

Supporting Information

The non-physiological reductant sodium dithionite and [FeFe] hydrogenase: influence on the enzyme mechanism

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Experimental Section

Protein production

Sodium dithionite (NaDT) was excluded from all protein preparations to avoid contamination by its oxidation products. *CrHydA1*, *DdHydAB* and *CpHydA1* were recombinantly produced in *E. coli* as apo-proteins (lacking the [2Fe]_H subcluster). Samples were then artificially matured with the [2Fe]_H precursor (Et₄N)₂[Fe₂(ADT)(CO)₄(CN)₂] (using a ⁵⁷Fe-labeled version for NRVS experiments) or (Et₄N)₂[Fe₂(PDT)(CO)₄(CN)₂], as previously described¹⁻³. (Et₄N)₂[Fe₂(ADT)(CO)₄(CN)₂] and (Et₄N)₂[Fe₂(ADT)(CO)₄(CN)₂] were prepared as previously described^{4,5}.

Sample preparation for FTIR Spectroscopy

Concentrated (2-5 mM) protein samples (in 25 mM Tris-HCl, 25 mM KCl, pH 8) were diluted (1 μL in 12 μL total volume) in 20 mM Na-acetate, 20 mM MES, 20 mM HEPES, 20 mM Tris, 20 mM glycine buffer (20 mM mixed buffer) adjusted to the desired pH with NaOH or HCl. Na₂SO₃, Na₂SO₄, Na₂S₂O₃ (all prepared in 20 mM mixed buffer) or oxidized NaDT (oxNaDT, prepared in water) were added at various concentrations (by substituting a fraction of the total 12 μL volume of the FTIR sample with a given volume of stock solution) as specified for each experiment. All solutions, except oxNaDT for some experiments where stated, were pH corrected before use. The solution of oxNaDT was prepared under aerobic conditions using fresh NaDT (1 M) and was left stirring at room temperature for 2 h under atmospheric oxygen to fully oxidize the dithionite anion. Before being added to the protein samples, the solution was degassed and moved into an anaerobic glovebox filled with N₂. To prepare *CrHydA1* in the H_{ox}-CO and H_{ox}H-CO states, samples (with or without Na₂SO₃) were flushed with 100% CO gas for 10 min before being loaded into the FTIR cell. For all Na₂SO₃ solutions, 1 M stock solutions were prepared in 20 mM mixed buffer, aerobically, and adjusted to the desired pH.

FTIR Spectroscopy

FTIR spectra were measured using a Bruker Vertex 80v FTIR spectrometer equipped with a mercury cadmium telluride detector cooled by liquid nitrogen. Spectra were recorded in the double-sided, forward-backward mode, with a resolution of 2 cm⁻¹, an aperture setting of 3 mm and scan velocity of 20 kHz. Spectra are the average of 100 scans. Samples (12 μL) were sealed between CaF₂ windows (20 mm x 4 mm, Korth Kristalle, Altenholz) separated by a 50 μm Teflon spacer coated with vacuum grease and closed in a homebuilt IR cell with rubber rings. FTIR data were processed using home-written routines in the MATLAB environment. For *CrHydA1*^{PDT} prepared under 2% H₂, the region between 1955–1920 cm⁻¹ (corresponding to the terminal CO on Fe_d) in each spectrum was fitted with pseudo-Voigt functions using QSoas 3.0 (ref ⁶) to identify the individual contributions from the H_{ox}, H_{red}, H_{ox}H and H_{red}'H states to the FTIR absorbance. Fit parameters (Table S3) were obtained for H_{ox} and H_{red} from spectra in the absence of Na₂SO₃. Parameters for H_{ox}H and H_{red}'H were obtained from the analysis of spectra at low pH and high Na₂SO₃ concentrations, where these were the dominant states, and by fixing the already determined parameters for H_{ox} and H_{red}. Fit parameters for the four states were then fixed in all subsequent fits (Table S3), allowing only the peak intensity to vary. Good agreement between simulated and experimental data was obtained for all spectra.

NRVS

NRVS spectra for [2⁵⁷Fe]_H-*CrHydA1* in 20 mM mixed buffer pH 6, with or without 1 M Na₂SO₃, were recorded at P01, PETRA-III, Hamburg.⁷ The operation of the storage ring was a 40-bunch mode (bunch separation of 192 ns) with a beam energy of 6 GeV. The experimental setup established at the beamline including a two-step monochromatization (energy resolution of ~ 1 meV) and detection by avalanche photo diodes was used.⁷ Samples, in home-built copper samples holders, were positioned in the beam on the top of a closed cycle helium cryostat with a copper cold finger, using a home-built copper adapter. The sample was then covered with a mylar cylinder and a metal cover with a beryllium window. The temperature at the

base of the sample was maintained at 10 K and the Stokes/anti-Stokes imbalance derived real sample temperatures were 10–20 K. The NIS data were collected during several scans within the energy range of –20 to 100 meV with a 0.25 meV step size and a measuring time of 5 s per energy step. Data were collected as the raw nuclear inelastic scattering (NIS) counts. Calculation of partial density of vibrational states (PVDOS) was performed using routines written in Python with a binning of 0.5 meV.

