

# Spectroscopic and Computational Evidence that [FeFe] Hydrogenases Operate Exclusively with CO-Bridged Intermediates

James A. Birrell,<sup>\*,†,¶</sup> Vladimir Pelmeshnikov,<sup>§,¶</sup> Nakul Mishra,<sup>‡</sup> Hongxin Wang,<sup>‡</sup> Yoshitaka Yoda,<sup>||</sup> Kenji Tamasaku,<sup>||</sup> Thomas B. Rauchfuss,<sup>#</sup> Stephen P. Cramer,<sup>⊥</sup> Wolfgang Lubitz,<sup>†</sup> and Serena DeBeer<sup>†</sup>

<sup>†</sup>Max Planck Institute for Chemical Energy Conversion, Stiftstrasse 34-36, 45470 Mülheim an der Ruhr, Germany

<sup>§</sup>Institut für Chemie, Technische Universität Berlin, Strasse des 17 Juni 135, 10623 Berlin, Germany

<sup>‡</sup>Department of Chemistry, University of California, Davis, One Shields Avenue, Davis, California 95616, United States

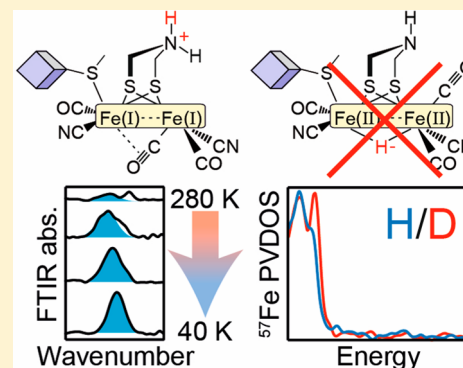
<sup>||</sup>JASRI Spring-8, 1-1-1 Kouto, Mikazuki-cho, Sayo-gun, Hyogo 679-5198, Japan

<sup>#</sup>School of Chemical Sciences, University of Illinois, 600 S. Mathews Avenue, Urbana, Illinois 61801, United States

<sup>⊥</sup>SETI Institute, Mountain View, California 94043, United States

## Supporting Information

**ABSTRACT:** [FeFe] hydrogenases are extremely active H<sub>2</sub>-converting enzymes. Their mechanism remains highly controversial, in particular, the nature of the one-electron and two-electron reduced intermediates called H<sub>red</sub>H<sup>+</sup> and H<sub>sred</sub>H<sup>+</sup>. In one model, the H<sub>red</sub>H<sup>+</sup> and H<sub>sred</sub>H<sup>+</sup> states contain a semibridging CO, while in the other model, the bridging CO is replaced by a bridging hydride. Using low-temperature IR spectroscopy and nuclear resonance vibrational spectroscopy, together with density functional theory calculations, we show that the bridging CO is retained in the H<sub>sred</sub>H<sup>+</sup> and H<sub>red</sub>H<sup>+</sup> states in the [FeFe] hydrogenases from *Chlamydomonas reinhardtii* and *Desulfovibrio desulfuricans*, respectively. Furthermore, there is no evidence for a bridging hydride in either state. These results agree with a model of the catalytic cycle in which the H<sub>red</sub>H<sup>+</sup> and H<sub>sred</sub>H<sup>+</sup> states are integral, catalytically competent components. We conclude that proton-coupled electron transfer between the two subclusters is crucial to catalysis and allows these enzymes to operate in a highly efficient and reversible manner.



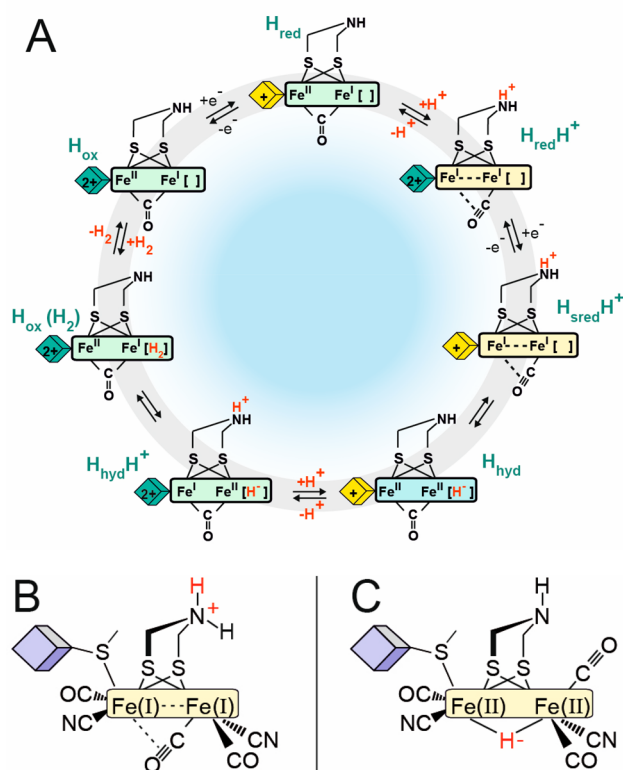
## INTRODUCTION

[FeFe] hydrogenases are highly active and efficient H<sub>2</sub> conversion catalysts with turnover frequencies up to 10 000 s<sup>-1</sup> for H<sub>2</sub> production.<sup>1,2</sup> The active site cofactor, the H-cluster, is composed of a unique [2Fe] subcluster ([2Fe]<sub>H</sub>) linked through a protein cysteine thiolate to a canonical [4Fe-4S] cluster ([4Fe-4S]<sub>H</sub>).<sup>3,4</sup> The two iron ions are connected by a bridging CO and a 2-azapropane 1,3-dithiolate (ADT) ligand.<sup>5,6</sup> Additionally, both irons are coordinated by terminal carbon monoxide (CO) and cyanide (CN<sup>-</sup>) ligands. The proximal iron (Fe<sub>p</sub>), vicinal to [4Fe-4S]<sub>H</sub>, is coordinatively saturated, while the distal iron (Fe<sub>d</sub>) features an open coordination site where substrates (H<sub>2</sub> and/or H<sup>+</sup>) and inhibitors (such as CO and O<sub>2</sub>) bind.<sup>7,8</sup> Some [FeFe] hydrogenases, including HydA1 from *Chlamydomonas reinhardtii* (CrHydA1), only contain the H-cluster.<sup>9,10</sup> Other enzymes, including HydAB from *Desulfovibrio desulfuricans* (DdHydAB) and HydA1 from *Clostridium acetobutylicum* (CaHydA1 or CaI), contain additional iron–sulfur clusters (F-clusters) for exchanging electrons with redox partner proteins.<sup>11,12</sup>

The catalytic cycle of [FeFe] hydrogenases (Figure 1A) remains controversial in spite of intensive efforts by many research groups.<sup>13–15</sup> Proton-coupled electron transfer (PCET) between the two subclusters of the H-cluster is believed to be essential for efficient and reversible catalysis.<sup>15,16</sup> It has been demonstrated with pH-dependent FTIR spectroelectrochemistry that, for CrHydA1, two forms of the one-electron reduced state exist where the reducing equivalent is localized on either [4Fe-4S]<sub>H</sub> or [2Fe]<sub>H</sub> (H<sub>red</sub> and H<sub>sred</sub>H<sup>+</sup>, respectively).<sup>15</sup> Electron transfer from [4Fe-4S]<sub>H</sub> to [2Fe]<sub>H</sub> is coupled to protonation of the bridgehead amine of the ADT ligand. This process was also shown to occur in DdHydAB, where it is enhanced by redox anticooperativity between [4Fe-4S]<sub>H</sub> and the F-cluster proximal to it.<sup>17</sup> PCET between the two subclusters is also thought to play a role in formation of H<sub>2</sub> from a terminal hydride-bound state, H<sub>hyd</sub>,<sup>16,18</sup> and activation of the oxygen-stable inactive state H<sub>inact</sub>.<sup>19</sup> PCET is disrupted in a sensory [FeFe] hydrogenase HydS from

Received: September 9, 2019

Published: December 10, 2019



**Figure 1.** (A) Putative catalytic mechanism for [FeFe] hydrogenase. The box represents the two Fe ions of the  $[2\text{Fe}]_{\text{H}}$  subsite, while the cube represents the  $[4\text{Fe-4S}]_{\text{H}}$  subsite. (B) The  $[2\text{Fe}]_{\text{H}}$  subcluster is in a homovalent  $\text{Fe(I)Fe(I)}$  configuration with the bridging CO shifted to a semibridging position on  $\text{Fe}_d$ , and the ADT ligand in the protonated  $-\text{NH}_2^+$  state, consistent with  $\text{H}_{(\text{s})\text{red}}\text{H}^+$ . (C) The  $[2\text{Fe}]_{\text{H}}$  subcluster is in a homovalent  $\text{Fe(II)Fe(II)}$  configuration with one of the CO ligands rotated into an apical position on  $\text{Fe}_d$  and a hydride in the bridging position. In the  $\text{H}_{\text{red}}\text{H}^+$  state  $[4\text{Fe-4S}]_{\text{H}}$  is oxidized to a 2+ state (green), and in  $\text{H}_{\text{sred}}\text{H}^+$  it is reduced to a 1+ state (yellow).

*Thermotoga maritima*<sup>20</sup> (*TmHydS*) or by an exchange of one of the cysteines ligating  $[4\text{Fe-4S}]_{\text{H}}$  with histidine in *CrHydA1*.<sup>21</sup>

An intriguing observation is that FTIR spectra of both the  $\text{H}_{\text{red}}\text{H}^+$  state and the two-electron reduced state  $\text{H}_{\text{sred}}\text{H}^+$  lack a peak corresponding to the bridging CO ligand.<sup>22–24</sup> In agreement with this, the crystal structure of *DdHydAB* under reducing conditions revealed a shift in the bridging CO ligand to a semibridging position on  $\text{Fe}_d$  ( $\text{Fe}_d-\mu\text{C}$  distance of 1.69 Å compared with the  $\text{Fe}_p-\mu\text{C}$  distance of 2.4 Å, shown schematically in Figure 1B).<sup>5</sup>

In the past few years, a number of observations have suggested an alternative structure for both the  $\text{H}_{\text{sred}}\text{H}^+$  and the  $\text{H}_{\text{red}}\text{H}^+$  states. In 2014, Leg er and co-workers demonstrated that *CrHydA1* undergoes reversible low potential inactivation<sup>25</sup> and attributed this conversion to the  $\text{H}_{\text{sred}}\text{H}^+$  state. However, the time-scale required for the inactivation is on the order of minutes, while the  $\text{H}_{\text{sred}}\text{H}^+$  state is routinely observed within seconds after reducing the enzyme.<sup>15,26</sup> In the same year, Haumann and co-workers suggested that the  $\text{H}_{\text{sred}}\text{H}^+$  (*sred* in their nomenclature, see supplementary discussion in the Supporting Information) state is best described as a bridging hydride based on site-selective X-ray absorption spectroscopy (XAS) and density functional theory (DFT) calculations (Figure 1C).<sup>27</sup> Subsequently, the same group suggested that the  $\text{H}_{\text{red}}\text{H}^+$  (*Hred* in their nomenclature) state

contains a bridging hydride based on DFT simulations of FTIR<sup>26</sup> and nuclear resonance vibrational spectra (NRVS).<sup>28</sup> In H-cluster model systems bridging hydrides are more stable than isomeric terminal hydrides.<sup>29–32</sup> This, and the fact that the former would not be adjacent to the proton relaying ADT, implies that bridging hydrides would be catalytically inactive.<sup>33</sup> As such, they have been proposed to serve a regulatory role.<sup>13,28</sup>

