model **64**. In the absence any of acid, model **64** is reduced via a concerted two-electron process. Contrary, when HOTs is added, the double-reduced state undergoes two successive protonation steps to yield a dihydride. Herein, each iron was suggested to bind a hydride affording [**64-2tH**]. Upon release of H_2 , this state reverts back to Fe^IFe^I . Another plausible route involves a one-electron reduction of [**64-2tH**] resulting in the monoanionic state with concomitant H_2 evolution. The former pathway is

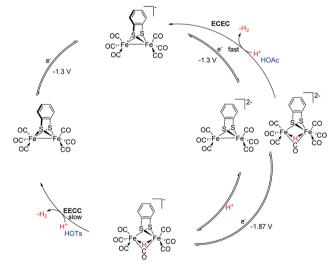


Fig. 100 Catalytic pathway for proton reduction by 149

considered to eliminate H_2 at slower rates due to the larger spatial separation of the two hydrides in [64-2tH].⁵⁰³

Along this line, **170** catalyses the proton reduction *via* two different pathways from HOTs.

- (1) Protonation of the reduced state leads to [170-H]. This protonation is followed by another reduction and protonation and results in [170-HH]. The thus obtained intermediate can now either directly release $\rm H_2$ by formation of 170 or undergo further reduction thereby releasing $\rm H_2$ from [170-HH]⁻ resulting in 170⁻.²³⁹
- (2) The naphthalene substituted imine model 302 herein follow a CECE mechanism to reduce protons from HOTs.²⁵¹ The acid protonates the imino substituent, resulting in imine protonated state [302–H]⁺, which is reduced at slightly positive potential than the parent model 302 resulting in [302-NH]⁺. [302-NH]⁺ undergoes a second protonation followed by reduction to give [302-HNH]⁺ which releases H₂.²⁵¹ Notably, the pyrazine modified models 303²²⁹ reduce protons from HOAc wherein the aromatic ring nitrogen acts as internal site of protonation. The protonated state undergoes reduction followed by another protonation at the nitrogen. Reduction of the diprotonated state yields the Fe⁰Fe⁰ state, which evolves dihydrogen and closes the catalytic cycle.

Another example wherein the acid strength influences the proton reduction pathway is the tetranuclear model **164**. When strong acids such as CX_3COOH (X = Cl, F) were employed as the proton source, the dianionic state was rapidly protonated and releases H_2 . Contrarily, in the presence of the weaker acid $CH_2ClCOOH$ the mono-protonated dianionic state either undergoes a further two-electron reduction resulting in the $[164]^{3-}$ state, which on further reaction with two protons releases H_2 from the $[164-HH]^{2-}$ (Fig. 101).

The influence of the ring-substituents on different catalytic pathways for the proton reduction can be demonstrated by **163** and **162** in the presence of HBF₄·OEt₂. In case of **163**, a direct protonation of the model is not observed and was explained by the interaction of the nitrogen lone-pair with the ring sulphur π -electrons. Thus, an EC mechanism for the proton reduction was proposed based on electrochemical data. In contrast, **162** reveals a protonation at the ring nitrogen. This protonation is followed by two 1e⁻-reduction steps leading to [Fe⁰Fe⁰N-NH]⁻, which then reacts back to the parent model upon liberating H_2 .

6.6.2 Influence of ligand substitution on the mechanism

6.6.2.1 Models with innocent ligands. To obtain an enhanced basicity at the metal centre and to avoid ligand protonation related complications, numerous models with innocent phosphine and carbene ligands have been designed (Section 4.7). These substituted models have more negative reduction

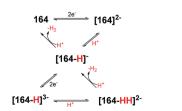


Fig. 101 Catalytic pathway displayed by tetrametallic complex 164.

potentials ($E_{\rm pe}$) than their corresponding hexacarbonyl complexes. Compensating this trend, their catalytic potentials ($E_{\rm cat}$) are, however, often anodically shifted. Due to the increased basicity of the metal core, the models are readily protonated at the metal centre resulting in terminal hydrides or along the metal–metal bond yielding bridging hydrides. These protonated intermediates undergo facile reduction. Notably, **219pdt** follows the CECE catalytic pathway involving the [**219pdt-\muH**] state. The P(OMe)₃ disubstituted complex and similar complexes follow a CE pathway for proton reduction.

Notably, the bridgehead amine in $Fe_2(adt^R)(CO)_{6-n}(L)_n$ (R = Ph, $C_3H_6COOH\ L$ = PMe₃) remains the favoured protonation site. ^{151,191,210,455} However, the electrochemical activity of the hexacarbonyl and phosphine disubstituted complexes differ significantly. In case of unsubstituted models, the protonation exclusively occurs at the bridgehead nitrogen, while in diphosphine substituted models the direct protonation at the Fe-Fe bond is possible as well resulting in potential catalysis CECE pathways. Interestingly, PMe₃ mono- and disubstituted analogues of **109** and **110**, with n- and isobutyric acid N-substitutions, were reported to even use water as proton source and revealed catalytic activity under neutral conditions at approx. $-2\ V$ (Fig. 106). ¹⁹¹

In the proposed catalytic enzymatic pathways, the terminal hydride state is of key relevance for the hydrogen development (Fig. 19-21). Nevertheless, this state is rarely observed for hexacarbonyl complexes. However, unsymmetric substitution patterns at the diiron centre, with electron donating ligand were shown to assist in achieving such terminal hydride (see Section 5.3). Barton and Rauchfuss investigated model 233pdt and found that addition of HBF₄·OEt₂ to a solution of 233pdt in DCM at 0 °C resulted in a terminal hydride state. Notably, this complex catalyses proton reduction at −1.49 V compared to -1.78 V for the bridging isomer. 154 Furthermore, the catalytic cycle proceeds with a second protonation of the one-electron reduced terminal hydride species followed by the release of H₂. Subsequent reduction at less negative potentials regenerates 233pdt. In contrast to the bridging hydride [233pdt-μH]⁺, the model [233pdt-tH]+ comprising a terminal hydride is capable of generating H₂ with a TOF of 5 s⁻¹ with HBF₄·OEt₂. ¹⁵⁴

Likewise, the adt counterpart, (233adt), was investigated for its catalytic activity in DCM using acids of different strengths (ClCH₂COOH (p K_a = 15.30), CF₃COOH (p K_a = 12.65) and $HBF_4 \cdot OEt_2$ (p $K_a > 2$)) at 0 °C (Fig. 102). ⁴⁵¹ In all cases, the catalytic cycle begins with a protonation of the bridgehead amine, followed by an intramolecular proton transfer to the metal centre to afford [233adt-tH]2+. With ClCH2COOH, the cycle proceeds with the reduction of [233adt-tH]+ to afford [233pdt-tH] which upon protonation and another reduction releases H₂. 451,504 With stronger acids (e.g. HBF₄·OEt₂ or CF₃COOH), the bridgehead amine of $[233adt-tH]^+$ is protonated further and the reduction of the double-protonated species gives a $[233adt-\eta^2-(H)_2]^+$ intermediate state. This state then returns to the Fe^IFe^I state upon reductive loss of H₂. Moreover, this study also highlights the catalytic importance of the terminal hydride states and shows that terminal-hydride species generate H₂ much faster (TON 5000 s⁻¹ at -1.49 V)

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Fig. 102 Catalytic pathways adopted by 233adt in presence of acids of different strength

as compared to the bridging congeners (TON = 20 at -1.72 V).451,504

Likewise, the phosphine and carbene substituted asymmetric model $Fe_2(pdt)(CO)_4(\kappa^2-I_{Me}(CH_2)_2PPh_2)$ (304) forms a terminal hydride at low temperature (-90 °C) with HBF₄·OEt₂. Notably, this terminal hydride is inaccessible at room temperature. 450 It was suggested that catalysis proceeds via the bridging hydride intermediate. Other carbene-substituted models, which have been studied for proton reduction, also proceed with a CE mechanism for H2 formation, majorly proceeding via µ-H state with moderately strong acids (CF₃COOH)^{197,363} as well with strong acids (HBF₄·OEt₂).²⁹¹

6.6.2.2 Models with protonable ligands. Besides the protonation of the bridgehead amine in adt-subsite models, a basic residue can be also present at the additional coordinating ligands. 152,505 Active site analogues having proton responsive ligands such as PNP, pyridine and bipyridine attracted interest and the pendant basic sites were anticipated to relay protons to the metal centre and aid the proton-hydride formation for efficient H₂ release (Fig. 103).