Electrochemistry

DdHydAB was covalently attached to a pyrolytic graphite electrode (0.031 cm², homemade using pyrolytic graphite from Momentive Materials) modified with 4-nitrobenzenediazonium salt as described previously.⁸ Once modified, the electrode was moved inside a glovebox filled with N₂ and subjected to 10 cycles of voltammetry from +0.241 to –0.659 V to remove oxygen adsorbed on the electrode surface. *DdHydAB* was immobilized by covering the electrode with 6 μL of protein solution (10 μM in 10 mM MES pH 5.8). The protein was left to adsorb for 20 min. Subsequently, N-hydroxysuccinimide (NHS, 4.5 μL of 128 mM solution) and N-(3-dimethylaminopropyl)-N'-ethylcarbodiimide (EDC, 5.5 μL of 210 mM solution), both dissolved in 10 mM MES pH 5.8, were added to the electrode and left to react for 90 min. The electrode was rinsed with water and then in the appropriate experimental buffer by rotating the electrode inside the buffer solution for 1 min. Electrochemical measurements were performed in a standard three-electrode electrochemical cell with a platinum wire as counter electrode and a saturated calomel reference electrode, separated from the main compartment in a sidearm containing 0.1 M KCl, connected to the main cell compartment by a Luggin capillary. The reference electrode potential was calibrated using (hydroxymethyl)ferrocene (+436 mV vs SHE) as a reference.⁹ The potential was controlled by a VersaSTAT 4–400 potentiostat. A set of mass flow-controllers was used to control the composition of the gas flushed through the electrochemical cell. A cell temperature of 25 °C was maintained by a waterjacket system and a water circulator. All the experiments were performed in 20 mM mixed buffer at the desired pH with addition of 0.1 M NaCl. All potentials are quoted versus the standard hydrogen electrode (SHE).

Supplementary Discussion: Origins of $H_{ox}H$ and $H_{red}'H$

As mentioned in the main text, the $H_{ox}H$ and $H_{red}'H$ states in Model 2 have been reported to accumulate at low pH in the presence of sodium dithionite (NaDT) and were proposed to be protonated versions of H_{ox} and H_{red}' , respectively, based on density functional theory (DFT) calculations.^{10, 11} $H_{ox}H$, originally identified during studies of the H_{hyd} state, exhibits an infrared (IR) spectrum similar to that of H_{ox} , but with vibrational frequencies shifted higher in energy by 4-6 cm^{-1} (Table S1).¹² Stripp and coworkers suggested that the $H_{ox}H$ state is protonated on the sulfur of one of the cysteine residues ligating $[4Fe-4S]_H$,¹¹ the same cysteine proposed to be protonated in H_{red}' ,¹⁰ which in Model 2 forms from H_{ox} via PCET at $[4Fe-4S]_H$. This protonation is thought to have a regulatory effect on catalysis, preventing the formation of the low-activity reduced states containing a μH^- .¹³⁻¹⁵ Recently, the presence of a μH^- in the $H_{red}H^+$ and $H_{sred}H^+$ states has been disputed,¹⁶⁻¹⁸ and the redox potential of $[4Fe-4S]_H$ has been shown to be pH independent,¹⁹ challenging the proposed PCET at $[4Fe-4S]_H$ in the formation of H_{red}' .

As the reductant NaDT is needed for the formation of $H_{ox}H$, this state was proposed to lie at the end of the catalytic cycle: after H_{red}' and H_{hyd} , where the proton on $[4Fe-4S]_H$ is retained. Deprotonation of $H_{ox}H$, which would reform H_{ox} , is hindered at low pH, potentially explaining why this state accumulates under acidic conditions. However, it is unclear why the H_{ox} state does not simply convert to $H_{ox}H$ at low pH. The $H_{red}'H$ state was reported to form upon reduction and protonation of $H_{ox}H$, under H_2 or via electrochemical reduction, but again only in the presence of NaDT at low pH.^{10, 11} Similar to H_{ox} vs $H_{ox}H$, the infrared spectrum of $H_{red}'H$ resembles the one of H_{red}' , but with bands shifted to slightly higher energies (Table S1). DFT calculations suggested that, in $H_{red}'H$, a second proton, in addition to the one proposed to be already bound in $H_{ox}H$ and H_{red}' , binds to another cysteine residue ligating the $[4Fe-4S]_H$ subcluster. Whether $H_{red}'H$ is an intermediate of the catalytic cycle was not clarified. Importantly, both $H_{ox}H$ and $H_{red}'H$ have not been reported to form with chemical reducing agents other than NaDT, including H_2 , or via electrochemical reduction in the absence of NaDT. A satisfactory explanation for this behavior is still lacking.

Supplementary Tables

Table S1 – IR bands in [FeFe] hydrogenase states observed in this and previous studies

Enzyme	State [#]	CN _p	CN _d	CO _p	CO _d	CO _b	Ref.
<i>CrHydA1</i> ^{ADT}	H _{ox}	2089	2072	1964	1940	1804	Here
		2088	2072	1964	1940	1802	Ref ¹¹
	H _{ox} H	2092	2076	1971	1947	1813	Here
		2092	2076	1971	1946	1812	Ref ¹¹
	H _{red} H _{red} '	2084	2067	1962	1933	1793	Here
		2084	2066	1962	1933	1792	Ref ¹¹
H _{red} 'H	2086	2068	1966	1938	1800	Ref ¹¹	
H _{ox} -CO	2092	2082	1964	2013/1969*	1810	Here	
	2091	2081	1962	2012/1968*	1808	Ref ¹¹	
H _{ox} H-CO	2094	2086	1966	2006/1972*	1816	Here	
	2094	2086	1966	2006/1972*	1816	Ref ¹¹	
<i>CrHydA1</i> ^{PDT}	H _{ox}	2090	2073	1966	1942	1811	Here
		2090	2073	1965	1941	1810	Ref ¹¹
	H _{ox} H	2091	2075	1970	1946	1813	Here
		2090	2075	1969	1945	1814	Ref ¹¹
H _{red} H _{red} '	2085	2066	1964	1935	1799	Here	
	2084	2066	1963	1934	1798	Ref ¹¹	
H _{red} 'H	2084	2068	1966	1939	1802	Here	
	2084	2068	1966	1938	1802	Ref ¹¹	
<i>DdHydAB</i> ^{ADT}	H _{ox}	2087	2078	1965	1940	1802	Here
		2090	2078	1965	1940	1802	Ref ¹¹
H _{ox} H	2096	2082	1972	1947	1811	Here	
	2098	2083	1972	1947	1812	Ref ¹¹	
<i>CpHydA1</i> ^{ADT}	H _{ox}	2082	2070	1970	1947	1801	Here
		2081	2069	1970	1947	1801	Ref ¹¹
H _{ox} H	2084	2073	1975	1953	1810	Here	
	2085	2074	1975	1953	1808	Ref ¹¹	

when two nomenclatures are given, the one underneath is from Stripp and Haumann²⁰

*In the H_{ox}-CO and H_{ox}H-CO states the two CO ligands bound to Fe_d are strongly vibrationally coupled giving rise to a higher energy symmetric stretching band and a lower energy antisymmetric stretching band.