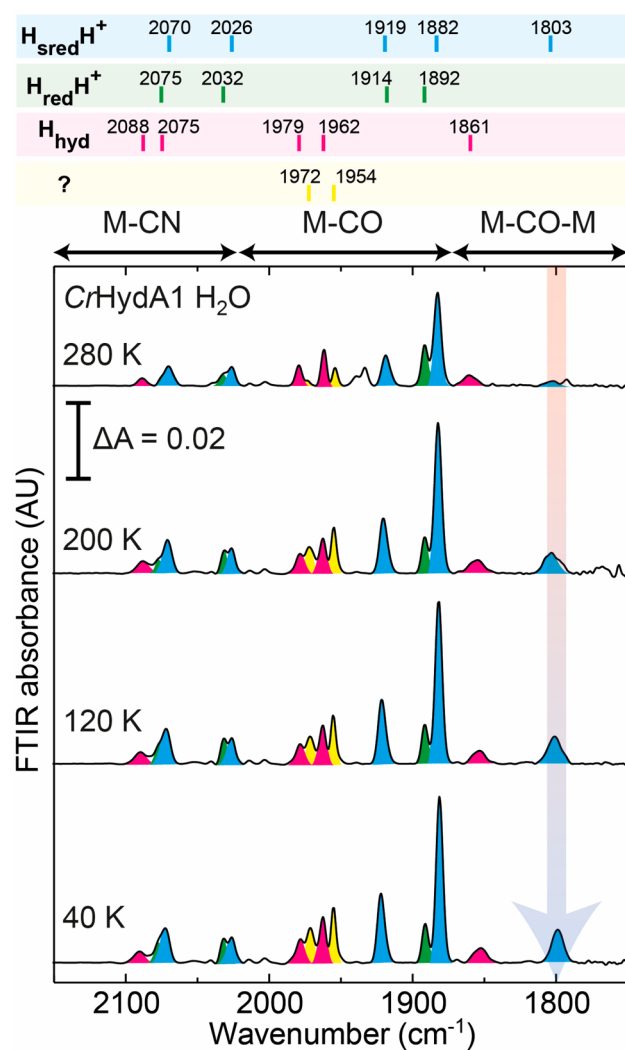
Despite the evidence in support of the bridging hydride structure of the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states and their inactive nature,<sup>13,27,28</sup> a few recent observations contradict this assignment. *TmHydS* can be poised in redox states with very similar FTIR spectra to the  $\text{H}_{\text{sred}}\text{H}^+$  and  $\text{H}_{\text{red}}\text{H}^+$  states but retaining the bridging CO.<sup>20</sup> The same appears to be true for the highly active, reversible [FeFe] hydrogenase, *CaI*.<sup>14</sup> Finally, a cysteine to histidine modification at  $[4\text{Fe-4S}]_{\text{H}}$  in *CrHydA1* increases the redox potential of  $[4\text{Fe-4S}]_{\text{H}}$ , resulting in an enzyme that does not stably form the  $\text{H}_{\text{red}}\text{H}^+$  state.<sup>21</sup> If the  $\text{H}_{\text{red}}\text{H}^+$  state is inactive, as has been proposed,<sup>13</sup> then the C362H variant of *CrHydA1* should show higher activity for  $\text{H}_2$  production than the wild-type enzyme. However, the C362H variant shows essentially no  $\text{H}_2$  production activity.

Clearly, new spectroscopic data are urgently required to resolve the involvement of the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states. To this end, this paper described an investigation of the nature of the bridging ligand, specifically whether a (semi)bridging CO or a bridging  $\text{H}^-$  ligand is present in the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states. Our studies focused on the best characterized [FeFe] hydrogenases, *CrHydA1* and *DdHydAB*. The former enzyme can be easily poised in the  $\text{H}_{\text{sred}}\text{H}^+$  state,<sup>34</sup> while the latter can be poised in the  $\text{H}_{\text{red}}\text{H}^+$  state,<sup>22</sup> both with almost complete purity. Crucially, low-temperature IR measurements reveal that both of these states feature bridging CO ligands. This result rules out the presence of a bridging hydride. Samples prepared in  $\text{H}_2\text{O}$  and  $\text{D}_2\text{O}$  show only subtle ( $<2\text{ cm}^{-1}$ ) differences in the  $\text{CN}^-$  and CO peak positions. At the same time, NRVS studies on samples prepared in  $\text{H}_2\text{O}$  and  $\text{D}_2\text{O}$  show only very subtle isotope effects, with no evidence for a bridging hydride. DFT calculations give better agreement with a bridging CO structure with a protonated ADT ligand than a structure in which a hydride is bridged between the two Fe ions of the  $[2\text{Fe}]_{\text{H}}$  subcluster. These findings have important consequences for the mechanism of  $\text{H}_2$  conversion by [FeFe] hydrogenases. Specifically, the presence of a bridging CO in the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states favors their inclusion into the catalytic cycle as catalytically competent states.

## RESULTS AND DISCUSSION

In order to study the spectral properties of the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states in more detail, we chose to apply variable-temperature FTIR spectroscopy. Initial experiments focused on sodium dithionite-reduced samples of *CrHydA1* at pH 8, in which the  $\text{H}_{\text{sred}}\text{H}^+$  state is highly populated.

**Low-Temperature FTIR Reveals the Bridging CO in  $\text{H}_{\text{sred}}\text{H}^+$ .** Figure 2 shows FTIR spectra of *CrHydA1* reduced with 20 mM sodium dithionite ( $E \approx -600\text{ mV}$ ) taken at various temperatures (see Figure S1 for the full temperature range and Figure S2 for data before spline curve fitting to remove additional background). At 280 K it can be seen that *CrHydA1* is predominantly in the  $\text{H}_{\text{sred}}\text{H}^+$  state with terminal CO bands at 1882 and 1919  $\text{cm}^{-1}$  and terminal  $\text{CN}^-$  bands at 2026 and 2070  $\text{cm}^{-1}$ . Historically, an additional terminal CO band has been identified at 1954  $\text{cm}^{-1}$ <sup>11,23,34,35</sup> and assumed to



**Figure 2.** Variable-temperature FTIR spectra of *CrHydA1* in  $H_2O$  buffer (50 mM MES, 50 mM HEPES, 300 mM KCl, pH 8) reduced with 20 mM sodium dithionite. Peaks corresponding to the  $H_{\text{sred}}H^+$  state are colored blue (60%). Smaller contributions from the  $H_{\text{red}}H^+$  state (25%), the  $H_{\text{hyd}}$  state (10%), and an uncharacterized state similar to  $H_{\text{hyd}}$  (5%) are colored green, pink, and yellow, respectively.

be the fate of the missing bridging CO peak. However, the intensity of this peak varies dramatically between different reports and is unlikely to be associated with the  $H_{\text{sred}}H^+$  state. Indeed, at 280 K we observed a low-intensity peak at 1954  $cm^{-1}$  (colored yellow).

In addition to the peaks from the  $H_{\text{sred}}H^+$  state, we observe the most intense terminal CO peak from the  $H_{\text{red}}H^+$  state at 1892  $cm^{-1}$  and some evidence for the  $CN^-$  bands of the same state as shoulders on the high-energy side of the  $H_{\text{sred}}H^+$   $CN^-$  peaks. Very minor contributions from the  $H_{\text{ox}}$  and  $H_{\text{red}}$  states can be identified from peaks at 1940 and 1934  $cm^{-1}$ , and a clear contribution from the  $H_{\text{hyd}}$  state can be observed as the pink peaks at 1861 (bridging CO) and 1962 and 1979  $cm^{-1}$  (terminal CO). The highest energy  $CN^-$  band from the  $H_{\text{hyd}}$  state is visible at 2088  $cm^{-1}$ , but the low-energy  $CN^-$  band from  $H_{\text{hyd}}$  and the high-energy  $CN^-$  bands from  $H_{\text{red}}H^+$  and  $H_{\text{sred}}H^+$  are overlapping in the 2070–2075  $cm^{-1}$  region.

As the temperature is lowered, the majority of the peaks observed do not significantly change (small shifts in peak position and line width are shown in Table S1). However, one

clear phenomenon can be observed: the appearance of a band at 1803  $cm^{-1}$ . We attribute this band to the bridging CO of the  $H_{\text{sred}}H^+$  state, the dominant state observed at all temperatures. This assignment is supported by its similarity to a band of the  $H_{\text{red}}H^+$  state observed for *CaI*.<sup>14</sup> It is also clear that at lower temperature the peaks at 1972 and 1954  $cm^{-1}$  have gained intensity. These changes also coincide with small changes to the  $H_{\text{hyd}}$  bridging CO peak (in the 1840–1870  $cm^{-1}$  region, Figure S3). The peak is also clearly visible in spectra that have not yet been additionally background corrected with spline curve fitting (Figure S2).

The same experiment was performed in  $D_2O$  and gave very similar results (Figures S4–S6). However, in  $D_2O$  all the peaks of the  $H_{\text{sred}}H^+$  state are shifted to slightly lower energies by 1–2  $cm^{-1}$  (Figure S7), suggesting subtle changes to the electron density on the  $[2Fe]_H$  subcluster or changes due to the vibrational coupling. These may be due to deuteration of the bridging ADT ligand, changes in hydrogen bond strength to  $[2Fe]_{HV}$ , or general changes in protein structure upon  $D_2O$  exchange. The  $D_2O$  sample also displayed much lower levels of additional states, yet retained similar intensity of the peak at 1803  $cm^{-1}$ , further supporting that this peak originates from the  $H_{\text{sred}}H^+$  state. The lack of a 1954  $cm^{-1}$  peak in the  $D_2O$  spectra supports our suggestion that this peak does not originate from  $H_{\text{sred}}H^+$ .

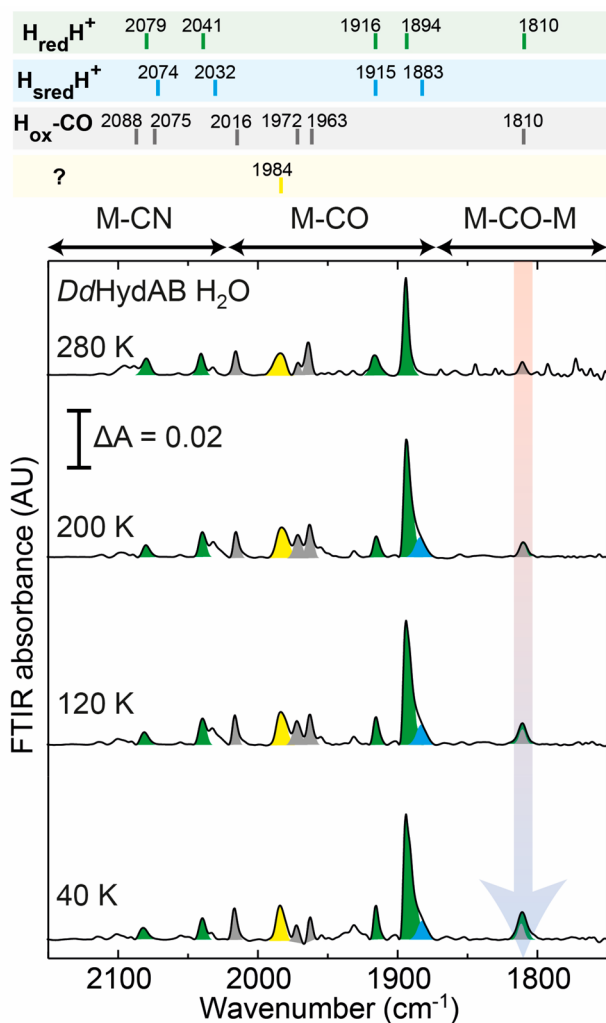
**Low-Temperature FTIR Reveals the Bridging CO in  $H_{\text{red}}H^+$ .** FTIR measurements on the  $H_{\text{red}}H^+$  state focused on the *DdHydAB* enzyme because *CrHydA1*-derived samples contain multiple states ( $H_{\text{ox}}$ ,  $H_{\text{sred}}H^+$ , and  $H_{\text{red}}$ ). Conveniently, the *DdHydAB* enzyme, when reduced with 20 mM sodium dithionite at pH 8, produced almost exclusively the  $H_{\text{red}}H^+$  state. This is because of redox-anticooperativity between the H-cluster and the proximal F-cluster, which lowers the intrinsic redox potential of  $[4Fe-4S]_H$ .<sup>17</sup> This effect destabilizes both  $H_{\text{red}}$  and  $H_{\text{sred}}H^+$  in *DdHydAB* and favors  $H_{\text{red}}H^+$ .