Herein, 294Me undergoes N-ligand protonation upon protonation with HBF4·OEt2 in acetone. Remarkably, in DCM a tautomerism resulting in μ -H was observed. ¹⁵² A similar proton

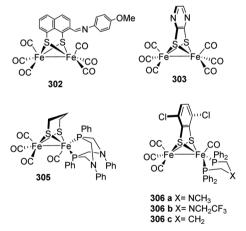


Fig. 103 Structure of selected models discussed in this section.

migration was observed for $Fe_2(pdt)(CO)_4(\kappa^2-L)$ (305, L = (PhP(CH₂)₂NPh)₂) with CX₃SO₃H (X = F, H) resulting in the catalytically active μ -H state. 461 In 294R (R = Ph, Me), the pendant basic site is rendered free to take up additional protons after transfer of proton to the metal centre. Although proton transfer reactions to the metal centres are observed and cause an anodic shift of the catalytic potential. Yet, the rate

constants for HER are low due to sluggish proton transfer and $\rm H_2$ release. 352

Along this line, 306a-c with similar diphosphine ligands $((Ph_2PCH_2)_2X, X = NCH_3 (a), NCH_2CF_3 (b), CH_2 (c))$ was examined. The study displays the proton directing influence of the chelating ligand. In case of 306a, the amine was protonated with HBF₄·OEt₂, while in 306b a hydride bridging the Fe centres was observed. In contrast, in case of 306c which lacks an amine group, protonation at the bridging sulphur was detected.²²⁵ Thus, it seems that pendant basic sites are not the first choice for protonation and the sequence of protonation depends on strength of acid and the corresponding basic site in the mimic. This preference was also illustrated by 307a,b $(Fe_2(pdt)(CO)_5(PPh_2(CH_2)_nPy) (n = 0 (a), 1 (b)),$ wherein protonation with HOTf occurs primarily at the Fe-Fe bond and only thereafter at the N-pyridyl in the phosphine ligand. ⁵⁰⁵ Also, it is to be noted that a secondary sphere protonation might not always be fruitful, e.g. while the PTA ligand (1,3,5-triaza-7phosphaadamantane) is protonated, this protonation is catalytically not very relevant. 302 Likewise, [Fe₂(pdt)(CO)₅(CN)]⁻ (215) undergoes protonation with HOTs at the cyanide ligand but displayed no significant catalytic activity. 153 In contrast, [Fe₂(adt)(CO)₅(CN)]⁻ (308) was found to be catalytically active with HOAc (as well with Cl₂CHCOOH). The detailed catalytic mechanism remains to be elucidated - however, the difference in catalytic activity of 215 and 308 could be reasoned by the involvement of bridgehead nitrogen. 478

Notably, the unsymmetrical model [Fe₂(pdt)(CO)₄(CN)-(PMe₃)] (214) exhibits proton reduction activity in the presence of acids (HOTs, H_2SO_4) at -1.4 V (-1.0 V vs. NHE). IR and ¹H-NMR spectroscopic investigation revealed direct protonation of the Fe-Fe bond affording a bridging hydride. On further reaction with HOTs, protonation of the coordinated cyanide likewise occurs. 150 The double-protonated species undergoes reduction resulting in mixed-valent state which heterolytically releases H2. Notably, no external protons were required to move the cycle forward from the [HFe^{1.5}₂(pdt)-(CO)₄(CNH)(PMe₃)] state. Thus, H₂ was released upon intramolecular hydride transfer. In contrast, the direct congener, ${\bf \left[219pdt\text{-}\mu H\right]}^{+423,424}$ requires additional protons to slowly form H_2 . Moreover, $[Fe_2(pdt)(CO)_4(P(OMe)_3)(CN)]^-$ (309) is protonated solely at the cyanide ligand by HOTs, while HBF4·OEt4 protonates the complex at the metal centre as well as the ligand. Yet, the model displayed no catalytic activity. 153 These examples clearly demonstrate the control of reactivity via the ancillary ligand protonation and highlights that the catalytic activity is achieved through a crucial electronic balance around the metal centres and efficient proton relays from non-remote transferring basic sites. 150,153

Evidently, for Fe₂(bdt)(CO)₆ the introduction of one or two pyridyl-appended phosphine ligands (PPy₃) proved to be extremely beneficial. Not only does the pyridyl site act as a potential site of protonation, but also solubilizes the complex in aqueous conditions. Having achieved a precise balance of the increased electron density at the Fe centre from phosphine ligands and a proximal protonation site, the mimic displays a TOF = $1.8 \times 10^7 \, \rm s^{-1}$

at -0.90 V vs. NHE for Fe₂(bdt)(CO)₅(PPy₃) (310) and 2.7×10^8 s⁻¹ at -0.97 V vs. NHE for Fe₂(bdt)(CO)₄(PPy₃)₂ (311) with dilute H₂SO₄. ⁵⁰⁶

6.6.2.3 Models with redox-active ligands. The active machinery of the enzyme comprises a cubic [4Fe-4S]-cluster attached *via* cysteinyl ligand to the [2Fe-2S]-core. Subsequently, surrogates for the cubic cluster were investigated. With an exception of the appended [4Fe-4S]-cluster to a [2Fe-2S]-model reported by Pickett *et al.*,⁴¹⁶ no such elaborate artificial systems have been reported. Instead, mimics with smaller and less complicated redox active ligands such as bipyridyl,^{371,507} phenanthroline⁵⁰⁷ and phosphole⁴⁹³ have been designed.

Also, in the IMes-substituted pdt complex 248 the presence of the NHC ligand alters the general pathway of the proton reduction. Due to involvement of the IMes ligand, 248 undergoes two successive reduction steps and the first reduction takes place at the metal centre, while the second occurs at the ligand. Hence, it was proposed that 248 follows an EECC mechanism and thus is able to generate H2 from weak acids. Following this double-reduction, protonation at the Fe site forms [H-Fe^{II}Fe^I]. This state undergoes a second protonation along with concomitant internal electron transfer regenerating a Fe^IFe^I state along with H₂ evolution. 362 A similar proton reduction mechanism was noted for in the presence of a bipyridyl ligand in complex 312.371 Herein, the generated dianion is protonated by AcOH. In contrast, in the presence of strong acids such as HBF4·Et2O and HOTs, 312 displays different of proton reductions behaviour (Fig. 104). Due to the bipyridyl ligand, the electron density at the FeFe centre is significantly increased and the metal-metal bond is protonated by the stronger acids resulting in a bridging hydride state $[312-\mu H]^{+}$. The reduced species $[312-\mu H]$ is then protonated,

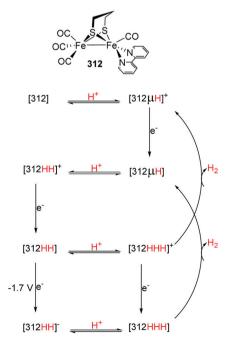


Fig. 104 Catalytic pathway for HER from HBF₄·OEt₂ by **312**.³⁷¹

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followed by another electron and proton uptake resulting in [312-3H]⁺ which on reductive elimination gives H₂. However, this pathway did not account for the catalytic peak observed at −1.7 V. Therefore, it was proposed that [312-HH] undergoes a single-electron reduction at this potential to yield [312-HH]-. This state is then converted to [312-3H] upon protonation which on releasing H₂ forms the [312-μH] state.³⁷¹ Hence, in the presence of strong acids either $[312-\mu H]^+$ is the proton reduction catalyst and operates via a ECEC or [312-μH] following a CECE mechanism.