Table S2 – Comparison of IR bands in [FeFe] hydrogenase in the H_{ox} vs H_{ox}-DT_i and H_{hyd:red} vs H_{hyd} states

Enzyme	State	CN _p	CN _d	CO _p	CO _d	CO _b	Ref.
<i>CrHydA1</i> ^{ADT}	H _{ox}	2089	2072	1964	1940	1804(2)	Here
		2088	2072	1964	1940	1802	Ref ¹¹
	H _{ox} -DT _i	2092	2076	1971	1947	1813	Here
		2092	2074	1970	1946	1812	Ref ¹¹
	H _{hyd:ox}	2092	2086	1983	1954	1865	Ref ¹⁸
H _{hyd:red}	2087	2078	1972	1954	1851	Ref ¹⁸	
H _{hyd}	2088	2076	1980	1960	1860	Ref ¹²	

Table S3 – Parameters used for Pseudo-Voigt fitting

Pseudo-Voigt fitting parameters			
State	x_i	w_i	μ_i
H _{ox}	1942.08	1.79	0.426
H _{red}	1935.23	2.36	0.535
H _{ox} H	1946.15	1.83	0.546
H _{red} 'H	1938.67	2.76	0.705

Where x_i, w_i and μ_i are parameters for each of the four pseudo-Voigt peaks fit to the following formula, while A_i was allowed to vary:

$$f(x) = Y_0 + \sum_i A_i \left[\frac{1 - \mu_i}{\sqrt{2\pi w_i^2}} \exp(-(x - x_i)^2 / 2w_i^2) + \frac{\mu_i}{w_i \pi (1 + ((x - x_i) / w_i)^2)} \right]$$

x_i = peak center, w_i = broadening factor, μ_i = profile shape factor

Supplementary Figures

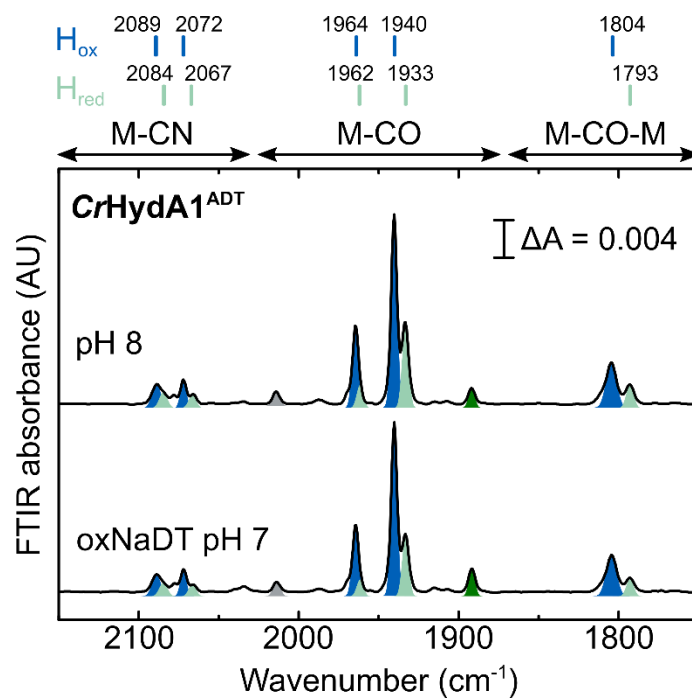


Figure S1 – Effect of oxNaDT on *CrHydA1*^{ADT} at pH 7. IR spectra of *CrHydA1* ($\approx 730 \mu\text{M}$) diluted in 25 mM Tris-HCl, 25 mM NaCl, pH 8 (top) and in a solution of oxidized NaDT (0.83 M final concentration) corrected to pH 7 (bottom). At neutral pH the H_{ox}H state does not form in the presence of the oxidation products of NaDT.