At room temperature, peaks can be observed at 1894, 1916, 2041, and 2079  $cm^{-1}$ , corresponding to the terminal CO and  $CN^-$  ligands (Figure 3; see also Figures S8 and S9). Small contributions can be observed from the  $H_{\text{ox}}$ -CO state (2088, 2075, 2016, 1972, 1963, and 1810  $cm^{-1}$ ) and some unknown states, including a large broad feature around 1984  $cm^{-1}$ . At lower temperatures the main change observed is the increase in intensity of the feature at 1810  $cm^{-1}$ , assigned to the bridging CO from the  $H_{\text{red}}H^+$  state, with some additional subtle changes to peak positions and line widths (Table S1). The peak occurs at 7  $cm^{-1}$  higher energy compared with the bridging CO in the  $H_{\text{sred}}H^+$  state of *CrHydA1*, similar to the other CO and  $CN^-$  peaks. The peak is, however, overlapping with the bridging CO from the  $H_{\text{ox}}$ -CO state, which is already an observable contribution at high temperature.

Samples prepared in  $D_2O$  behaved similarly to those in  $H_2O$  (Figures S10–S12), but with a smaller  $H_{\text{ox}}$ -CO contribution. However, the more intense bridging CO peak at 1810  $cm^{-1}$  further highlights that this peak originates from the  $H_{\text{red}}H^+$  state. Again, only small shifts in the peaks of the  $H_{\text{red}}H^+$  state were observed following  $D_2O$  exchange, this time to higher energy (Figure S13).

To summarize, both the  $H_{\text{sred}}H^+$  and  $H_{\text{red}}H^+$  states of the  $[FeFe]$  hydrogenase appear to contain a bridging CO ligand. Meanwhile, none of the other peaks undergo particularly large shifts concomitant with the appearance of the bridging CO peak, excluding the possibility that a large conformational change occurs upon cooling the sample. Even if this were the

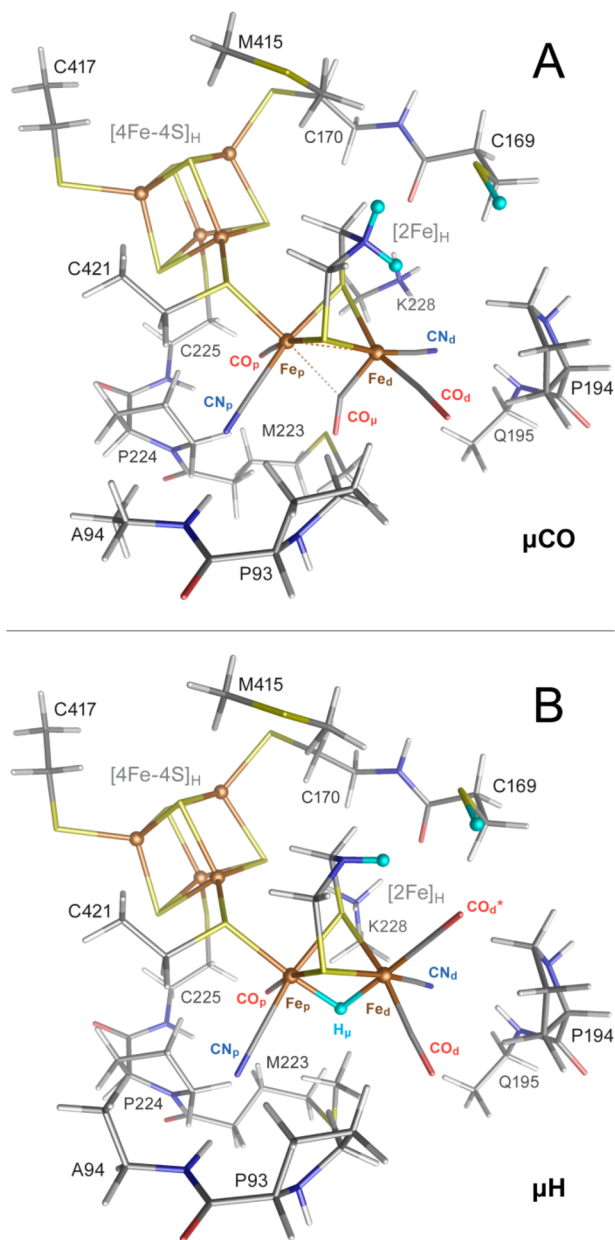




**Figure 3.** Variable-temperature FTIR spectra of *DdHydAB* in  $H_2O$  buffer (50 mM MES, 50 mM HEPES, 300 mM KCl, pH 8) reduced with 20 mM sodium dithionite. The peaks corresponding to the  $H_{\text{red}}H^+$  state are colored green. Small contributions from the  $H_{\text{sred}}H^+$  state, the  $H_{\text{ox}}\text{-CO}$  state, and an uncharacterized state are colored blue, gray, and yellow, respectively.

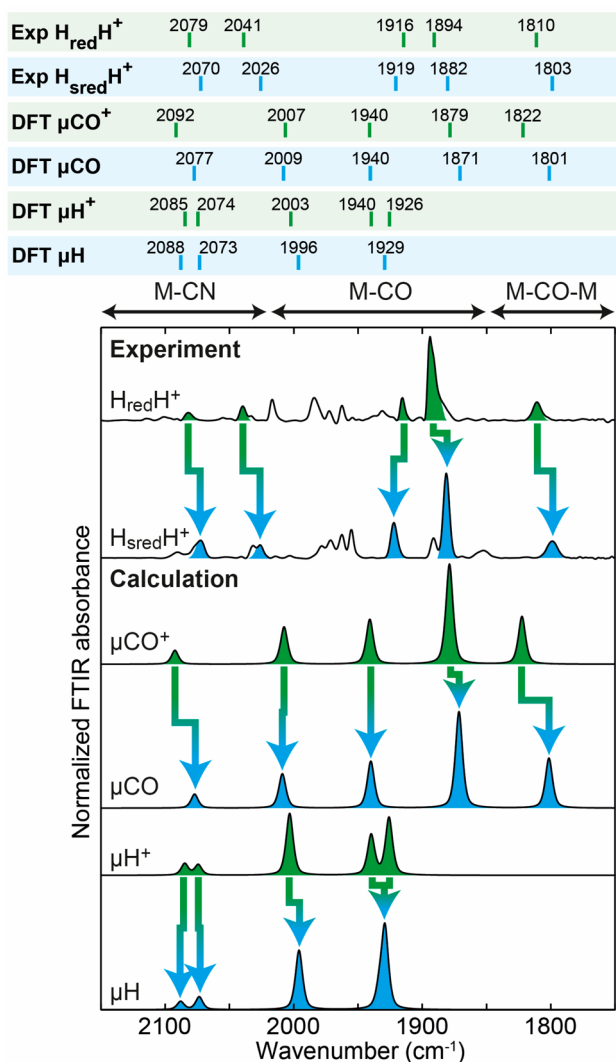
case, the previous XAS and NRVS studies, which were used to propose the bridging hydride, were also performed on frozen samples under cryogenic conditions and should, therefore, contain a bridging CO.<sup>27,28</sup> An IR H/D isotope effect, like that observed in the  $H_{\text{hyd}}$  state,<sup>16</sup> which is well known to contain a terminal Fe hydride,<sup>18,36–38</sup> could not be observed here. This further suggests that the  $H_{\text{red}}H^+$  and  $H_{\text{sred}}H^+$  states do not contain a bridging hydride.

**DFT Calculations Reproduce Experimental FTIR Spectra of  $H_{\text{red}}H^+$  and  $H_{\text{sred}}H^+$  Using a Bridging CO Model.** DFT calculations using bridging CO ( $\mu\text{CO}$  for  $H_{\text{sred}}H^+$  and  $\mu\text{CO}^+$  for  $H_{\text{red}}H^+$ ) and bridging hydride ( $\mu\text{H}$  for  $H_{\text{sred}}H^+$  and  $\mu\text{H}^+$  for  $H_{\text{red}}H^+$ ) models (Figure 4) were used to calculate FTIR spectra (Figure 5). These DFT models are compared in Figures S14–S17, with their additional characterization provided in Tables S2–S5. Notably, the models  $\mu\text{CO}$  and  $\mu\text{CO}^+$  produced vanishingly small spin populations at the two  $[2\text{Fe}]_{\text{H}}$  Fe(I) sites, indicating the Fe(I)–Fe(I) metal–metal bonding as depicted in Figure 1B. The calculated bands for  $H_{\text{sred}}H^+$  using the  $\mu\text{CO}$  model are at 2077 and 2009  $\text{cm}^{-1}$



**Figure 4.** Optimized structures from DFT calculations on the  $\mu\text{CO}$  (A) and  $\mu\text{H}$  (B) models shown in tube representation. Additionally, in ball representation are indicated the Fe sites and the three H-to-D exchangeable protons (light blue). Single-letter amino acid labeling corresponds to the *CrHydA1* enzyme sequence. Element colors are C (gray), H (white), N (blue), O (red), Fe (brown), and S (yellow). For extra details and DFT models of  $\mu\text{CO}^+$  and  $\mu\text{H}^+$ , see Figures S14–S17.

for the  $\text{CN}^-$  ligands, 1940 and 1871  $\text{cm}^{-1}$  for the terminal CO ligands, and 1801  $\text{cm}^{-1}$  for the bridging CO ligand. The 1871 and 1801  $\text{cm}^{-1}$  C–O modes display a minor degree of vibrational coupling (see Supporting Information for the normal mode animations). These values are quite close to the experimental values of 2070, 2026, 1919, 1882, and 1803  $\text{cm}^{-1}$ . While the C–O/N stretch vibrational energies of the terminal ligands assigned to  $\text{Fe}_p$  are somewhat overestimated, the energies of the terminal ligands assigned to  $\text{Fe}_d$  are underestimated (Table S4). Notably, the relative IR intensities of the observed bands were reproduced. Given the limitations of DFT, we find the agreement with the experiment to be very



**Figure 5.** Experimental and DFT-calculated IR spectra of the  $H_{\text{red}}H^+$  and  $H_{\text{sred}}H^+$  states in  $H_2O$ . The experimental spectra are the 40 K spectra from the  $H_{\text{red}}H^+$  state of *DdHydAB* (Figure 3) and *CrHydA1* in the  $H_{\text{sred}}H^+$  state (Figure 2). The calculated  $\mu\text{CO}$  and  $\mu\text{CO}^+$  spectra were generated from DFT calculations on a bridging CO model (Figure 4A). The  $\mu\text{H}$  and  $\mu\text{H}^+$  spectra were generated from DFT calculations on a bridging hydride model (Figure 4B). The peaks corresponding to the  $H_{\text{red}}H^+$  and  $H_{\text{sred}}H^+$  states are colored green and blue, respectively. See Table S4 for comparison of experimental and calculated IR frequencies and assignment to the ligands on the proximal and distal Fe.

good. Importantly, while the calculations predict that the CO ligand retains its bridging character in the  $H_{\text{sred}}H^+$  (and the  $H_{\text{red}}H^+$ ) model, its carbon shifts 0.1 Å closer to  $\text{Fe}_d$  and 0.2 Å away from  $\text{Fe}_p$  when compared to, for example, the  $H_{\text{hyd}}$  state, as detailed in Table S2 and schematized in Figure 1B.