Another example wherein the ligand actively participates in the catalytic pathway is 313 modified with a phosphole. The complex displays three cathodic peaks - the first corresponds to the reduction of the ligand. 493 The other two peaks are associated with two single-electron transfers to the diiron centre resulting into Fe⁰Fe⁰. According to spectroelectrochemical and theoretical modelling data, the catalytic cycle is initiated by a proton-coupled electron transfer and protonation occurring at -1.44 V. Here, protonation occurs at the Fe centre, leading to a mixed valent bridging hydride state [Fe^{II}Fe^Iµ-H] and another at the pyridine nitrogen of the ligand, which serves as the second protonation site. This double-protonated state undergoes a second two-electron reduction at -2.00 V accompanied by another ligand protonation (Fig. 105). This state was determined to be the resting state of the catalyst. The protonation and subsequent release of H₂ from this state is the rate-determining step of the reaction. Hence, in this cycle the ligand serves as an electron reservoir and is involved in electron transfer to the metal centre. Additionally, 313 also catalyses proton reduction from dilute H₂SO₄ with an overpotential of 0.66 V and TOF $7 \times 10^4 \text{ s}^{-1}.^{493}$

In addition to the aforementioned models, many other mimics with redox active ligands such (bma = 2,3-bis(diphenylphosphino)

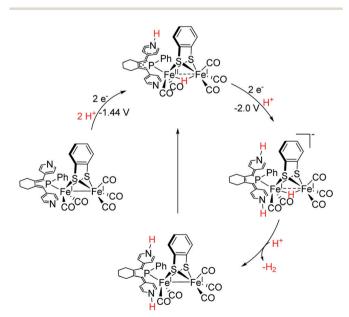


Fig. 105 HER by 313 with an appended redox active ligand from Et₃NHBF₄ in DCM.

maleic anhydride (bma), bpcd (4,5-bis(diphenylphosphino)-4cyclopenten-1,3-dione)), 311,313,314,508 fullerene, 509 ferrocene 312,349 and carborane bis-phosphine⁵¹⁰ have been reported. However, involvement of these redox active moieties in catalytic mechanisms is either questionable or remains to be investigated in detail.

Based upon numerous detailed investigations of hydrogen evolution activity by diverse Fe^IFe^I subsite models, the observed trends and generic behaviour can be summarised as follows:

- To enable H2 evolution from weak acids by hexacarbonyl models, these complexes are required to undergo reduction before being protonated.
- CO-substitution by stronger donating ligands such as phosphines or carbenes cause the diiron core to attain increased basicity. Thus, the catalytic cycle may begin via direct metal protonation.
- If there is an accessible basic site available in the model complex and an acid of suitable strength is employed, the catalytic cycle is facilitated by its protonation.
- Involvement of a redox active ligand can cause a double reduction to occur, i.e. one at the metal centre and another at the attached ligand before the model is protonated.
- In case of bidentate substitution at the Fe centre, the chelating isomer is reported to be more efficient for H2 generation than the symmetrically substituted bridging counterpart.
- pK_a of acid used as the proton source determines the adopted catalytic pathway for H2 evolution and has profound influence on the catalytic activity and over-potentials.

Conclusively, extensive electrochemical analysis revealed various mechanisms (EECC, ECEC, ECCE, CECE, CEEC) and shed light on the involved iron centres redox behaviour within the H₂ evolution activity in acidic media.

6.6.3 Influence of water as solvent on the mechanism. Numerous model complexes with modified secondary sphere have been designed to realise catalysis in aqueous electrolytes. 165,511-514 Many of these systems are also photocatalytically active and will be discussed in Section 7. Thus for the Fe^IFe^I models to achieve electrocatalysis in presence of water, the bridgehead groups were modified (Section 4.1) or ligands assisting in solubility and stability were appended. 302,486 Herein, introduction of functionalized sugar on the diiron core hexacarbonyl, resulted in 33 & 34. These models were soluble in H₂O:CH₃CN (5:1) and were found to be catalytically active in presence of acetic acid at potential of ca. -1.6 V for 38 and ca. -1.8 V for 39 (vs. $Ag^{+/0}$ 0.01 M $AgNO_3$). 133

Attempts to achieve proton reduction catalysis in water are not limited to the modifications of the bridgehead substituents but have been extended to systems with modified ligands such as phosphines 129,130 and peptides. 306 Herein, tris(morpholino)phosphine (TMP) mono and disubstituted pdt models were tested for their proton reduction capabilities in different water: acetonitrile mixtures in the presence of AcOH. 129 Both models proved to be electrocatalytically active operating via ECCE mechanism. Similar effects were reported for the introduction of a charged, quarternary ammonium-modified PNP ligand, enabling H₂ production from HOAc in MeCN following the same mechanism. 130 First, the reduced monoanion [Fe^IFe⁰] is protonated resulting in [H-Fe^{II}Fe^I] which on further

Fig. 106 Proton reduction in H₂O by model 110PMe₃. 191

protonation produces [HH-Fe^{II}Fe^I]⁺. This species generates H₂ upon reduction and the starting complex is regenerated. Notably, an increased catalytic activity (TON_{bridging} = 14.4 in MeCN vs. 25.7 in MeCN: $H_2O(3:2)$) in mixed-solvent systems was reported for these complexes. 129,130 Likewise, electrochemical properties of peptide modified models were significantly affected by presence of water and showed an up to 0.1 V anodic shift in their reduction potentials along with increased catalytic currents.306

Furthermore, adt-models 109 to 111 with carboxylic acid functionalities were synthesised and studied for their redox activity to benefit from the hydrogen bonding properties of their carboxylic acid residues. Cathodic scans displayed reduction peaks (ca. -1.6 V) which shift about 400 mV towards less negative potentials upon addition of one or two equivalents of HOTf suggesting protonation of nitrogen of the adt-bridge. Notably, complex 109 displayed an additional reduction peak, which was attributed to the proton coupled one-electron reduction process assisted by the carboxylic acid group. 191

Furthermore, improved, second generation mimics were achieved upon replacing two carbonyls with PMe3 and were tested for their catalytic capabilities in water. Model 110PMe₃ showed exclusive enhancement in reductive peak upon addition of one equivalent of water. However, in case of model 109PMe₃ a small increasing peak at -1.7 V was observed, with subsequent addition of water which shifted to -1.5 V. Herein, the carboxylate group was involved in stabilising the protonated amine (Fig. 106) via hydrogen bonding. Distant orientation of carboxylate in 111PMe₃ might, however limit this stabilisation and hence no comparable catalysis for this model was recorded. 191

Immobilisation of H-cluster models on electrodes

At the end of Section 4.2.3.1 we already mentioned that carboxylic acid functionalised [FeFe]-hydrogenase models are

suitable for the attachment to surfaces. This is based on their reactivity towards amines that might be covalently bound to e.g. glassy carbon or fluorine doped tin oxide (FTO) electrodes, forming amides.141 This general concept to attach a homogeneous catalyst to electrode surfaces to generate immobilised systems was utilised by several groups. 142,515,516 Interestingly the results are diverse. A gold or carbon immobilised pdt-model and a FTO immobilised bdt-model catalyse proton reduction from HBF₄·Et₂O acidic and chloroacetic acid acidic acetonitrile solutions, respectively but are rapidly inactivated by the loss of carbonyl ligands, catalyst leaching or hydrolysis under these conditions. 515,516 Conversely, no catalytic proton reduction was found for an o-xyldt-model that was immobilised via a carbon surface bound p-tolylformamide. 142 Other immobilisation techniques include formation of triazoles via Cu(1)-catalysed Huisgen addition or formation of an Au-S bond via thiols.517-519 Latter includes a pdt-like model that comprises a -PH₂(CH₂)₂SH ligand for the attachment and is therefore the only surface attached model that is not bound via the bridging disulphide. Interestingly, thus study revealed that the parent compound 314 is inactive regarding proton reduction in solution, while its immobilised counterpart shows catalytic proton reduction upon addition of acetic acid. This phenomenon was explained by the inability of 314 to undergo a reduction process to a Fe⁰Fe^I or Fe⁰Fe⁰ species, which was however shown to be at -1.87 V and -2.24 V vs. Fc^{0/+}, respectively by Liu et al. in 2009215 and is also visible in the respective cyclic voltammograms in the works by Darensbourg and coworkers. 518,519 Four years later, Zaffaroni et al. described a 160 mV anodically shifted operational overpotential for the FTO immobilised bdt-model compared to its homogenous counterpart. 516 This kinetic effect due to the binding of an homogeneous catalyst to an electrode surface might have been the reason for the observation of Darensbourg and coworkers

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Fig. 107 Schematic presentation of covalent attachment for the [FeFe]-hydrogenase model **314** on modified graphite surfaces. Reproduced from ref. 517 with permission from The Royal Society of Chemistry.

as well. In 2017, Dey and coworkers reported on the first adt-like model that was covalently linked to a graphite electrode surface via Huisgen addition (Fig. 107). In the opposite to the systems described above, this system is reported to be stable for several hours under turn-over conditions and shows a high faradaic yield of 90.7% for $\rm H_2$ evolution after 1200 s. $\rm ^{517}$

6.8 Perspective for the electrocatalytic proton reduction by H-cluster models

The electrochemical investigations coupled with theoretical calculations as well as spectroscopic investigations significantly contributed to describe different pathways for proton reduction by the [FeFe]-hydrogenase models. It has to be noted that the strength of the acid employed significantly affects the proton reduction pathways. The models possessing intrinsic bases, such as the bridgehead amine, conduct protonation at milder potentials in comparison to the models lacking such sites. Furthermore, the donating capabilities of the ligands also influences the attainable redox states in the proton reduction catalytic cycles. When the carbonyl ligands are substituted by donor ligands, the reduction potential of the models display cathodic shifts. Accompanying this cathodic shift, the protonation is facilitated due to increased electron density at the metallic centre. A few terminal hydride bearing species have been observed and it has been ascertained that the terminal hydride species are more efficient catalysts in comparison to the thermodynamically more stable bridging hydride state.