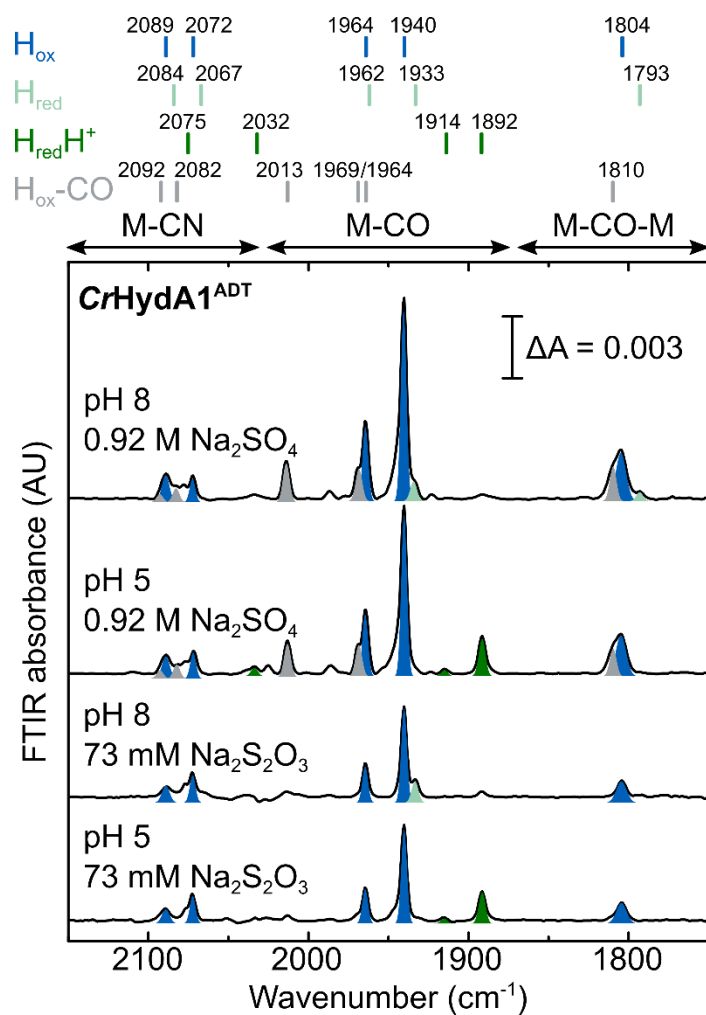


Figure S2 – Effect of Na₂SO₄ and Na₂S₂O₃ on CrHydA1^{ADT}. IR spectra of CrHydA1 (0.2 mM) diluted in 20 mM mixed buffer (see experimental section) at pH 8 and pH 5 in the presence of 0.92 M Na₂SO₄ or 73 mM Na₂S₂O₃.

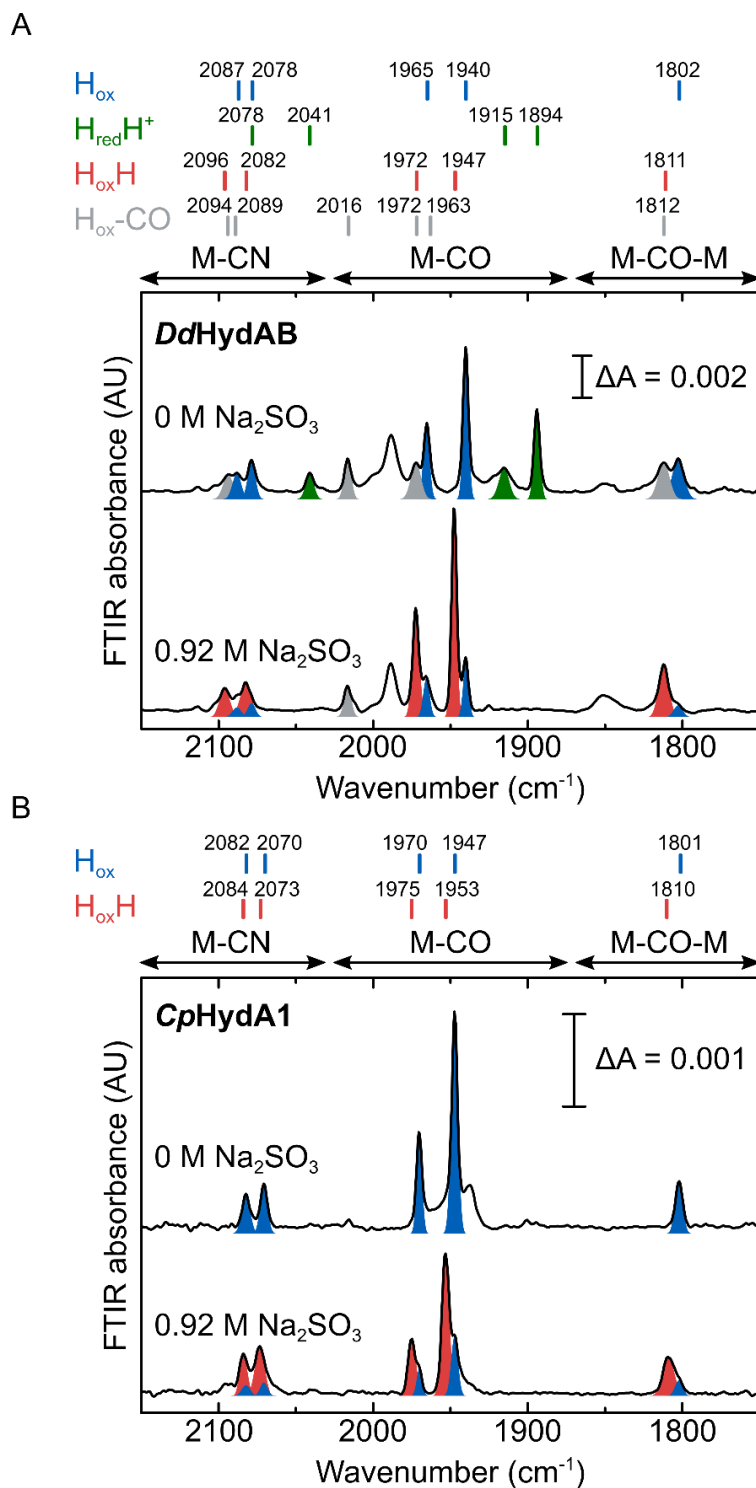


Figure S3 – Formation of the $H_{ox}H$ state in *DdHydAB* and *CpHydA1* with Na_2SO_3 and low pH. **A)** IR spectra of *DdHydAB* ($\approx 250 \mu M$) in 20 mM mixed buffer (see experimental section), pH 6 (top) and in 0.92 M Na_2SO_3 pH 6 (bottom). **B)** Same as **A** but with *CpHydA1* ($\approx 60 \mu M$). The unassigned peaks at 1988 cm^{-1} and 1950 cm^{-1} in the *DdHydAB* spectra are often observed for this enzyme and appear to be related to the longer artificial maturation process compared with *CpHydA1* and *CrHydA1*. Further investigation of the origin of these peaks will be the subject of ongoing investigation.