Calculations on a bridging hydride variant ( $\mu\text{H}$ ) with two terminal CO ligands on  $\text{Fe}_d$  give radically different FTIR spectra with CN bands at 2088 and 2073  $\text{cm}^{-1}$  and terminal CO bands at 1996 and 1929  $\text{cm}^{-1}$ . The high values compared with the bridging CO model are due to the fact that hydride formation leaves the two Fe ions formally in Fe(II) oxidation states (Figure 1C). The prediction of just two CO bands is due to an overlay of the terminal (at  $\text{Fe}_p$  and apical at  $\text{Fe}_d$ ) C–O modes at  $\sim 1929 \text{ cm}^{-1}$ , which become significantly mixed in the absence of the bridging CO ligand. The calculated spectra

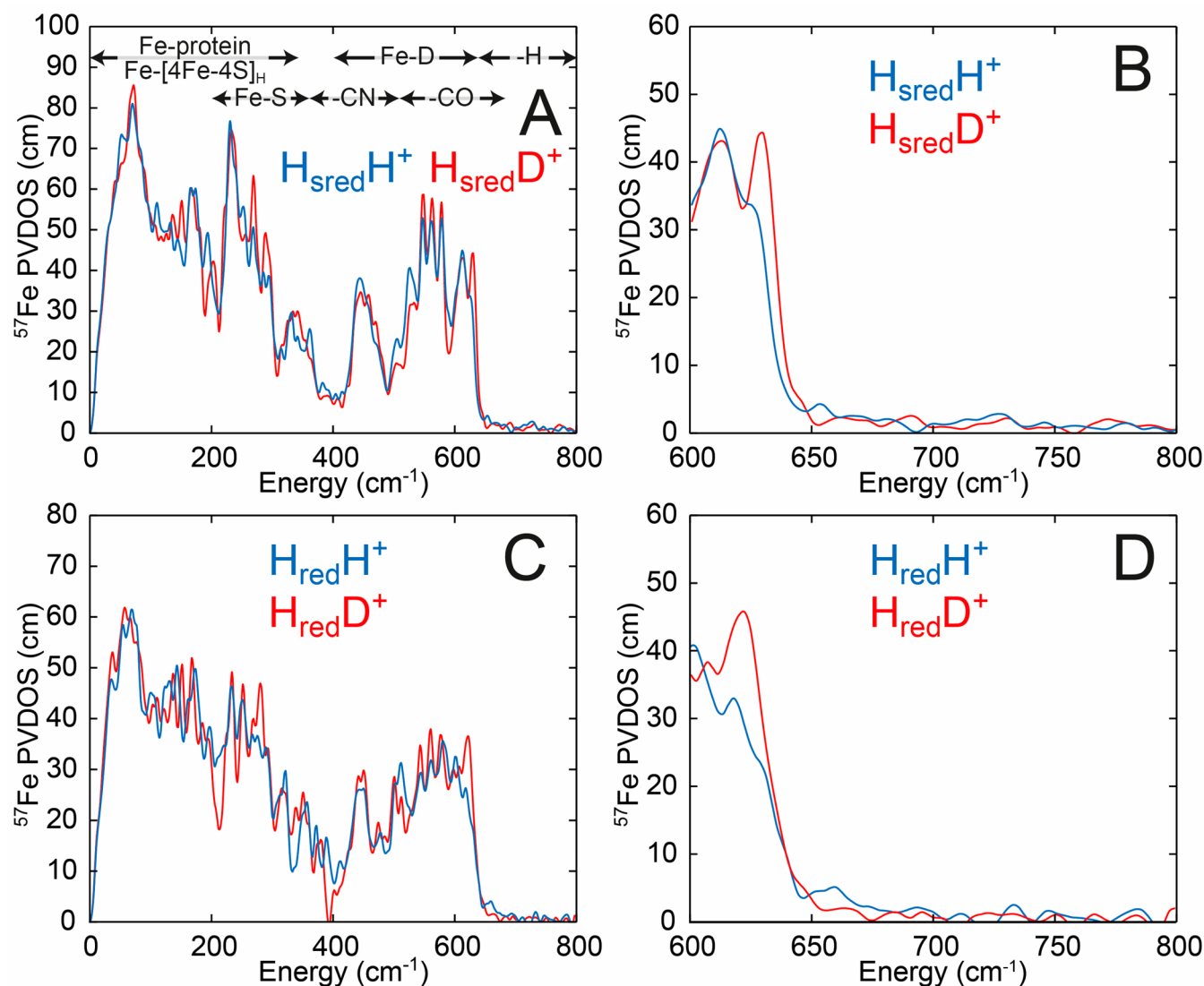
for the deuterated forms were only very subtly different from those of the protonated forms (Figure S18).

The calculated bands for  $H_{\text{red}}H^+$  using a bridging CO ( $\mu\text{CO}^+$ ) model are at 2092 and 2007  $\text{cm}^{-1}$  for the CN<sup>−</sup> ligands, 1940 and 1879  $\text{cm}^{-1}$  for the terminal CO ligands, and 1822  $\text{cm}^{-1}$  for the bridging CO ligand. These are quite similar to the calculated bands for  $H_{\text{sred}}H^+$  but shifted to higher energy by up to 21  $\text{cm}^{-1}$ . The band positions, relative intensities, and most of the shifts relative to  $H_{\text{sred}}H^+$  capture the trends observed in the experimental spectra. In both the  $\mu\text{H}^+$  vs  $\mu\text{H}$  and  $\mu\text{CO}^+$  vs  $\mu\text{CO}$  spectra comparisons, the IR band shifts are governed solely by the redox state of the  $[4\text{Fe-4S}]_{\text{H}}$  subcluster, while the  $[2\text{Fe}]_{\text{H}}$  subcluster harboring the CO/CN ligands remains in the same formal state. Our calculations indicate that, upon reduction of  $\mu\text{CO}^+$  to  $\mu\text{CO}$ , the  $[4\text{Fe-4S}]_{\text{H}}$  fragment charge shifts by  $\sim 0.9$  units, and  $[2\text{Fe}]_{\text{H}}$  obtains the remaining  $\sim 0.1$  units of the added electron density (Table S5). Calculations with a bridging hydride variant  $\mu\text{H}^+$  gave IR bands similar to those produced for the  $\mu\text{H}$  model. However, the vibrational energies of the terminal CO ligands are better separated in  $\mu\text{H}^+$  giving bands at 1940 and 1926  $\text{cm}^{-1}$ . Importantly, the bridging hydride models do not produce IR spectra remotely reminiscent of the experimental spectra.

The complete assignment of the bands for the  $H_{\text{red}}H^+$  and  $H_{\text{sred}}H^+$  states makes clear that the bridging CO does not move to a terminal apical position on  $\text{Fe}_d$  giving an FTIR peak around 1950–1960  $\text{cm}^{-1}$ . Therefore, the origin of the 1954  $\text{cm}^{-1}$  peak in the reduced *CrHydA1* and *DdHydAB* samples needs to be explained. We observed that the increase in intensity of the 1954  $\text{cm}^{-1}$  peak in reduced *CrHydA1* correlated with an increase in intensity of a peak at 1972  $\text{cm}^{-1}$  and changes to the bridging CO peak associated with the  $H_{\text{hyd}}$  state. Therefore, we believe these peaks arise from an alternative form of the  $H_{\text{hyd}}$  state. A thorough assignment will require obtaining this state in higher purity, probably by performing cryogenic FTIR experiments on *CrHydA1* samples under various reducing conditions. An important outcome of the present work is that samples measured at room temperature (e.g., with FTIR) cannot be directly compared with samples measured at cryogenic temperatures. A number of recent results will need to be revisited in light of this finding. Recently there have been numerous spectroscopic investigations into the  $H_{\text{hyd}}$  state, identifying it as a terminal hydride intermediate.<sup>16,18,36–39</sup> However, IR measurements have never been performed at similarly low temperatures to those used for EPR, Mössbauer, and NRVs spectroscopies and could provide further insight.

**NRVS and DFT Calculations Support a Bridging CO Structure for the H-Cluster in  $H_{\text{sred}}H^+$  and  $H_{\text{red}}H^+$ .** To test our assignment of the  $H_{\text{red}}H^+$  and  $H_{\text{sred}}H^+$  states, NRVs measurements were conducted on samples of *CrHydA1* and *DdHydAB*. Samples were poised under identical conditions as for the FTIR samples, transferred to NRVs cells, and frozen in liquid nitrogen. To ensure high sensitivity, the  $[2\text{Fe}]_{\text{H}}$  subsite was selectively labeled with  $^{57}\text{Fe}$ .