Even after the aforementioned advances, challenges to develop systems with matching efficiencies to the natural system exists. Unlike the enzyme, the majority of the models are catalytically active in organic solvents and utilize organic acids to generate H_2 . They do not follow the bio-catalytic

pathway. Nonetheless the electrochemical studies on the synthetic systems impressively uncovered the interplay between the basicity and reduction potentials of the models and shed light on the hydrogen generation mechanism.

VII Photocatalytic proton reduction by H-cluster models

The development of artificial systems that efficiently mimic the natural photosynthetic pathways and allow for a conversion of solar energy into storable and useable forms has gained major attention. Due to their unprecedented H₂ generation capacity, [FeFe]-hydrogenases are considered a suitable choice for engineering sustainable photocatalytic machineries although these natural enzymes are not photocatalytically active by themselves. Nonetheless, the need to develop renewable alternatives for H₂ production, drives the hydrogenase community to take profound interest in designing such systems. Numerous models of the [FeFe]-hydrogenase have been previously described, which in conjunction with suitable light harvesting systems could form promising systems for light-to-fuel conversion technologies (Fig. 108).

Photocatalytic hydrogen producing assemblies are commonly comprised of a photosensitizer (PS) capable of harvesting light along with a proton reducing catalyst, a sacrificial electron donor, which is capable of restoring the actual redox state of the sensitizer, and a proton donor. For diironcomplexes in particular, a potential mechanism for the H₂ formation involves photon absorption by the sensitizer (organic moieties, semiconductors), followed by an electron transfer from the photoexcited sensitizer to the diiron catalytic centre.

Fig. 108 Potential mechanism for the photosensitised hydrogen evolution PS = photosensitizer, SED = sacrificial electron donor.

Subsequently, protonation of the reduced catalyst takes place. This process is followed by another electron transfer to the bimetallic catalytic centre resulting in $\rm H_2$ formation, thereby closing the cycle. $\rm ^{521,523}$

A variety of architectures with different photocatalytic abilities have been designed and will be discussed herein. These systems either vary in the choice of photosensitizer or the proton reduction centre. In general, two approaches were adopted to link photosensitizers to the active site models. In the first strategy, attachment of the chromophore to the bridgehead atom through chemical modifications was performed. Furthermore, the photosensitive moiety can be directly linked to the diiron site.

7.1 Covalent attachment of photosensitizers to H-cluster models

7.1.1 Attachment to the dithiolate bridge. Following the above-mentioned strategy, Ott and coworkers designed elaborate systems. The first example reported by the researches is comprised of an adt-model functionalised with a modified $[Ru(terpy)_2]^{2+}$ complex (terpy = 2,6-bis(2-pyridyl)pyridine) connected *via* an ethynyl linker (315, Fig. 109). The linker provides

separation of the photosensitizer and the catalytic centre along with enhancing the excited state lifetime of $[Ru(terpy)_2]^{2^+}$. In comparison to the non-derivatised Ru sensitizer, this system displays a 30% quenching of the MLCT state. Nonetheless, electron transfer to the catalytic site was not observed and the amount of H_2 generated was low.⁵²⁴ Additionally, a modified PDT model (Fig. 109) was linked to the $[Ru(bpy)_3]^{2^+}$ sensitizer through an amide linkage to afford 316. However, no electron relay between the photosensitizer and the diiron site was observed, which was explained by the high flexibility of the linker in homogenous solutions. The electron transfer from the excited photosensitizer to the diiron site thus becomes thermodynamically unfavourable.⁵²⁵

Following these reports, Song *et al.* incorporated a tetraphenylporphyrin group (TPP) on the bridgehead nitrogen of $Fe_2(adt^R)(CO)_6$ (R=p- C_6H_4CHO), *via* covalent bonding resulting in 317. ⁵²⁶ With the improved light absorbance of TPP and the longer excited state lifetime, it was anticipated to be favourable for the electron transfer processes towards the diiron site. Indeed, the electron was shown to be readily transferred from the photoexcited sensitizer to the diiron center. ⁵²⁶

Subsequently, a three-component system, 318 built from $Fe_2(adt^R)(CO)_6$, a Re-photosensitizer and ferrocene was established. Here, the catalytic site and the sensitizer were separated by an ethynyl linker that provided appropriate separation for fast electron transfer without electron recombination at the sensitizer. Utilizing ascorbic acid as proton source and as the sacrificial electron donor, the system displayed a TON of $0.35.^{527}$

Despite of their obvious potentials noble metal-based photosensitizers are not particularly desirable due to their expensive production and involvement in complex deteriorating

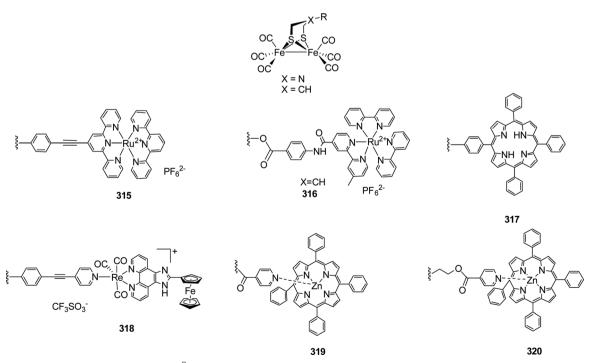


Fig. 109 Models 315 to 320. Examples of Fe₂(xdt^R)(CO)₆ models modified for photocatalytic purposes.

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pathways. As a noble metal free alternative, a zinc tetraphenylporphyrin (ZnTPP) unit linked to ADT via the amine bridge (319) to achieve considerable photochemical HER activity. As expected, significant fluorescence quenching was observed suggesting an intramolecular electron transfer from ZnTPP to the diiron centre. However, the exact electron transfer pathway for such systems remains elusive. 188 Along this line, complex 320 was established and investigated for its H2 production capability. Interestingly, the complex 320 revealed a faster electron transfer from the TPP moiety to the diiron unit as compared with the covalently linked systems. The newly established complex also allowed to overcome charge recombination required for fast catalytic conversion processes but only showed low turnover numbers. This behaviour was ascribed to the photo instability of the catalyst. 528 Along this modification strategy, model 181 was reported comprising a covalently attached ZnTPP/naphthalene unit. A thermodynamically favourable electron transfer from the sensitizer to the catalytic site was suggested based on electrochemical investigations with the first reduction potential of the mimic being less negative than the oxidation potential of the photoexcited ZnTPP unit (-1.65 V vs. -1.74 V). In addition, Wasielewski and coworkers investigated the reactivity of 321a-c. It was observed that employing a second electron donor in these systems is beneficial, increasing the lifetime of the reduced diiron centre and facilitates the catalytic activity. Likewise, upon extending the linker length between the photosensitizer and the catalytic site, the electron transfer to the diiron centres is favoured and the electron recombination time period increases by a factor of 7.5. Hence, only 321c displayed significant photocatalytic activity with the additional phenyl group disfavouring quenching due to efficient energy transfer to the ferrocene moiety. However, the observed photocatalytic activity is limited by degradation of the diiron centre upon irradiation (Fig. 110).²⁵²

7.1.2 Attachment by CO-ligand substitution. As the secondary amine of the adt-bridge was believed to play a crucial role in the proton reduction, modifying this position was

321c R=Ph-4-Fc

Fig. 110 Modified models of 170 for photocatalytic activity

Fig. 111 $\,$ Model 322 with direct attachment of the photosensitiser to metal centre.

anticipated to severely influence the hydrogen evolution. Likewise, the electron relay machinery is linked to the iron centre *via* a cysteinyl ligand in the natural system. Therefore, a direct linkage of the photoactive groups to the iron centre emerged as an attractive design strategy. A first approach to realize such a system was provided by the synthesis of **322**. Subsequently, electrochemical studies revealed that the bridging acetylenic ligand (Fig. 111) is the most easily reduced component. Furthermore, the generated oxidised diiron site is unstable, which accounts for the light sensitivity of the dyad and limits further photochemical investigations and catalytic applications.⁵²⁹