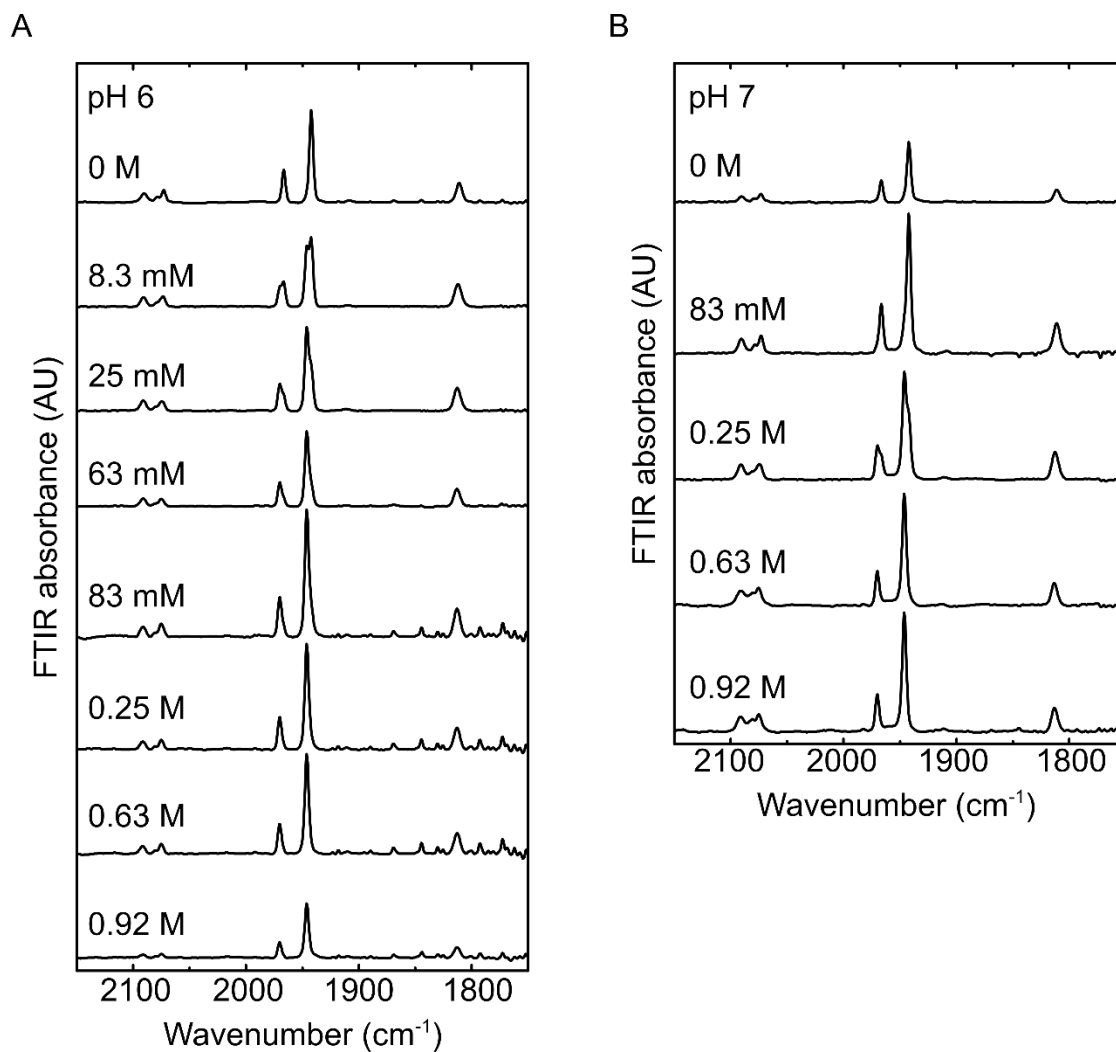


Figure S4 – Full titration of *CrHydA1*^{PDT} with Na₂SO₃ at pH 6 and pH 7 under N₂. IR spectra of *CrHydA1*^{PDT} ($\approx 420 \mu\text{M}$) diluted in 20 mM mixed buffer (see experimental section) at pH 6 (**A**) and pH 7 (**B**) with various concentrations of Na₂SO₃. Selected spectra are shown in **Figure 3** (main text).

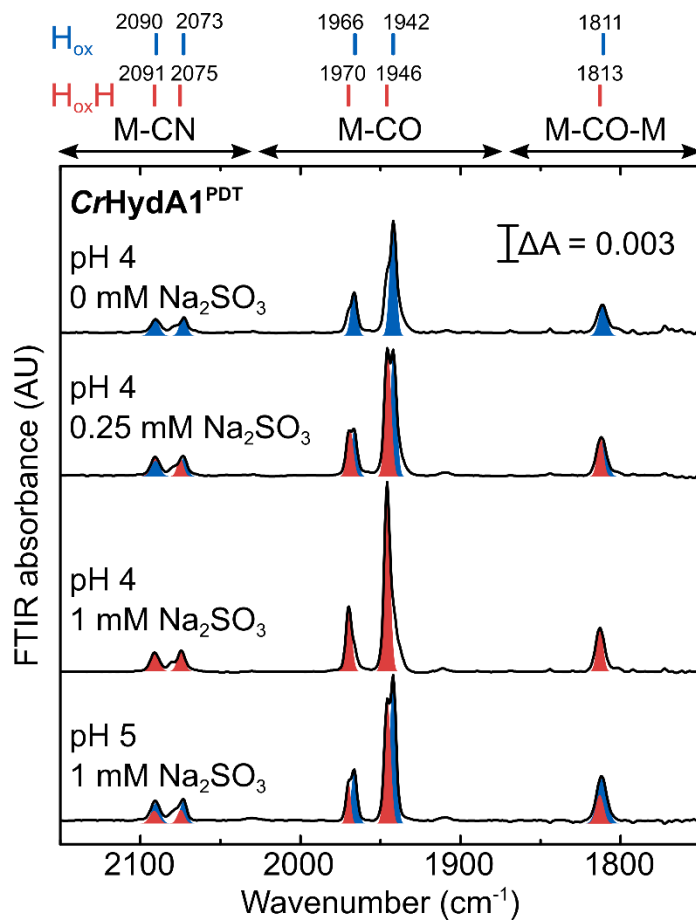


Figure S5 – Titration of CrHydA1^{PDT} with Na₂SO₃ at pH 4 and pH 5 under N₂. IR spectra of CrHydA1^{PDT} ($\approx 420 \mu\text{M}$) diluted in 20 mM mixed buffer (see experimental section) at pH 4 and pH 5 with various concentrations of Na₂SO₃.