Figure 6 (and Figure S19) shows NRVs spectra for *CrHydA1* and *DdHydAB* in the  $H_{\text{sred}}H^+$  and  $H_{\text{red}}H^+$  states, respectively, under  $\text{H}_2/\text{H}_2\text{O}$  (blue traces) and  $\text{D}_2/\text{D}_2\text{O}$  (red traces). Spectra measured in  $\text{H}_2\text{O}$  and  $\text{D}_2\text{O}$  in both enzymes look very similar. Importantly, at high energy (600–800  $\text{cm}^{-1}$ ) no peaks can be observed in the  $\text{H}_2\text{O}$  samples. Previous studies on the terminal hydride  $H_{\text{hyd}}$  state of the  $[\text{FeFe}]$  hydrogenase<sup>18,37,38</sup> and the bridging hydride Ni-R state of the

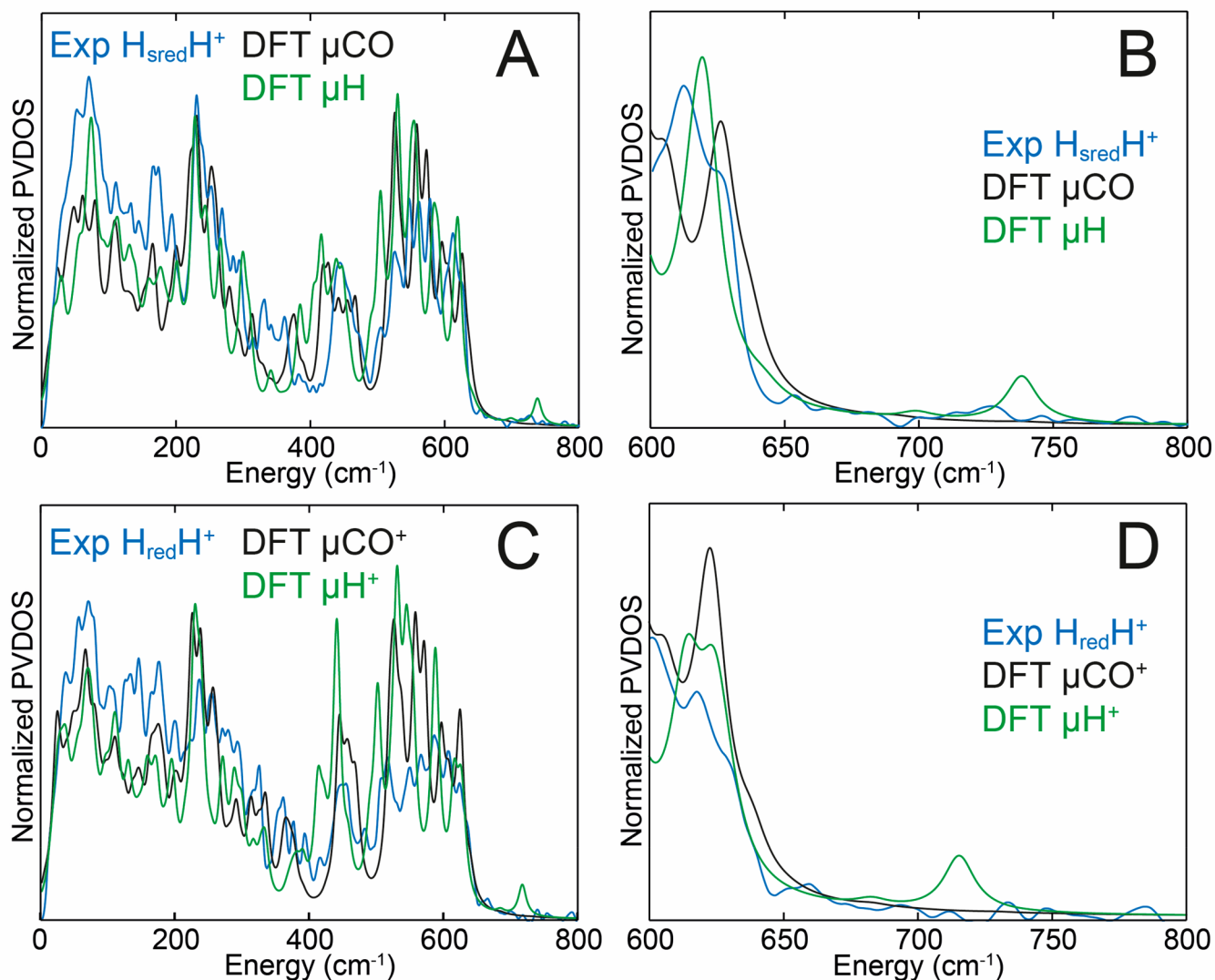


**Figure 6.** NRVs of CrHydA1 in the  $H_{\text{sred}}H^+$  state (A and B) in  $H_2O$  buffer (50 mM MES, 50 mM HEPES, 300 mM KCl, pH 8) (blue traces) and  $D_2O$  buffer (50 mM MES, 50 mM HEPES, 300 mM KCl, pD 8) (red traces) reduced with 20 mM sodium dithionite. NRVs of DdHydAB in the  $H_{\text{red}}H^+$  state (C and D) in  $H_2O$  buffer (50 mM MES, 50 mM HEPES, 300 mM KCl, pH 8) (blue traces) and  $D_2O$  buffer (50 mM MES, 50 mM HEPES, 300 mM KCl, pD 8) (red traces) reduced with 20 mM sodium dithionite. B and D are expansions of the high-energy region from A and C, respectively. Error bars have been omitted for clarity. The regions of the spectrum corresponding to Fe-protein, Fe-[4Fe-4S]<sub>H</sub>, Fe-S, Fe-CN, Fe-CO, Fe-D, and Fe-H are indicated in A. Data including error bars are presented in Figure S19.

[NiFe] hydrogenase,<sup>40</sup> as well as either terminal or bridging hydride containing diiron model complexes,<sup>41</sup> show distinct peaks in this region. For CrHydA1 with an oxadithiolate (ODT) bridge the peaks at 670 and 727  $\text{cm}^{-1}$  were assigned to bending motions of the terminal hydride correspondingly out-of-plane and in-plane, with respect to the pseudosymmetry plane of the  $[2Fe]_H$  subcluster, respectively.<sup>38</sup> CrHydA1 with the native ADT cofactor showed the same pair of peaks but shifted to higher energy (675 and 744  $\text{cm}^{-1}$ ), as did DdHydAB (675 and 747  $\text{cm}^{-1}$ , see Figure S20).<sup>18</sup> In the C169S variant of CrHydA1, the in-plane peak is shifted to even higher energy (772  $\text{cm}^{-1}$ ), while the out-of-plane peak is barely perturbed by the amino acid substitution (673  $\text{cm}^{-1}$ ).<sup>37</sup> For the [NiFe] hydrogenase in the Ni-R state a peak was observed at 675  $\text{cm}^{-1}$ ,<sup>40</sup> assigned to wagging motion of the bridging hydride. For a diiron model complex synthesized with a hydride in either terminal or bridging positions, Fe–H bending and Fe–H–Fe wagging NRVs intensities were observed in the 720–

800 and 670–700  $\text{cm}^{-1}$  regions.<sup>41</sup> Upon H-to-D isotope exchange, these previously identified iron-hydride vibrations show large (up to  $\sim 200 \text{ cm}^{-1}$ ) shifts to lower energies, where they additionally perturb high-intensity Fe–CO bands.<sup>42</sup> The fact that no such  $H_2O$  vs  $D_2O$  NRVs signatures are presently observed in either the  $H_{\text{sred}}H^+$  or  $H_{\text{red}}H^+$  samples directly contradicts the idea that Fe–H bonds exist in these states.

Figure 7A and B show an overlay of DFT calculations of the  $H_{\text{sred}}H^+$   $H_2O$  spectra generated using both a bridging CO model ( $\mu\text{CO}$ ) and a bridging hydride model ( $\mu\text{H}$ ). The experimental spectra are best reproduced by the  $\mu\text{CO}$  model. Notably, the  $\mu\text{H}$  model predicts an NRVs-detectable Fe–H–Fe mode at 738  $\text{cm}^{-1}$ , which is not experimentally observed. The calculations indicate that this peak is above the level of noise in the experimental data (Figure S21). H-to-D isotopically labeled (as specified in Figure 4) models  $\mu\text{CO}^D$  and  $\mu\text{D}$  were used to calculate the  $D_2O$  sample spectra, and again the  $\mu\text{CO}^D$  model gave better agreement to the



**Figure 7.** DFT calculations of the NRVS spectra in the  $H_{\text{sred}}H^+$  state (A and B) and  $H_{\text{red}}H^+$  state (C and D). The experimental NRVS spectra of CrHydA1 in the  $H_{\text{sred}}H^+$  state (A and B) and DdHydAB in the  $H_{\text{red}}H^+$  state (C and D) are shown in blue, while the DFT-calculated spectra generated using bridging CO ( $\mu\text{CO}$  and  $\mu\text{CO}^+$ ) models (Figure 4A) are shown in black, and the DFT-calculated spectra generated using bridging hydride ( $\mu\text{H}$  and  $\mu\text{H}^+$ ) models (Figure 4B) are shown in green. B and D are expansions of the high-energy regions of A and C. Experimental error bars have been omitted for clarity. Data including error bars are presented in Figures S21 and S23. See Figure S20 for an alternative comparison of the DFT-calculated and NRVS-observed  $^{57}\text{Fe}$ -PVDOS spectra.

experimental data. The  $\mu\text{CO}/\mu\text{CO}^{\text{D}}$  models suggest only very minor changes upon H/D exchange (Figure S22A), while the  $\mu\text{H}/\text{D}$  models suggest much larger changes (Figure S22B), particularly in the Fe-CO region (400–650  $\text{cm}^{-1}$ ), which are not experimentally observed. Likewise, DFT calculations on  $\mu\text{CO}^+$  and  $\mu\text{H}^+$  (now with an oxidized  $[4\text{Fe-4S}]_{\text{H}}^{2+}$  subcluster) models were used to calculate the spectra of  $H_{\text{red}}H^+$  (Figure 7C and D and Figure S23). The  $\mu\text{CO}^+$  model provided better agreement with experimental data than the  $\mu\text{H}^+$  model. The  $\mu\text{H}^+$  model predicts an NRVS-detectable Fe–H–Fe mode now at 716  $\text{cm}^{-1}$ ; however, a corresponding band is not produced by the  $H_{\text{red}}H^+$  sample. Additionally, the  $\mu\text{CO}^+/\mu\text{CO}^{\text{D}+}$  models, similarly to  $\mu\text{CO}/\mu\text{CO}^{\text{D}}$ , suggest only small changes upon H/D exchange (Figure S24A), while the  $\mu\text{H}^+/\mu\text{D}^+$  models suggest larger changes (Figure S24B). Notably, the bridging hydride models, formally different only in the  $[4\text{Fe-4S}]_{\text{H}}$  oxidation level ( $[4\text{Fe-4S}]_{\text{H}}^{2+}$  in  $\mu\text{H}^+$  vs  $[4\text{Fe-4S}]_{\text{H}}^{1+}$  in  $\mu\text{H}$ ), display a substantial 22  $\text{cm}^{-1}$  deviation in the Fe–H–

Fe wagging mode position. This highlights the sensitivity of the iron-hydride bands in  $[\text{FeFe}]$  hydrogenase to even subtle changes in the electronic structure and environment, as previously noted.<sup>18,37</sup> A supplementary discussion in the Supporting Information additionally compares the NRVS-observed and DFT-calculated  $^{57}\text{Fe}$ -PVDOS spectra of  $H_{\text{red}}H^+$  and  $H_{\text{sred}}H^+$  vs that of the earlier reported  $H_{\text{hyd}}$  state<sup>18</sup> (Figure S25), adding consistency to the present results.

To summarize, FTIR and NRVS spectra coupled with DFT calculations show clear evidence for the presence of a bridging CO in the  $H_{\text{red}}H^+$  and  $H_{\text{sred}}H^+$  states of the  $[\text{FeFe}]$  hydrogenase. These data exclude the possibility of a bridging hydride in these states. Our DFT rationalization of the IR signature with the prominent 1800–1820  $\text{cm}^{-1}$  feature appears to be satisfactory only when using the bridging CO H-cluster model. An absence of this low-frequency band in the IR reference may support a “rotated” isomer model having terminal CO ligands only, thus implying the bridging



hydride.<sup>26–28</sup> Furthermore, the previous analysis assumed that peaks in the 1950–1970  $\text{cm}^{-1}$  region derive from the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states. In contrast, we suggest that these peaks are attributed to an alternative  $\text{H}_{\text{hyd}}$  state, and instead we find a peak in the 1800–1820  $\text{cm}^{-1}$  region, which we assign to the bridging CO in the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states. We believe that if the present findings were accounted for in Haumann and co-workers' model, they may find that a  $\mu\text{CO}$  model is in better agreement with experimental data than a  $\mu\text{H}$  model.