In a comparable approach, complex 323 was prepared. Herein, the platinum(π)-polypyridyl alkynyl sensitizer is attached via an isonitrile group to the Fe₂(pdt)(CO)₅ moiety. The resulting system displayed long-lived MLCT states. However, the important reduction of the Fe¹Fe⁰ state is thermodynamically unfeasible and thus explains the overall low catalytic efficiency. Furthermore, a series of cyanide ligand modified hydrogenase mimics was reported (324a–c, Fig. 112) using rhenium photosensitizers as prospective catalyst systems but revealed comparable problems. 531

Kluwer and coworkers reported a macromolecular system carrying two photosensitizers covalently attached to the Fe centres *via* phosphine groups [Fe₂(pdt)(CO)₄{PPh₂(4-py)}₂] and modified this complex with zinc(π)porphyrin chromophores **325** (Fig. 113). However, this design strategy proved to be unsatisfactory as only two turnovers per catalyst under optimized conditions were observed.⁵³² Rauchfuss and coworkers reported on the photocatalytic activity of [**219pdt-μH**]⁺ and [**231pdt-μH**]⁺ in the presence of ferrocene as recyclable electron donor. Strikingly, this system was shown to overcome the limitation of the otherwise required high reduction potentials for the hexacarbonyl systems (-1.33, -1.4 V νs. -1.66 V for **20**).⁵³³

7.2 Photocatalytic multi-component systems

Although a significant number of reports exists for photosensitizer-bound proton reduction sites, all systems display severe limitations and none of them allows for sufficient H₂ generation. An alternative strategy to build photocatalytic hydrogenase mimics is the use of multicomponent systems *i.e.* consisting of unlinked photosensitizers and H₂-formation sites. One of the early reports in this field describes the incorporation of quantum dots (QDs) as potential photosensitizers. Quantum dots possess superior visible light absorption

Fig. 112 Cyanide modified photocatalytic systems with Pt- and Re-complexes

Fig. 113 Porphyrin modified supramolecular models relevant for photocatalysis.

375b R₁=(4-MeO)Ph

abilities and importantly, are stable under aqueous conditions. Notably, 3-mercaptopropionic stabilised nanocrystal quantum dots in the presence of a [2Fe-2S]-cluster and ascorbic acid revealed promising turnover numbers and turnover frequencies of up to 505 and 50 h⁻¹ within 10 h of illumination.⁵³⁴

Likewise, the utilization of Ru-based light harvesters was suggested as a potential replacement for the biologically relevant [4Fe-4S]-clusters in an artificial system. Due to its similar reduction potential compared to that of the diiron dithiolate hexacarbonyl complex, $[Ru(bpy)_3]^{2+}$ was proposed as an efficient photo sensitizer. Upon adopting the diethyldithiocarbamate anion (dtc⁻) as an electron donor, the challenge of reverse electron transfer was surmounted, as the electron transfer generated thiyl radicals, from the dithiolate anions, which underwent quick dimerization and thus extend the excited state lifetime of [Ru(bpy)₃]⁺. Still, this system suffered from photo-bleaching and protonation of the anionic quencher. 535 Multiple phosphine variants of pdt (20) and adt^{Bn} (91) models were subsequently investigated utilizing ascorbic acid and [Ru(bpy)₃]²⁺ and achieved a maximal TON of 4.3 under optimum conditions. 536

Especially the chlorine substituted bdt model 150 is a suitable candidate for studying the photochemical H₂ production due to its less negative reduction potential in combination with its high stability and capability to undergo reversible reductions. Along this line, a composite system consisting of this particular mimic, [Ru(bpy)₃]²⁺ as photosensitizer and

ascorbic acid as proton and electron donor was subsequently investigated and revealed a TON of 200 along with a TOF of 2.7 min^{-1} in DMF/H₂O solutions at pH of $5.5.^{537}$

7.3 Confinement of the photocatalytic system

Although chemists have succeeded in designing active site analogues, in which the photosensitizer is attached to the catalytic site, yet such systems are far away from any application as these systems are limited in stability, solubility and H2 generation. An alternative strategy to achieve an appreciable activity similar to that of the natural system, the skilful alteration of the secondary coordination sphere of the catalyst was anticipated to tune the solubility and stability.

Notably, the microenvironment influences the photocatalytic performance considerably - to this end, catalytic diiron models were either embedded in a polymeric scaffold or incorporated into nanomaterials, generating improved heterogeneous proton reduction assemblies that will be described in the following sections. 165,187,511,538-541

7.3.1 Polymer support

7.3.1.1 Synthetic polymers. Polymers are one potential support to incorporate the catalytic centre and photosensitizer as they allow to shape the environment of the catalyst and allow for the encapsulation of additional substrates. For instance, the sulphonate modified adt-model Fe₂(adt^R)(CO)₆ (326, R = p-C₆H₄SO₃⁻Na⁺) along with Ru-polypyridine based photosensitizers were grafted onto a phospholipid membrane (DOPC 1,2-dioleoyl-sn-glycero-3-phosphocholine). This self-assembled system displayed a significantly increased photocatalytic activity under acidic conditions (pH 2.6) in comparison to the nonimmobilised models. A reason for this performance enhancement was found in the close proximity of the relay groups involved in electron transfer processes. The highest photocatalytic activity was reported utilizing ascorbic acid with a TON of 57.

In addition, activity tests for catalysts embedded in lipids were conducted as well and an activity enhancement was observed with lipid layers with higher order such as DOPC and DMPC (1,2-dimyristoyl-sn-glycero-3-phosphocholine). However, the activity of these membrane assemblies is yet still limited by photosensitizer degradation.542

Furthermore, a series of macro models was established upon reacting Fe₂S₂(CO)₆ with a Fréchet-type dendron in Chem Soc Rev Review Article

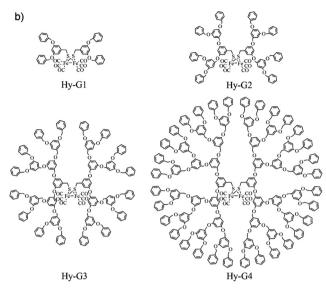


Fig. 114 Structures of dendrimer based [2Fe]_H mimics. Reprinted from ref. 511 with permission from John Wiley and Sons, Copyright 2013.

THF (Fig. 114). The resulting dendrites were likewise studied for their photocatalytic activity in the presence of $[Ir(ppy)_2-(bpy)]PF_6$ (ppy = 2-phenylpyridine, bpy = 2,2'-bipyridine) as the photosensitizer and triethylamine as the sacrificial electron donor in water/acetone (1:9) mixtures. Notably, the largest dendrimer (Fig. 114, Hy-G4) displayed a high TON of 22 200 with a TOF of 7240 h⁻¹. The hydrophobic environment created by the dendrite around the catalytic site was herein suggested to allow for a close interaction of the neutral sensitizer and the charged photosensitizer and reasoned to be origin of the high activity. 511

Furthermore, a three-component assembly – branched polyethyleneimine functionalised with an isocyanide ligand modified pdt mimic (327), MPA–CdSe quantum dots and ascorbic acid – was shown to be photocatalytically active in water over a broad pH range and provided high TON values of up to 10 600 at neutral pH. Herein, incorporating MPA–CdSe QDs into the systems was found to be beneficial as these are easily dispersed in water and have intense photon absorbing properties. Likewise, the branched polymer has numerous amine groups, which are capable of binding to the specific mimic and potentially act as proton relays to the catalytic site. Most important, the polymer stabilizes the CdSe quantum dots against aggregation and thus contributes significantly to the high activity and stability.⁵³⁸

A breakthrough study for the photocatalytic $\rm H_2$ production was reported on systems containing an isocyanide PDT model which was attached to polyacrylic acid (PAA) (Fig. 115). Here, the hydrophilic PAA assists in the solubilisation of the otherwise insoluble catalyst system in water and TONs of up to 27 135 and TOF 3.6 s⁻¹ were obtained at pH 4. Notably, due to the surrounding PAA, the 2-mercaptobenzoic acid (MAA) stabilized CdSe quantum dots (QDs) were protected from aggregation. Furthermore, the polymer brings the sensitizer and the catalytic site in close proximity to each other and thereby allows for an enhanced electron transfer. 165

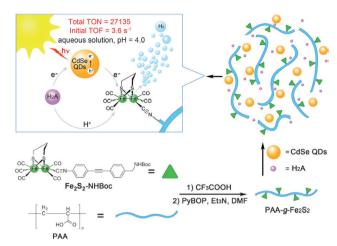


Fig. 115 Synthetic pathway towards modified pdt and representation of catalytic activity. Reprinted from ref. 165 with permission from John Wiley and Sons, Copyright 2013.