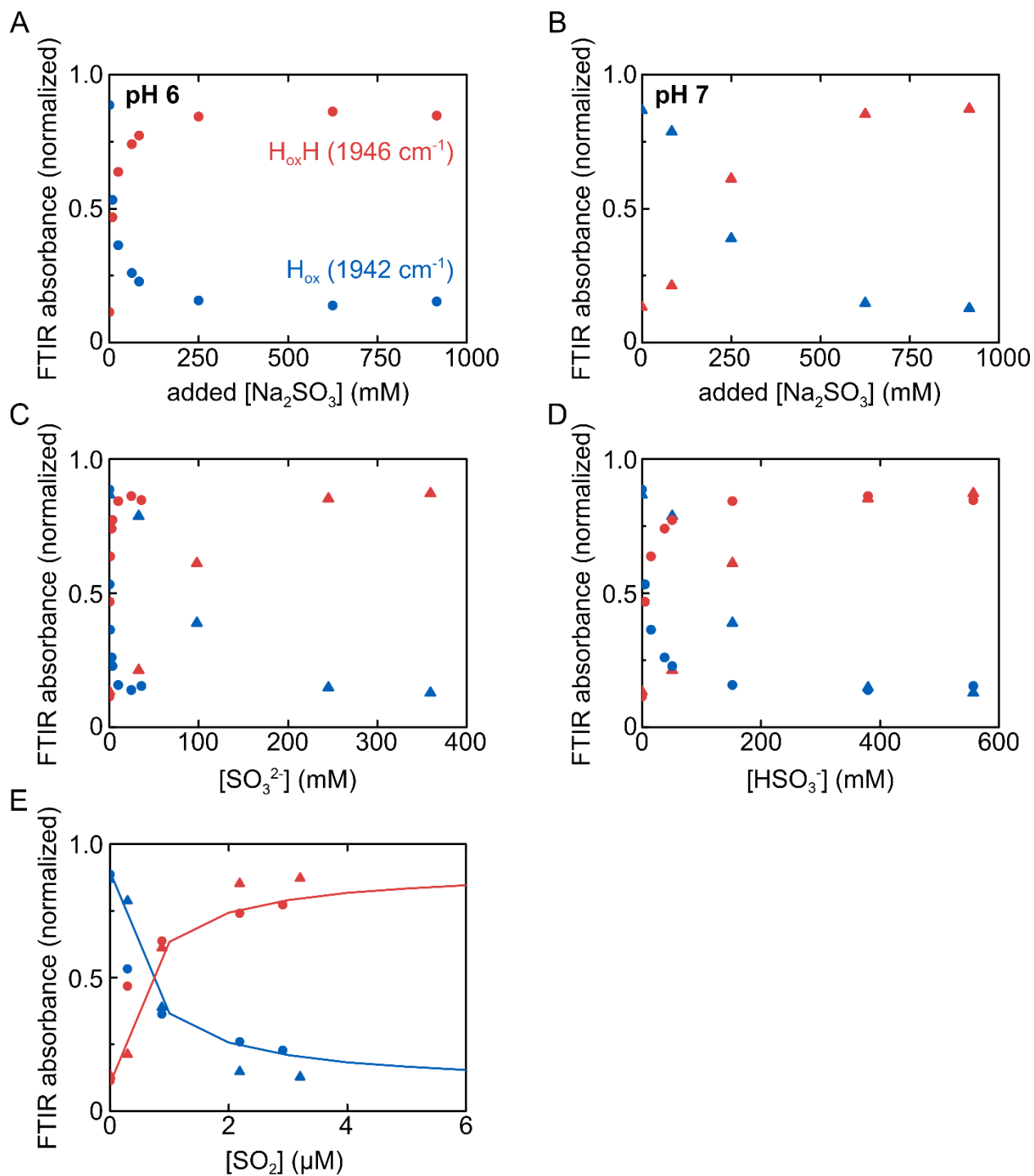


Figure S6 – Titration curves of *CrHydA1*^{PDT}. **A, B** Variation in the intensity of the 1942 cm^{-1} (H_{ox}) and 1946 cm^{-1} ($\text{H}_{\text{ox}}\text{H}$) peaks with the concentration of added Na_2SO_3 at pH 6 (**A**) and pH 7 (**B**). **C, D** as in **A** and **B**, but the data (triangles for pH 7, circles for pH 6) are plotted as a function of the estimated concentration of SO_3^{2-} (**C**) and HSO_3^- (**D**). **E** Expanded version of the region from 0 to 6 μM SO_2 in **Figure 3C** (main text).