We contend that there are no strong data indicating that the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states should be considered inactive. A number of observations hint that these states are, in fact, likely to be catalytically relevant. Under  $\text{H}_2$ , *CrHydA1* exists in a mixture of  $\text{H}_{\text{red}}\text{H}^+$ ,  $\text{H}_{\text{sred}}\text{H}^+$ , and  $\text{H}_{\text{hyd}}\text{H}^+$  states,<sup>34</sup> while *DdHydAB* is almost completely in the  $\text{H}_{\text{red}}\text{H}^+$  state.<sup>22</sup> Neither enzyme shows significant lag-phases during  $\text{H}_2$  oxidation<sup>43,44</sup> or  $\text{H}_2$  production,<sup>2,10</sup> suggesting that these states are catalytically active. Otherwise, the published activities of 1000–150 000  $\text{s}^{-1}$ <sup>2,10,43,44</sup> must represent the low activity of the inactive  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states, which seems unlikely. Formation of  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  requires reducing conditions at neutral to low pH, which are the conditions used for optimal  $\text{H}_2$  production. *DdHydAB* has a turnover frequency of 10 000  $\text{s}^{-1}$  for  $\text{H}_2$  production<sup>2</sup> and 150 000  $\text{s}^{-1}$  for  $\text{H}_2$  oxidation<sup>43</sup> and forms almost exclusively the  $\text{H}_{\text{red}}\text{H}^+$  state under reducing conditions at neutral pH.<sup>22</sup> Thus, if  $\text{H}_{\text{red}}\text{H}^+$  is essentially inactive, then the true activity of the *DdHydAB* should be much higher than 10 000  $\text{s}^{-1}$  for  $\text{H}_2$  production and much higher than 150 000  $\text{s}^{-1}$  for  $\text{H}_2$  oxidation, which also seems unlikely. Recently, time-resolved IR measurements of the rate of formation and decay of the various catalytic intermediates demonstrated the catalytic relevance of the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states in *CrHydA1*.<sup>45</sup>

The reason that the bridging CO has been so difficult to observe in the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states at room temperature is unclear. It may be related to flexibility of this ligand or coupling with other vibrational modes. Either of these two effects could potentially affect the oscillating dipole moment and thus the extinction coefficient for the IR transition. Since the local structure around the H-cluster may change slightly as the temperature is lowered, the bridging CO ligand could become more rigid, leading to a larger oscillating dipole moment and hence increased IR intensity. Notable in this regard is that the present DFT methodology for the normal-mode analysis and spectral simulations best corresponds to zero-point vibrations at absolute zero. Studies on CO-bound heme proteins have shown similar temperature-dependent peak broadening and have interpreted this as increased conformational flexibility at higher temperature, populating multiple conformational substates.<sup>46,47</sup> In agreement with this, the bridging CO can still be observed in the  $\text{H}_{\text{red}}^*$  and  $\text{H}_{\text{sred}}^*$  states of *TmHydS* at room temperature.<sup>20</sup> This thermostable [FeFe] hydrogenase may have decreased conformational flexibility around the H-cluster and thus fewer conformational substates.

Another interesting feature of the bridging CO ligands in the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states is their relatively high vibrational energy. The IR bands from the terminal CO and  $\text{CN}^-$  ligands generally shift to lower energy as  $[\text{2Fe}]_{\text{H}}$  becomes more reduced, due to increased metal–ligand  $\pi$  back-donation. However, the bridging CO in the  $\text{H}_{\text{red}}\text{H}^+$  state has a higher energy IR vibration (1810  $\text{cm}^{-1}$ ) than the  $\text{H}_{\text{ox}}$  state (1802  $\text{cm}^{-1}$ ). This may also reflect structural changes to the bridging

CO ligand upon  $[\text{2Fe}]_{\text{H}}$  reduction that prevent the increased  $\pi$  back-donation. Further IR studies coupled with DFT calculations and studies on amino acid variants may provide answers to these questions.

## CONCLUSIONS

This paper clarifies the structure of two controversial states invoked for the [FeFe] hydrogenases, a large family of highly active  $\text{H}_2$  conversion catalysts. Specifically, new spectroscopic and computational evidence is presented that both  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  contain bridging CO ligands, the presence of which precludes bridging hydride ligands.<sup>13,26–28</sup> We propose that the  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{sred}}\text{H}^+$  states are catalytically relevant intermediates. The results support the overall hypothesis that [FeFe] hydrogenases operate via terminal hydride-containing intermediates, which are rendered hydridic or acidic by proton-coupled electron transfer between the two halves of the H-cluster.<sup>15,17,21</sup>

## METHODS

Protein overproduction and artificial maturation were performed as described previously.<sup>48–50</sup>  $(\text{Et}_4\text{N})_2[\text{Fe}_2(\text{ADT})(\text{CO})_4(\text{CN})_2]$ ,  $((\text{Et}_4\text{N})_2[\text{2Fe}]^{\text{ADT}})$  and  $(\text{Et}_4\text{N})_2[\text{57Fe}_2(\text{ADT})(\text{CO})_4(\text{CN})_2]$ ,  $((\text{Et}_4\text{N})_2[\text{257Fe}]^{\text{ADT}})$  were synthesized following published procedures.<sup>51</sup> FTIR experiments were performed using a Bruker Vertex 80v FTIR spectrometer equipped with a nitrogen-cooled Bruker mercury cadmium telluride detector, liquid helium cooled OPTIA Optistat CF continuous flow cryostat system, liquid helium transfer line, and an Oxford Instruments ITC503 temperature controller. In an anaerobic chamber, samples (5  $\mu\text{L}$ ) of 1–2 mM *CrHydA1* or *DdHydAB* in 50 mM MES, 50 mM HEPES, and 300 mM KCl, pH 8, containing 20 mM sodium dithionite were loaded between  $\text{CaF}_2$  windows separated by a 50  $\mu\text{M}$  Teflon spacer and closed inside a brass holder with rubber rings. Spectra were collected in the double-sided, forward–backward mode with a resolution of 2  $\text{cm}^{-1}$ , an aperture setting of 1.5 mm, and a scan velocity of 20 Hz. Data were processed using home-written routines in the MATLAB environment.

NRVS spectra for  $[\text{257Fe}]_{\text{H}}\text{-CrHydA1}$  and  $[\text{257Fe}]_{\text{H}}\text{-DdHydAB}$  in 50 mM MES, 50 mM HEPES, and 300 mM KCl, pH/D 8, containing 20 mM sodium dithionite were recorded at Spring-8 BL09XU and BL19LXU. BL09XU uses a Si(111) double crystal in a high heat load monochromator to produce 14.414 keV radiation with  $\sim 1.0$  eV resolution, followed by a high energy resolution monochromator [Ge(422)x2Si(975)] to increase the resolution to  $\sim 0.8$  meV. The beam flux was  $\sim 2.5 \times 10^9$  photons/s, and the beam size was about 0.6 (height)  $\times$  1 (width)  $\text{mm}^2$ . A  $2 \times 2$  avalanche photodiode detector array was used to collect the delayed nuclear fluorescence and the  $K\delta$  fluorescence following nuclear excitation. The temperature at the base of the sample was maintained at 10 K with a closed cycle He cryostat. The Stokes/anti-Stokes imbalance derived real sample temperatures were 40–70 K. The setup at BL19LXU is similar to that at BL09XU except the average photo flux was  $\sim 5.4 \times 10^9$  photons/s and a liquid He cryostat was used. NRVS spectral analysis was performed using the PHOENIX software package executed through Spectratools.<sup>52</sup> Energy scale calibration was performed with a standard sample of  $[\text{NET}_4][\text{57FeCl}_4]$  with a prominent peak at 380  $\text{cm}^{-1}$ . Scans were divided into segments with different data collection times. In general, 1 s per point (s/p) was used for  $-200$  to 400  $\text{cm}^{-1}$ , 10 s/p for 400–600  $\text{cm}^{-1}$ , and 30 s/p for 600–800  $\text{cm}^{-1}$ . The scan ranges are all relative to the resonance energy.

The initial coordinates used for the DFT calculations were based on the 1.73 Å resolution 5BYQ X-ray data for semisynthetic *Clostridium pasteurianum* [FeFe] hydrogenase (CpI) matured with the ODT variant of the  $[\text{2Fe}]_{\text{H}}$  subcluster (CpI<sup>ODT</sup>).<sup>53</sup> A single serine side chain (S232 in CpI) had to be modified to alanine (A94 in *CrHydA1*). The approach used in the current work is analogous to our H-cluster DFT model construction described previously<sup>18,37</sup> and



denoted as  $L'$  (Large prime).  $L'$  corresponds to a molecular system including (i) a  $[2\text{Fe}]_{\text{H}}$  subcluster, (ii) its immediate protein environment, and (iii) the  $[4\text{Fe}-4\text{S}]_{\text{H}}$  subcluster as shown in Figures S14–S17. The structural optimizations and subsequent normal-mode calculations were performed using GAUSSIAN 09 Revision D.01<sup>54</sup> based on the densities exported from single-point calculations performed by JAGUAR 9.4,<sup>55</sup> which provided a high-quality initial guess. The PBE0<sup>56,57</sup> hybrid functional was employed in its unrestricted open-shell formulation. The LACV3P\*\* basis set as implemented in JAGUAR was employed. For the first- and second-row elements, LACV3P\*\* implies 6-311G\*\* triple- $\zeta$  basis sets including polarization functions. For the Fe atoms, LACV3P\*\* consists of a triple- $\zeta$  basis set for the outermost core and valence orbitals and the quasi-relativistic Los Alamos effective core potential for the innermost electrons. The molecular systems environment was considered using a self-consistent reaction field polarizable continuum model and integral equation formalism (IEF-PCM)<sup>58</sup> as implemented in GAUSSIAN 09, with the static dielectric constant set to  $\epsilon = 4.0$  as often used for proteins, and the remaining IEF-PCM parameters at their default values for water. Furthermore, the computational scheme included two-body D3 empirical dispersion correction by Grimme et al. in its original formulation.<sup>59</sup> The  $^{57}\text{Fe}$ -PVDOS and IR intensities were extracted from normal-mode outputs using an in-house program, Q-SPECTOR, successfully applied to simulate the  $[\text{FeFe}]$  hydrogenase NRVS spectra in previous works.<sup>18,37,38</sup> To empirically account for the experimental line shape, the computed intensities were broadened by Lorentzian convolution with a full width at half-maximum (fwhm) =  $14\text{ cm}^{-1}$  for  $^{57}\text{Fe}$ -PVDOS and by pseudo-Voigtian convolution with an fwhm =  $8\text{ cm}^{-1}$  for IR. For  $^{57}\text{Fe}$ -PVDOS, empirical scaling by 0.94 was applied to the calculated frequencies in the  $>400\text{ cm}^{-1}$  region. For IR, the calculated CO/CN frequencies from the representative  $\mu\text{CO}^+$  and  $\mu\text{CO}$  models were fit linearly to the observed  $\text{H}_{\text{red}}\text{H}^+$  and  $\text{H}_{\text{red}}\text{H}^+$  FTIR band positions, producing the 0.877 slope and  $+140.88\text{ cm}^{-1}$  offset parameters. For the H-cluster model as described above, H/D isotope exchange implied proton(s) of the ADT bridgehead, thiol proton of the adjacent cysteine (C169/178 in the CrHydA1/DdHydAB sequences, respectively), and the hydride if present, as indicated in Figure 4.