7.3.1.2 Peptides and proteins. As an alternative to synthetic polymers, a combination of active site models with biological scaffolds was used to tune the secondary coordination sphere and with it to optimize catalyst solubility and activity – for example, 219pdt was encapsulated in a low molecular weight hydrogelator (Fmoc–Leu–Leu). This secondary sphere prevents the polymerization of 219pdt and is hence positively affecting its stability (Fig. 116). Notably, a network of H-bonds was observed when the model is implemented within the gel confirming a significantly higher stability in aqueous media. 512

In addition, the covalent attachment of modified Fe₂(pdt)(CO)₆ to the cavity of a β -barrel protein apo-nitrobindin (apo-NB) was reported. Model **328** bearing a maleimide group covalently binds to cysteine residue (Q96C mutant) within the protein and this assembly along with [Ru(bpy)₃]²⁺ and ascorbate in 50 mM Tris/HCl at pH 4.0 and 25 °C allowed for the generation of hydrogen (TON up to 130 for 6 h).⁵⁴³

Furthermore, employing cytochrome C which functions as native electron relay in the natural system, an artificial metalloenzyme was constructed via attaching the [2Fe–2S]-site to the CXXC motif (Cys14, Cys17) of the apo-enzyme, giving a Fe₂(S-Cys)₂(CO)₆ unit. Subsequently, this synthetic machinery was studied for photocatalytic H₂ development using [Ru(bpy)₃]²⁺ and ascorbic acid. The assembly in aqueous media generates H₂

Fig. 116 Model 328 for attachment to the β -barrel protein.

with a TON of 80 over 2 h at pH 4.7.⁵⁴⁴ To further investigate the electron transfer process between the photosensitizer and the catalytic centre, a more elaborate system was designed, wherein the ruthenium sensitizer was coordinated to the adjacent histidine residue (CXXCH sequence) of cytochrome C_{556} . After 2 h of irradiation, catalytic activity for the photo-induced hydrogen development with a TON of 9 at pH 8.5 was observed.⁵⁴⁵

Although Ru-based photosensitisers and pdt-derived model 23 were non-covalently encapsulated in a hydrophobic (horse spleen) apo-ferritin cavity, this confinement enhances the solubility of the components along with providing a close proximity for an improved electron exchange. However, the highest recorded TON for such a system was 5 and the photosensitizer [Ru(bpy)₂(dpqp)]²⁺ was found to limit the activity of the system. ⁵⁴⁶ In contrast, if the sensitizer [Ru(bpy)₃]²⁺ was not incorporated in the ferritin pocket and rather homogenised in solution, a TON of 31 was reported. ⁵⁴⁷

Recently, Li and coworkers reported self-assembled ovalbumin (OVA) nanogels which incorporate **20**. This incorporation improved the photocatalytic efficiency for H_2 evolution in acid aqueous solution by 15% under optimized conditions. It is important to mention that acid induced structural changes (unfolding of α -helix to β -sheets) within the nanogels contribute massively to the overall performance. ⁵⁴⁸ **20** was likewise linked to an artificial dithiol amino acid, which was in turn incorporated into an alpha helical peptide scaffold. This assembly, using $[Ru(bpy)_3]^{2+}$ and ascorbate generates H_2 with a TON of 84 in acidic aqueous solutions under light irradiation. ⁵¹³

7.3.1.3 Polysaccharides and oligosaccharides. To imitate the protein cavity of hydrogenases and to favour catalytically active intermediates Darensbourg and coworkers incorporated 326 into a β-dextrin. The anionic sulfonate group herein assists in the dissolution of the complex within dextrin. Notably, upon protonation with HOAc in water, the system displayed an electrocatalytic peak at -1.4 V vs. Ag/AgCl. However, ¹³C-NMR studies revealed that the encapsulation of the model into dextrin is rather dynamic and not sufficient enough to artificially reproduce the natural protein environment. 131 Also, an improved version of this model with various phosphine ligands (PTA, PMe₃, P(OMe)₃, PPh₃) was reported, but these models display rather unsatisfactory reduction properties. 189 Following this research Sun et al. studied the photocatalysis of this encapsulated system using Rose Bengal as well as Eosin Y as sensitizers. 549 These organic sensitizers also participated in the host-guest interaction with β - and γ -dextrins and allowed for a faster electron transfer processes from the photosensitizer to the diiron centre. This improved electron transfer resulted in an enhanced quantum efficiency for the system.

In another attempt, chitosan – a naturally occurring polysaccharide – was used to greatly improve the solubility and photo dependent H_2 producing ability of $\bf 91$ in the presence of CdTe quantum dots capped with 3-mercaptopropionic acid (MPA–CdTe QDs) and ascorbic acid. Along this line, chitosan can be converted into a multi cationic system when its surface amine and phenolic groups are exposed to acidic conditions. These cationic species show great affinity for the negatively charged MPA-CdTe QDs and in return favour the electron transfer process to the imbedded catalyst. Here, the confined environment provided by chitosan allows for a close interaction of the catalyst, photosensitizer and the proton donor and results in an unprecedented TON of up to 52 800 and a TOF of 1.40 s⁻¹ Seemingly, a modified environment to increase the stability of the catalyst and to overcome limitations of the non-modified systems is of utmost importance.⁵³⁹

7.3.2 Micelle systems. Molecules bearing amphiphilic groups undergo self-assembly to form water-soluble clusters. This assembly allows the application of hydrogenase mimics to conduct catalysis under aqueous conditions. To achieve photocatalysis in water, the adt^{Ph} and adt^{Bn} complexes 90 and 91 were both incorporated into sodium dodecyl sulphate (SDS) micelles along with Re^I(4,4'-dimethylbpy)(CO)₃Br or Re^I(1,10-phenanthroline)(CO)₃Br as photosensitizer. However, this attempt only afforded low amounts of H₂ (TON 0.13). ⁵⁵⁰ In a similar approach, 20 was embedded in a self-assembling system by using an amphiphilic P-NB polymer, ^{132,551} resulting in an active system in water and displaying a TON of 133 in the presence of [Ru(bpy)₃]²⁺ and ascorbic acid. The close proximity of the electron donor and the catalytic site, established due to micellar incorporation, facilitates the electron transfer. ¹³²

Along this line of modification, **149** displayed reasonable catalytic activity (TON 117) in water under basic conditions (at pH 10.5) upon incorporation into sodium dodecyl sulphate (SDS) micelles. Eosin Y was herein utilized as a photosensitizer and triethyl amine as electron donor. In the absence of the SDS micelles, the catalytic activity of the complex is lowered to one fourth of the maximum activity.⁵¹⁴

7.3.3 Metal-organic frameworks and molecular sieves. Another strategy to enhance the photoactive performance is to use metal-organic frameworks (MOFs). MOFs are 3D systems built from covalent linkage of metal ions and organic moieties. They are capable of loading the catalysts on top of their surfaces and also possess the advantages of easy product separation, reusability, and enhanced stability due to their heterogeneous nature. Moreover, they can provide structural support along with their organic groups imitating the second coordination spheres of natural systems. Following this strategy, a novel assembly was described with 329 attached to a robust zirconium-porphyrin based metal-organic framework (ZrPF). Along this line, the zinc-porphyrin [tetrakis(4-carboxy-phenyl porphyrin) zinc (ZnTCPP)] was linked to the Zr cations of the framework via carboxylic groups. Here, the ZnTCPP moiety acts as anchor for the diiron site within its cavity along with its function as photosensitizer. The zirconium-porphyrin MOF ZrPF is a chemically stable, rigid framework with a high surface area and enables light dependent H2 activity. After irradiating (>420 nm) the catalytic system for 120 min in the presence of ascorbic acid in water at pH 5 about 3.5 µmol H₂ were detected.552

Utilizing the same conceptual approach, 330 loaded onto the zirconium terephthalic acid MOF UiO-66(Zr) revealed an improved photocatalytic activity in presence of ascorbic acid