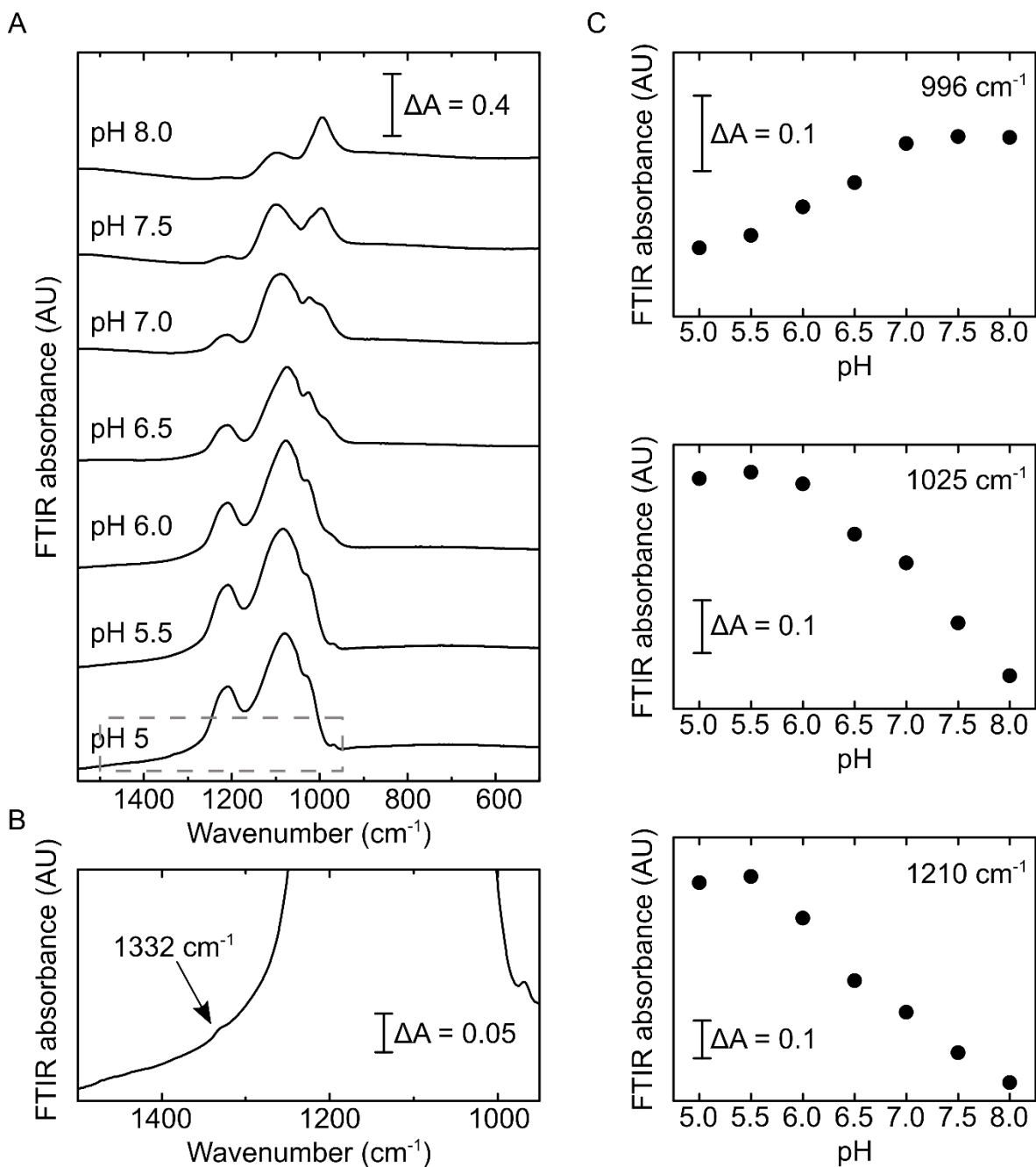


Figure S7 – Speciation of SO_3^{2-} as a function of pH monitored by IR spectroscopy. **A)** Spectra of Na_2SO_3 (1 M) at different pH values. **B)** Enlargement of the pH 5 spectrum (dashed rectangle in **A**), indicating the presence of dissolved SO_2 (ν_3 band at 1332 cm^{-1} , ref²¹). **C)** Titration of the features at 996 cm^{-1} , 1025 cm^{-1} , 1210 cm^{-1} with pH. The feature at 996 cm^{-1} was assigned to SO_3^{2-} , while the ones at 1025 cm^{-1} , 1210 cm^{-1} to HSO_3^- or one of its dimers as reported in ref²¹.