## ■ ASSOCIATED CONTENT

### ■ Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/jacs.9b09745>.

Supplementary FTIR, NRVS, and DFT, supplementary discussion, and references (PDF)

Coordinates of the DFT models as XYZ files (ZIP)

Animated vibrational normal modes as GIF files (ZIP)

## ■ AUTHOR INFORMATION

### Corresponding Author

\*james.birrell@cec.mpg.de

### ORCID

James A. Birrell: 0000-0002-0939-0573

Vladimir Pelmenschikov: 0000-0002-0523-4418

Thomas B. Rauchfuss: 0000-0003-2547-5128

Stephen P. Cramer: 0000-0002-3751-7623

Wolfgang Lubitz: 0000-0001-7059-5327

Serena DeBeer: 0000-0002-5196-3400

### Author Contributions

<sup>†</sup>J.A.B. and V.P. contributed equally to this work.

### Notes

The authors declare no competing financial interest.

## ■ ACKNOWLEDGMENTS

This work was supported by NIH GM-65440 (S.P.C.) and NIH GM-61153 (T.B.R.), by the Deutsche Forschungsgemeinschaft (DFG, German Research Foundation) under Germany's Excellence Strategy – EXC 2008/1 (UniSysCat) – 390540038 (V.P.), and by the Max Planck Society (J.A.B., W.L., and S.D.). S.D. and J.A.B. acknowledge funding from the DFG SPP 1927 “Iron–Sulfur for Life” project (Project Nos. DE 1877/1-1 and BI 2198/1-1). NRVS data collection was supported by a long-term proposal at BL09XU [2018B0141] and a general proposal at BLXU19 [2018B1379] at SPring-8. The authors thank Nina Breuer for help with enzyme preparation.

## ■ REFERENCES

- (1) Lubitz, W.; Ogata, H.; Rüdiger, O.; Reijerse, E. *Hydrogenases*. *Chem. Rev.* **2014**, *114* (8), 4081–4148.
- (2) Glick, B. R.; Martin, W. G.; Martin, S. M. Purification and properties of the periplasmic hydrogenase from *Desulfovibrio desulfuricans*. *Can. J. Microbiol.* **1980**, *26* (10), 1214–1223.
- (3) Peters, J. W.; Lenz, W. N.; Lemon, B. J.; Seefeldt, L. C. X-ray crystal structure of the Fe-only hydrogenase (CpI) from *Clostridium pasteurianum* to 1.8 angstrom resolution. *Science* **1998**, *282* (5395), 1853–1858.
- (4) Nicolet, Y.; Piras, C.; Legrand, P.; Hatchikian, C. E.; Fontecilla-Camps, J. C. *Desulfovibrio desulfuricans* iron hydrogenase: the structure shows unusual coordination to an active site Fe binuclear center. *Structure* **1999**, *7* (1), 13–23.
- (5) Nicolet, Y.; de Lacey, A. L.; Vernède, X.; Fernandez, V. M.; Hatchikian, E. C.; Fontecilla-Camps, J. C. Crystallographic and FTIR spectroscopic evidence of changes in Fe coordination upon reduction of the active site of the Fe-only hydrogenase from *Desulfovibrio desulfuricans*. *J. Am. Chem. Soc.* **2001**, *123* (8), 1596–1601.
- (6) Silakov, A.; Wenk, B.; Reijerse, E.; Lubitz, W.  $^{14}\text{N}$  HYSCORE investigation of the H-cluster of  $[\text{FeFe}]$  hydrogenase: evidence for a nitrogen in the dithiol bridge. *Phys. Chem. Chem. Phys.* **2009**, *11* (31), 6592–6599.
- (7) Lemon, B. J.; Peters, J. W. Binding of exogenously added carbon monoxide at the active site of the iron-only hydrogenase (CpI) from *Clostridium pasteurianum*. *Biochemistry* **1999**, *38* (40), 12969–73.
- (8) Swanson, K. D.; Ratzloff, M. W.; Mulder, D. W.; Artz, J. H.; Ghose, S.; Hoffman, A.; White, S.; Zadovnyy, O. A.; Broderick, J. B.; Bothner, B.; King, P. W.; Peters, J. W.  $[\text{FeFe}]$ -hydrogenase oxygen inactivation is initiated at the H cluster 2Fe subcluster. *J. Am. Chem. Soc.* **2015**, *137* (5), 1809–1816.
- (9) Happe, T.; Naber, J. D. Isolation, characterization and N-terminal amino acid sequence of hydrogenase from the green alga *Chlamydomonas reinhardtii*. *Eur. J. Biochem.* **1993**, *214* (2), 475–481.
- (10) Kamp, C.; Silakov, A.; Winkler, M.; Reijerse, E. J.; Lubitz, W.; Happe, T. Isolation and first EPR characterization of the  $[\text{FeFe}]$ -hydrogenases from green algae. *Biochim. Biophys. Acta, Bioenerg.* **2008**, *1777* (5), 410–416.
- (11) Adams, M. W.; Eccleston, E.; Howard, J. B. Iron-sulfur clusters of hydrogenase I and hydrogenase II of *Clostridium pasteurianum*. *Proc. Natl. Acad. Sci. U. S. A.* **1989**, *86* (13), 4932–4936.
- (12) Grande, H. J.; Dunham, W. R.; Averill, B.; Van Dijk, C.; Sands, R. H. Electron paramagnetic resonance and other properties of hydrogenases isolated from *Desulfovibrio vulgaris* (strain Hildenborough) and *Megasphaera elsdenii*. *Eur. J. Biochem.* **1983**, *136* (1), 201–7.
- (13) Haumann, M.; Stripp, S. T. The molecular proceedings of biological hydrogen turnover. *Acc. Chem. Res.* **2018**, *51* (8), 1755–1763.
- (14) Ratzloff, M. W.; Artz, J. H.; Mulder, D. W.; Collins, R. T.; Furtak, T. E.; King, P. W. CO-bridged H-cluster intermediates in the catalytic mechanism of  $[\text{FeFe}]$ -hydrogenase CaI. *J. Am. Chem. Soc.* **2018**, *140* (24), 7623–7628.