Chem Soc Rev Review Article

Fig. 117 Subsite models for photocatalytic HER systems

and $[Ru(bpy)_3]^{2+}$ as compared to the individual components and likewise circumvents the undesired charge recombination.⁵⁵³ As studies on this system revealed a potential clogging of channels by ion pairs generated by reduction of catalyst, **159** was loaded onto *e.g.* the chromium MOF MIL-101 which revealed enlarged pore sizes of 29–34 Å as compared to UiO-66(Zr) with 8–11 Å. Subsequent photochemical investigation with different catalyst to MOF ratios revealed that H_2 production is proportional to catalyst loading in the MOF thus following first order kinetics.⁵⁵⁴

In addition, bis(2-phenylpyridine)(2,2/-bipyridine)iridium(III) chromophores were introduced on a diiron hydrogenase mimic by click-chemistry. The modified chromophores were then incorporated into a K⁺-exchanged molecular sieve, MCM-41. When examined for their photocatalytic activity in presence of triethylamine (TEA), CH₃CN and H₂O (9:1) at pH 10, the composite enabled the formation of 11.8 μ L H₂ with 5.5 mg composite and a loading 19.1 μ mol g⁻¹. The enhanced activity of the incorporated system was attributed to the complex stabilisation by the molecular sieve. ⁵⁵⁵

7.4 Attachment of the photocatalytic system to nanoparticles

To overcome the limitations of high cost and insufficient stability associated with above mentioned complexes, semiconductors can likewise be employed as photosensitizers. Prerequisites of an ideal semiconductor are a large band gap possessing a conduction band with a redox potential capable of reducing protons as well as good water solubility. In that sense, two sub-site models, 90 and 331 were incorporated onto the surface of ZnS. ZnS is a noble metal-free and highly photoactive material. However, it suffers from a rapid recombination of the generated electron/holes pairs. 556,557 To this end, when the diiron subsite models are adsorbed on the ZnS surface and triethanolamine (TEOA) was used as the electron donor, photocatalytic experiments showed a TON of 3400 and 4950 for C1&ZnS and C2&ZnS in DMF: H₂O system (9:1).¹⁸⁷ Similarly, aniline functionalised 332 adsorbed on ZnS nanoparticles was stable for up to 38 h of irradiation and displayed a TON of 2607 and initial TOF of 100 h⁻¹ in presence of ascorbic acid at pH 4.6.540

Furthermore, **159** was attached to the surface of an oligoethylene glycol shell modified ZnO along with the natural pigment betanin as the light harvester. The modified ZnO

nanocrystals revealed an extended excited electron lifetime in the conduction band and showed enhanced charge separation as well as accumulation of reactive electrons for the photocatalytic process. With trifluoroacetic the system displayed a TON of 11 and stability up to 6 h. Under otherwise unmodified catalyst conditions, solely a stability of up to 2 h was observed. Furthermore, with modified nano cathodes *i.e.* cross-linked indium phosphide nanocrystal array grafted with ${\rm Fe_2S_2(CO)_6}$ (1), a photoelectrochemical efficiency of more than 60% was witnessed at -0.9 V νs . SCE (Fig. 117). 558

7.5 Tailoring the photocatalytic proton reduction by H-cluster models

These studies report significant developments in tailoring artificial scaffolds for the light-dependent hydrogen generation and support the crucial role of the microenvironment in tuning the properties of these systems. Also, remarkable progress was achieved in obtaining functional systems under aqueous conditions or systems which avoid the use of noble metals for hydrogen generation. Presently, investigations of mechanistic pathways are under progress to provide detailed explanation of the functioning of these systems. Nonetheless, to match the activities of the natural systems, further modulations are required to generate systems which are suitable for incorporation in technological devices.

VIII Conclusion

In conclusion, we herein highlight the development in the [FeFe]-hydrogenase research over the last decades. While initially, structural alterations of mimics along with spectroscopic investigations were performed to understand the natural enzyme, pinpoint its various states and relate them to electronic and structural changes in the active centre, later investigations focused mainly on the electrochemical properties of such systems with the aim to find suitable proton reduction catalysts. Recently, the understanding of the enzymatic properties as well as direct alteration of the entire enzyme became the focus of many investigations due to the possibility to implement artificial synthetic mimics into the natural environment. However, concerning the enzyme and its functional mechanism, important questions remain to be answered. Up to date, no

conclusive evidence was presented allowing for the determination of an exact consecutive sequence of proton/electron transfer events in the catalytic cycle. Hence, several plausible catalytic cycles were presented, and future investigations will have to provide evidence in favour of one over the others. Furthermore, understanding of the functional role of the enzyme backbone surrounding the H-cluster is deficient, e.g. reflected in the potential role of a second substrate/proton channel in the native enzymes. Along this line, artificial surroundings of the H-cluster such as polymers are expected to be a growing field of interest. In general, however, we expect hydrogenase-research to become more applications oriented. For example, enzymes, entire cells or well-designed mimics may serve as efficient catalysts in energy storage applications such as fuel cells or electrolysers for water splitting. Furthermore, a functional coupling to sustainable energy supplies such as photovoltaics appears to be a promising approach. With these ideas in mind, we believe that hydrogenase research is and will be a vivid field of enzymatic research that is now on the verge of advancing to the exploration of potential technological applications. At the same time smart ideas are demanded to advance to the level of understanding further to a yet unprecedent detail. We hope that this review gives the interested reader an overview on this topic and allows him to find hitherto unresolved research questions to fuel this interesting research with novel ideas worth pursuing

List of abbreviations

in the near future.

Review Article

Apical			
Basal			
Bovine serum albumin			
Electron nuclear double resonance			
Anodic peak potential			
Cathodic peak potential			
Electron paramagnetic resonance			
Distal (or rotated) iron atom in $[2Fe]_H$ referred to $[4Fe]_H$			
Proximal (or unrotated) iron atom [2Fe] _H referred to [4Fe] _H			
Fluorine doped tin oxide			
Hydrogen evolution reaction			
Hyperfine coupling			
Hyperfine sublevel correlation			
infrared			
Metal-to-ligand charge transfer			
Nuclear resonance vibrational spectroscopy			
Normal hydrogen electrode			
Open circuit potential			
Proton-coupled electron transfer			
Photon-driven electron transfer			
Photosensitizer			
Proton transfer pathway			
Quantum dots			
Room temperature			

SDM	Site directed mutagenesis
SHE	Standard hydrogen electrode
SEC	Spectroelectrochemistry
TOF	Turnover frequency
TON	Turnover number
WT	Wild type
SZATE	V man abasemtian and anticalan

XAE X-ray absorption and emission

XRD X-ray diffraction

Proteins

HydG	Maturase protein
HydE	Maturase protein
HydF	Maturase protein

apo-HydA Hydrogenase lacking [2Fe]_H
HydA Maturated hydrogenase
HydS Sensory hydrogenase

CrHydA1(XDT) [FeFe]-hydrogenase from C. reinhardtii with

an

xdt bridged [2Fe]_H

CpI Hydrogenase I from C. pasteurianum
CpII Hydrogenase II from C. pasteurianum
DdH Hydrogenase from D. desulfuricans

Compounds and groups

bda	benzylideneacetone/	(E)	4-phen	ylbut-3-ene-2-

one

BIAN-R Bis(arylimino)acenaphthene

Bn Benzyl

DCM Dichloromethane

DDQ 2,3-Dichloro-5,6-dicyano-1,4-benzoquinone

DHG Dehydroglycine
DMF Dimethylformamide
DMPC 1,2-Dimyristoyl-*sn-glycero*-3-

phosphatidylcholine

DOPC 1,2-Dioleoyl-sn-glycero-3-

phosphatidylcholine

dtc⁻ Diethyldithiocarbamate

Fc Ferrocene

Fc* Decamethylferrocene

Fmoc Fluorenylmethoxycarbonyl protecting group

hmds Hexamethyldisilazide

HOAc Acetic acid

HOTf Trifluoromethanesulfonic acid

HOTs *p*-Toluenesulfonic acid

 $\begin{array}{ll} HOB & Hydroxybenzyl \\ H_4 folate & Tetrahydrofolate \end{array}$