- (15) Sommer, C.; Adamska-Venkatesh, A.; Pawlak, K.; Birrell, J. A.; Rüdiger, O.; Reijerse, E. J.; Lubitz, W. Proton coupled electronic rearrangement within the H-cluster as an essential step in the catalytic cycle of [FeFe] hydrogenases. *J. Am. Chem. Soc.* **2017**, *139* (4), 1440–1443.
- (16) Mulder, D. W.; Ratzloff, M. W.; Bruschi, M.; Greco, C.; Koonce, E.; Peters, J. W.; King, P. W. Investigations on the role of proton-coupled electron transfer in hydrogen activation by [FeFe]-hydrogenase. *J. Am. Chem. Soc.* **2014**, *136* (43), 15394–15402.
- (17) Rodríguez-Maciá, P.; Pawlak, K.; Rüdiger, O.; Reijerse, E. J.; Lubitz, W.; Birrell, J. A. Intercluster redox coupling influences protonation at the H-cluster in [FeFe] hydrogenases. *J. Am. Chem. Soc.* **2017**, *139* (42), 15122–15134.
- (18) Pelmenschikov, V.; Birrell, J. A.; Pham, C. C.; Mishra, N.; Wang, H.; Sommer, C.; Reijerse, E.; Richers, C. P.; Tamasaku, K.; Yoda, Y.; Rauchfuss, T. B.; Lubitz, W.; Cramer, S. P. Reaction coordinate leading to H<sub>2</sub> production in [FeFe]-hydrogenase identified by nuclear resonance vibrational spectroscopy and density functional theory. *J. Am. Chem. Soc.* **2017**, *139* (46), 16894–16902.
- (19) Rodríguez-Maciá, P.; Reijerse, E. J.; van Gestel, M.; DeBeer, S.; Lubitz, W.; Rüdiger, O.; Birrell, J. A. Sulfide protects [FeFe] hydrogenases from O<sub>2</sub>. *J. Am. Chem. Soc.* **2018**, *140* (30), 9346–9350.
- (20) Chongdar, N.; Birrell, J. A.; Pawlak, K.; Sommer, C.; Reijerse, E. J.; Rüdiger, O.; Lubitz, W.; Ogata, H. Unique spectroscopic properties of the H-cluster in a putative sensory [FeFe] hydrogenase. *J. Am. Chem. Soc.* **2018**, *140* (3), 1057–1068.
- (21) Rodríguez-Maciá, P.; Kertess, L.; Burnik, J.; Birrell, J. A.; Hofmann, E.; Lubitz, W.; Happe, T.; Rüdiger, O. His-ligation to the [4Fe-4S] subcluster tunes the catalytic bias of [FeFe] hydrogenase. *J. Am. Chem. Soc.* **2019**, *141* (1), 472–481.
- (22) Pierik, A. J.; Hulstein, M.; Hagen, W. R.; Albracht, S. P. A low-spin iron with CN and CO as intrinsic ligands forms the core of the active site in [Fe]-hydrogenases. *Eur. J. Biochem.* **1998**, *258* (2), 572–578.
- (23) Roseboom, W.; de Lacey, A. L.; Fernandez, V. M.; Hatchikian, E. C.; Albracht, S. P. The active site of the [FeFe]-hydrogenase from *Desulfovibrio desulfuricans*. II. Redox properties, light sensitivity and CO-ligand exchange as observed by infrared spectroscopy. *JBIC, J. Biol. Inorg. Chem.* **2006**, *11* (1), 102–118.
- (24) Katz, S.; Noth, J.; Horch, M.; Shafaat, H. S.; Happe, T.; Hildebrandt, P.; Zebger, I. Vibrational spectroscopy reveals the initial steps of biological hydrogen evolution. *Chem. Sci.* **2016**, *7* (11), 6746–6752.
- (25) Hajj, V.; Baffert, C.; Sybirna, K.; Meynial-Salles, I.; Soucaille, P.; Bottin, H.; Fourmond, V.; Léger, C. FeFe hydrogenase reductive inactivation and implication for catalysis. *Energy Environ. Sci.* **2014**, *7* (2), 715–719.
- (26) Mebs, S.; Senger, M.; Duan, J.; Wittkamp, F.; Apfel, U. P.; Happe, T.; Winkler, M.; Stripp, S. T.; Haumann, M. Bridging hydride at reduced H-cluster species in [FeFe]-hydrogenases revealed by infrared spectroscopy, isotope editing, and quantum chemistry. *J. Am. Chem. Soc.* **2017**, *139* (35), 12157–12160.
- (27) Chernev, P.; Lambert, C.; Brunje, A.; Leidel, N.; Sigfridsson, K. G.; Kositzki, R.; Hsieh, C. H.; Yao, S.; Schiwon, R.; Driess, M.; Limberg, C.; Happe, T.; Haumann, M. Hydride binding to the active site of [FeFe]-hydrogenase. *Inorg. Chem.* **2014**, *53* (22), 12164–12177.
- (28) Mebs, S.; Duan, J.; Wittkamp, F.; Stripp, S. T.; Happe, T.; Apfel, U. P.; Winkler, M.; Haumann, M. Differential protonation at the catalytic six-iron cofactor of [FeFe]-Hydrogenases revealed by <sup>57</sup>Fe nuclear resonance X-ray scattering and quantum mechanics/molecular mechanics analyses. *Inorg. Chem.* **2019**, *58* (6), 4000–4013.
- (29) Bruschi, M.; Greco, C.; Kaukonen, M.; Fantucci, P.; Ryde, U.; de, G. L. Influence of the [2Fe]<sub>H</sub> subcluster environment on the properties of key intermediates in the catalytic cycle of [FeFe] hydrogenases: hints for the rational design of synthetic catalysts. *Angew. Chem., Int. Ed.* **2009**, *48* (19), 3503–3506.
- (30) Filippi, G.; Arrigoni, F.; Bertini, L.; De Gioia, L.; Zampella, G. DFT dissection of the reduction step in H<sub>2</sub> catalytic production by [FeFe]-hydrogenase-inspired models: can the bridging hydride become more reactive than the terminal isomer? *Inorg. Chem.* **2015**, *54* (19), 9529–9542.
- (31) Zampella, G.; Fantucci, P.; Gioia, L. D. Unveiling how stereoelectronic factors affect kinetics and thermodynamics of protonation regiochemistry in [FeFe] hydrogenase synthetic models: a DFT investigation. *J. Am. Chem. Soc.* **2009**, *131* (31), 10909–10917.
- (32) Pullen, S.; Maji, S.; Stein, M.; Ott, S. Restricted rotation of an Fe(CO)<sub>2</sub>(PL<sub>3</sub>)-subunit in [FeFe]-hydrogenase active site mimics by intramolecular ligation. *Dalton Trans.* **2019**, *48* (18), 5933–5939.
- (33) Finkelmann, A. R.; Stiebritz, M. T.; Reiher, M. Inaccessibility of the  $\mu$ -hydride species in [FeFe] hydrogenases. *Chem. Sci.* **2014**, *5* (1), 215–221.
- (34) Adamska, A.; Silakov, A.; Lambert, C.; Rüdiger, O.; Happe, T.; Reijerse, E.; Lubitz, W. Identification and characterization of the “super-reduced” state of the H-cluster in [FeFe] hydrogenase: a new building block for the catalytic cycle? *Angew. Chem., Int. Ed.* **2012**, *51* (46), 11458–11462.
- (35) Silakov, A.; Kamp, C.; Reijerse, E.; Happe, T.; Lubitz, W. Spectroelectrochemical characterization of the active site of the [FeFe] hydrogenase HydA1 from *Chlamydomonas reinhardtii*. *Biochemistry* **2009**, *48* (33), 7780–7786.
- (36) Mulder, D. W.; Guo, Y.; Ratzloff, M. W.; King, P. W. Identification of a catalytic iron-hydride at the H-Cluster of [FeFe]-hydrogenase. *J. Am. Chem. Soc.* **2017**, *139* (1), 83–86.
- (37) Pham, C. C.; Mulder, D. W.; Pelmenschikov, V.; King, P. W.; Ratzloff, M. W.; Wang, H.; Mishra, N.; Alp, E. E.; Zhao, J.; Hu, M. Y.; Tamasaku, K.; Yoda, Y.; Cramer, S. P. Terminal hydride species in [FeFe]-hydrogenases are vibrationally coupled to the active site environment. *Angew. Chem., Int. Ed.* **2018**, *57* (33), 10605–10609.
- (38) Reijerse, E. J.; Pham, C. C.; Pelmenschikov, V.; Gilbert-Wilson, R.; Adamska-Venkatesh, A.; Siebel, J. F.; Gee, L. B.; Yoda, Y.; Tamasaku, K.; Lubitz, W.; Rauchfuss, T. B.; Cramer, S. P. Direct observation of an iron-bound terminal hydride in [FeFe]-hydrogenase by nuclear resonance vibrational spectroscopy. *J. Am. Chem. Soc.* **2017**, *139* (12), 4306–4309.
- (39) Winkler, M.; Senger, M.; Duan, J.; Esselborn, J.; Wittkamp, F.; Hofmann, E.; Apfel, U. P.; Stripp, S. T.; Happe, T. Accumulating the hydride state in the catalytic cycle of [FeFe]-hydrogenases. *Nat. Commun.* **2017**, *8*, 16115.
- (40) Ogata, H.; Krämer, T.; Wang, H.; Schilter, D.; Pelmenschikov, V.; van Gestel, M.; Neese, F.; Rauchfuss, T. B.; Gee, L. B.; Scott, A. D.; Yoda, Y.; Tanaka, Y.; Lubitz, W.; Cramer, S. P. Hydride bridge in [NiFe]-hydrogenase observed by nuclear resonance vibrational spectroscopy. *Nat. Commun.* **2015**, *6*, 7890.
- (41) Carlson, M. R.; Gray, D. L.; Richers, C. P.; Wang, W.; Zhao, P.-H.; Rauchfuss, T. B.; Pelmenschikov, V.; Pham, C. C.; Gee, L. B.; Wang, H.; Cramer, S. P. Sterically stabilized terminal hydride of a diiron dithiolate. *Inorg. Chem.* **2018**, *57* (4), 1988–2001.
- (42) Pelmenschikov, V.; Guo, Y.; Wang, H.; Cramer, S. P.; Case, D. A. Fe–H/D stretching and bending modes in nuclear resonant vibrational, Raman and infrared spectroscopies: Comparisons of density functional theory and experiment. *Faraday Discuss.* **2011**, *148* (0), 409–420.
- (43) Hatchikian, E. C.; Forget, N.; Fernandez, V. M.; Williams, R.; Cammack, R. Further characterization of the [Fe]-hydrogenase from *Desulfovibrio desulfuricans* ATCC 7757. *Eur. J. Biochem.* **1992**, *209* (1), 357–365.
- (44) Siebel, J. F.; Adamska-Venkatesh, A.; Weber, K.; Rumpel, S.; Reijerse, E.; Lubitz, W. Hybrid [FeFe]-hydrogenases with modified active sites show remarkable residual enzymatic activity. *Biochemistry* **2015**, *54* (7), 1474–1483.
- (45) Sanchez, M. L. K.; Sommer, C.; Reijerse, E.; Birrell, J. A.; Lubitz, W.; Dyer, R. B. Investigating the kinetic competency of CrHydA1 [FeFe] hydrogenase intermediate states via time-resolved infrared spectroscopy. *J. Am. Chem. Soc.* **2019**, *141* (40), 16064–16070.
- (46) Hong, M. K.; Braunstein, D.; Cowen, B. R.; Frauenfelder, H.; Iben, I. E.; Mourant, J. R.; Ormos, P.; Scholl, R.; Schulte, A.

Steinbach, P. J.; et al. Conformational substates and motions in myoglobin. External influences on structure and dynamics. *Biophys. J.* **1990**, *58* (2), 429–436.

(47) Kaposi, A. D.; Vanderkooi, J. M.; Stavrov, S. S. Infrared absorption study of the heme pocket dynamics of carbonmonoxyheme proteins. *Biophys. J.* **2006**, *91* (11), 4191–4200.

(48) Kuchenreuther, J. M.; Grady-Smith, C. S.; Bingham, A. S.; George, S. J.; Cramer, S. P.; Swartz, J. R. High-yield expression of heterologous [FeFe] hydrogenases in *Escherichia coli*. *PLoS One* **2010**, *5* (11), No. e15491.

(49) Birrell, J. A.; Wrede, K.; Pawlak, K.; Rodríguez-Maciá, P.; Rüdiger, O.; Reijerse, E. J.; Lubitz, W. Artificial maturation of the highly active heterodimeric [FeFe] hydrogenase from *Desulfovibrio desulfuricans* ATCC 7757. *Isr. J. Chem.* **2016**, *56* (9–10), 852–863.

(50) Esselborn, J.; Lambertz, C.; Adamska-Venkatesh, A.; Simmons, T.; Berggren, G.; Noth, J.; Siebel, J.; Hemschemeier, A.; Artero, V.; Reijerse, E.; Fontecave, M.; Lubitz, W.; Happe, T. Spontaneous activation of [FeFe]-hydrogenases by an inorganic [2Fe] active site mimic. *Nat. Chem. Biol.* **2013**, *9* (10), 607–609.

(51) Li, H.; Rauchfuss, T. B. Iron carbonyl sulfides, formaldehyde, and amines condense to give the proposed azadithiolate cofactor of the Fe-only hydrogenases. *J. Am. Chem. Soc.* **2002**, *124* (5), 726–727.

(52) Sturhahn, W. CONUSS and PHOENIX: Evaluation of nuclear resonant scattering data. *Hyperfine Interact.* **2000**, *125* (1), 149–172.

(53) Esselborn, J.; Muraki, N.; Klein, K.; Engelbrecht, V.; Metzler-Nolte, N.; Apfel, U. P.; Hofmann, E.; Kurisu, G.; Happe, T. A structural view of synthetic cofactor integration into [FeFe]-hydrogenases. *Chem. Sci.* **2016**, *7* (2), 959–968.

(54) Frisch, M. J. T. G. W.; Schlegel, H. B.; Scuseria, G. E.; Robb, M. A.; Cheeseman, J. R.; Scalmani, G.; Barone, V.; Mennucci, B.; Petersson, G. A.; Nakatsuji, H.; Caricato, M.; Li, X.; Hratchian, H. P.; Izmaylov, A. F.; Bloino, J.; Zheng, G.; Sonnenberg, J. L.; Hada, M.; Ehara, M.; Toyota, K.; Fukuda, R.; Hasegawa, J.; Ishida, M.; Nakajima, T.; Honda, Y.; Kitao, O.; Nakai, H.; Vreven, T.; Montgomery, J. A., Jr.; Peralta, J. E.; Ogliaro, F.; Bearpark, M.; Heyd, J. J.; Brothers, E.; Kudin, K. N.; Staroverov, V. N.; Kobayashi, R.; Normand, J.; Raghavachari, K.; Rendell, A.; Burant, J. C.; Iyengar, S. S.; Tomasi, J.; Cossi, M.; Rega, N.; Millam, J. M.; Klene, M.; Knox, J. E.; Cross, J. B.; Bakken, V.; Adamo, C.; Jaramillo, J.; Gomperts, R.; Stratmann, R. E.; Yazyev, O.; Austin, A. J.; Cammi, R.; Pomelli, C.; Ochterski, J. W.; Martin, R. L.; Morokuma, K.; Zakrzewski, V. G.; Voth, G. A.; Salvador, P.; Dannenberg, J. J.; Dapprich, S.; Daniels, A. D.; Farkas, Ö.; Foresman, J. B.; Ortiz, J. V.; Cioslowski, J.; Fox, D. J. *Gaussian 09*, Revision D.01; Gaussian Inc.: Wallingford, CT, 2009.

(55) *Jaguar*, version 9.4; Schrodinger, Inc.: New York, NY, 2016.

(56) Adamo, C.; Barone, V. Toward reliable density functional methods without adjustable parameters: The PBE0 model. *J. Chem. Phys.* **1999**, *110* (13), 6158–6170.

(57) Perdew, J. P.; Burke, K.; Ernzerhof, M. Generalized Gradient Approximation Made Simple. *Phys. Rev. Lett.* **1996**, *77* (18), 3865–3868.

(58) Tomasi, J.; Mennucci, B.; Cammi, R. Quantum Mechanical Continuum Solvation Models. *Chem. Rev.* **2005**, *105* (8), 2999–3094.

(59) Grimme, S.; Antony, J.; Ehrlich, S.; Krieg, H. A consistent and accurate ab initio parametrization of density functional dispersion correction (DFT-D) for the 94 elements H–Pu. *J. Chem. Phys.* **2010**, *132* (15), 154104.