MAA 2-Mercaptobenzoic acid

MeCN Acetonitrile

MOF metal-organic framework
MPA 3-Mercaptopropionic acid

NaDT Sodium dithionite

NB Nitrobindin

NMI Naphthalene monoimide

OVA Ovalbumin
PAA Polyacrylic acid
PLP Pyridoxal phosphate

Chem Soc Rev **Review Article**

POEGMA Poly(oligo(ethyleneglycol)methyl ether methacrylate) [2Fe] _H [2Fe-2S]-cluster in the active site hydrogenases ppy 2-Phenylpyridine [4Fe] _H [4Fe-4S]-cluster in the active site	
ppy 2-Phenylpyridine [4Fe] _H [4Fe-4S]-cluster in the active site	of [FeFe]-
	of [FeFe]-
py Pyridyl hydrogenases	[]
SAM S-Adenosyl-L-methionine H-cluster Active site of [FeFe]-hydrogenases	
SDS Sodium dodecyl sulphate	
TEA Triethylamine Ligands	
TEOA Triethanolamine bma 2,3-Bis(diphenylphosphino)maleic	
TFA Trifluoroacetic acid anhydride	
THF Tetrahydrofuran bpcd 4,5-Bis(diphenylphosphino)-4-cycle	nnenten-
TMP Tris(morpholino)phosphine 1,3-dione	penten
tol Methylphenyl bpy 2,2'-Bipyridine	
5'-DA• 5'-Deoxyadenosyl radical Cp Cyclopentadienyl	
5'-DAH 5'-Deoxyadenosine dcpm Bis(dicyclohexylphosphino)methan	ne
[BAr ^F] tetrakis(3,5- dmpe 1,2-Bis(dimethylphosphino)ethane	
bis(trifluoromethyl)phenyl)borane or dppb 1,4-Bis(diphenylphosphino)butane	
tetra(pentafluorophenyl)borane dppe 1,2-Bis(diphenylphosphino)ethane	
dppf 1,1'-Bis(diphenylphosphino)ferroce	
	ciic
dppn 18-Ris(diphenylphosphino)panhth	nalene
adse Bis(seienidometnyi)amine dnpp 1.2 Pic(diphenylphocphino)propag	
adt Bis(suindometnyi)amine dnny (7)-1 2-Ris(diphenylphosphino)ethe	
adt Borane Dis(suindometnyi)amine IMe 1 3-Dimethylimidagol-2-ylidana	ciic
aut /v-Benzylbis(sumdometnyr)amine // / / / / / / / / / / / / / / / / /	ne
adt N-Metnylbis(suindometnyl)amine IMes 1 3-Dimesitylimidazol-2-ylidene	iic
adt Borane N-methylbis(suindomethyl)amine Me dppm 1 1-Ris(diphenylphosphino)-1-meth	hylethane
adt - Propylbis(suindometnyi)amine NHC N heterogyelic carbene	lyiculanc
adt ^{SMe} N-(Methylthio)ethylbis(sulfidomethyl)amine NMI Naphthalene monoimide	
Dat 1,2-Benzeneditniolate DND W.L.Phoenhingamine ligands	
3-Methyl-1,2-benzenedithiolate	Δ.
bdt ^{4Me} 4-Methyl-1,2-benzenedithiolate terpy 2,6-Bis(2-pyridyl)pyridine	
Cl ₂ Ddt 3,6-Dichloro-1,2-benzenedithiolate TDD Tetraphenylporphyrin	
codt 3-Carboxybenzene-1,2-diffilolate triphos Phenyl	
dcbdt 1,4-Dicarboxybenzene-2,3-dithiolate his(diphenylphosphinoethyl)phosp	nhine
edt 1,2-Ethanedithiolate	, iiiiic
pdt ^{Et} 2,2-Diethyl-1,3-propanedithiolate Amino acids	
pdt ^{Me} 2,2-Dimethyl-1,3-propanedithiolate Ala (A) Alanine	
odt Dis(sulfidomethyl)ether	
odt Bis(sulfidomethyl)ether $\Delta ror(R)$ $\Delta rorinine$	
o-yyldt 1 2-Ris(sulfidomethyl)henzene Arg (R) Arginine	
o-xyldt 1,2-Bis(sulfidomethyl)benzene Asp (D) Aspartic acid PdSe 1,3-Propanediselenolate Cvs (C) Cysteine	
o-xyldt 1,2-Bis(sulfidomethyl)benzene Arg (R) Arginine pdSe 1,3-Propanediselenolate Cys (C) Cysteine pdt 1,3-Propanedithiolate Cys (C) Cysteine	
o-xyldt 1,2-Bis(sulfidomethyl)benzene Asp (R) Arginine pdSe 1,3-Propanediselenolate Asp (D) Aspartic acid pdt 1,3-Propanedithiolate Cys (C) Cysteine pdtMeSBn 2-Methyl-2-((benzylthio)methyl)propane-1 3- pdtMeSBn 2-Methyl-2-((benzylthio)methyl)propane-1 3-	
o-xyldt 1,2-Bis(sulfidomethyl)benzene pdSe 1,3-Propanediselenolate pdt 1,3-Propanedithiolate pdt 2-Methyl-2-((benzylthio)methyl)propane-1,3- dithiolate Asp (R) Arginine Asp (R) Arginine Asp (D) Aspartic acid Cys (C) Cysteine EPA 3-(Diethylphosphorothioyl)alanine Glutamine	
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o-xyldt 1,2-Bis(sulfidomethyl)benzene pdSe 1,3-Propanediselenolate pdt 1,3-Propanedithiolate pdt ^{MeSBn} 2-Methyl-2-((benzylthio)methyl)propane-1,3- dithiolate pdt ^{MeSMe} 2-Methyl-2-((methylthio)methyl)propane-1,3- dithiolate pdt Bis(sulfidomethyl)sulfide sdt Bis(sulfidomethyl)sulfide Asp (D) Aspartic acid Cys (C) Cysteine EPA 3-(Diethylphosphorothioyl)alanine Glu (E) Glu (E) Glutamic acid Gly (G) Glycine His (H) HistidineI	
o-xyldt 1,2-Bis(sulfidomethyl)benzene pdSe 1,3-Propanediselenolate pdt 1,3-Propanedithiolate pdt ^{MeSBn} 2-Methyl-2-((benzylthio)methyl)propane-1,3- dithiolate pdt ^{MeSMe} 2-Methyl-2-((methylthio)methyl)propane-1,3- dithiolate pdt Bis(sulfidomethyl)sulfide sdt Bis(sulfidomethyl)sulfide pdt 1,3-Propanedithiolate pdt ^{MeSBn} 2-Methyl-2-((methylthio)methyl)propane-1,3- dithiolate pdt Bis(sulfidomethyl)sulfide sdt Bis(sulfidomethyl)sulfide pdt 1,3-Propanediselenolate Cys (C) Cysteine 3-(Diethylphosphorothioyl)alanine Gln (Q) Glutamine Glu (E) Gly (G) Glycine His (H) HistidineI pdt 3-(Diisopropylphosphorothioyl)alanine sdt Dithiolate ligand	
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o-xyldt 1,2-Bis(sulfidomethyl)benzene Asp (D) Aspartic acid pdSe 1,3-Propanediselenolate Cys (C) Cysteine pdt 1,3-Propanedithiolate EPA 3-(Diethylphosphorothioyl)alanine pdtMeSBn 2-Methyl-2-((benzylthio)methyl)propane-1,3- dithiolate Glu (E) Glutamine Glu (E) pdtMeSMe 2-Methyl-2-((methylthio)methyl)propane-1,3- dithiolate Gly (G) Glycine sdt Bis(sulfidomethyl)sulfide His (H) HistidineI xdt Dithiolate ligand Leu (L) Leucine Met (M) Methionine Iron-sulfur clusters PPA 3-(Diphenylphosphorothioyl)alanine Asp (R) Argmine Asp (R) Aspartic acid Cys (C) Cysteine 3-(Diethylphosphorothioyl)alanine Glu (E) Glutamine Gly (G) Glycine His (H) HistidineI Iron-sulfur clusters PPA 3-(Diphenylphosphorothioyl)alanine Asp (D) Aspartic acid Asp (D) Cysteine BPA 3-(Diethylphosphorothioyl)alanine Asp (D) Aspartic acid Asp (D) Cysteine BPA 3-(Diethylphosphorothioyl)alanine Asp (D) Aspartic acid	nine
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D. desulfuricans Desulfovibrio desulfuricans (Gram-negative)

E. coli Escherichia coli (Gram-negative)

T. maritima Thermotoga maritima (Gram-negative)

S. obliquus Scenedesmus obliquus

Conflicts of interest

There are no conflicts to declare.

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