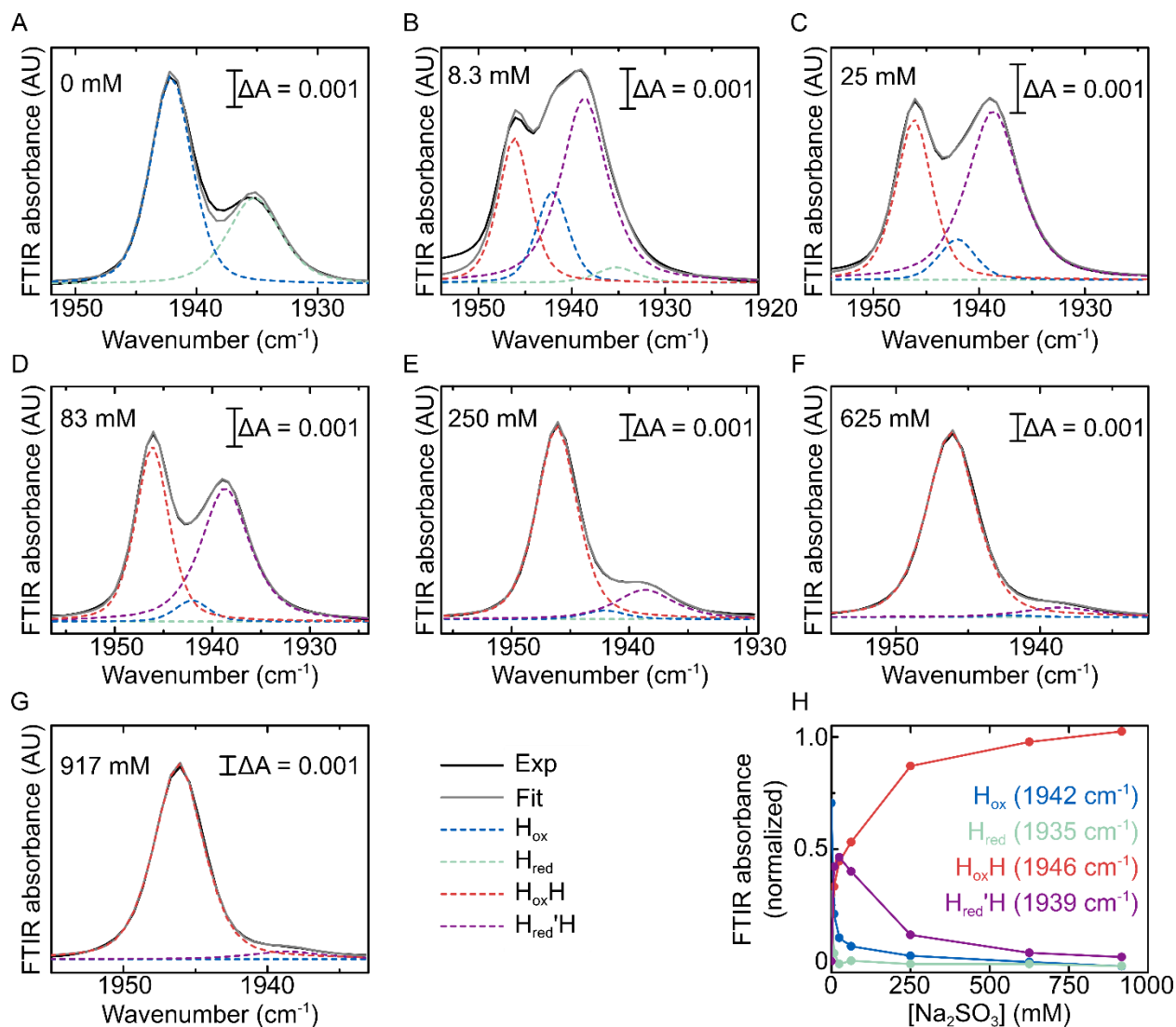


Figure S8 – Full titration of *CrHydA1*^{PDT} with Na₂SO₃ under 2% H₂ at pH 6. A-G) Pseudo-Voigt peak-fitting analysis of the 1955-1920 cm⁻¹ in IR spectra of *CrHydA1*^{PDT} measured under 2% H₂ at pH 6. **H)** Variation in the intensity of the 1942 cm⁻¹ (H_{ox}), 1935 cm⁻¹ (H_{red}), 1946 cm⁻¹ (H_{ox}H) and 1939 cm⁻¹ (H_{red}'H) peaks from **A-G** with the Na₂SO₃ concentration at pH 6. The lines connecting the points in **H** are for visual purposes only.

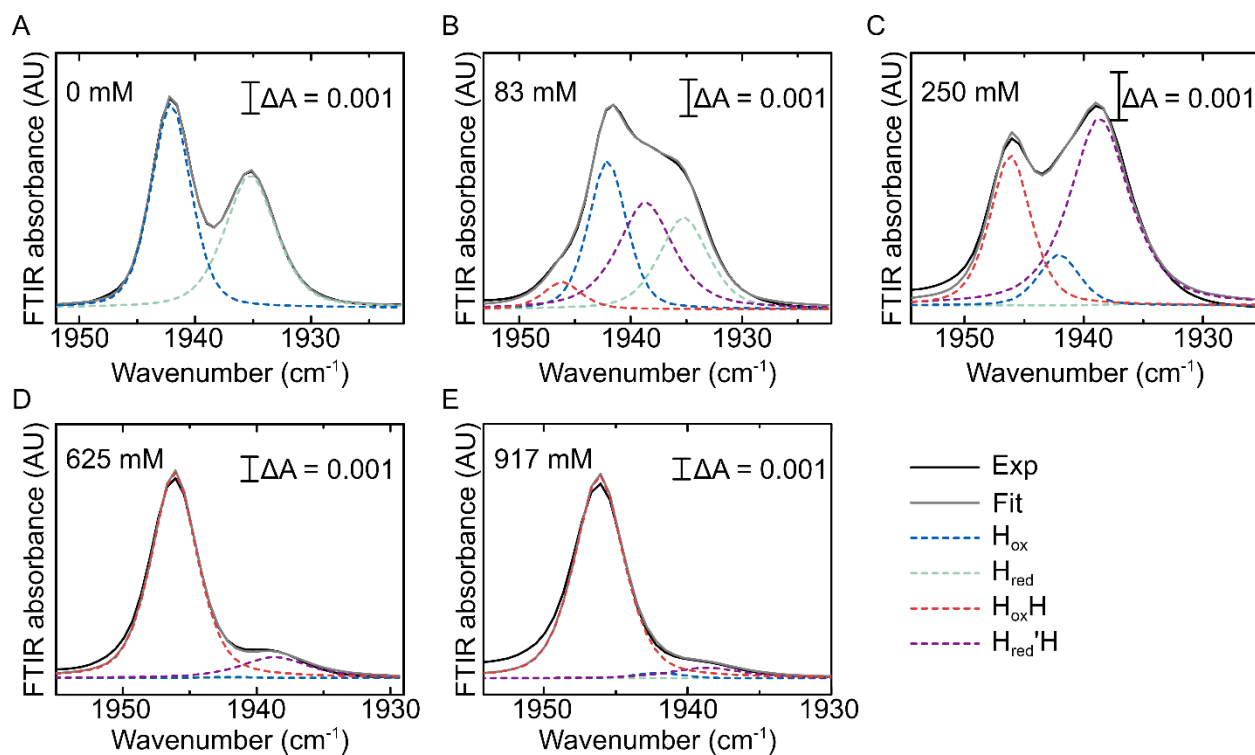


Figure S9 – Full titration of *CrHydA1*^{PDT} with Na₂SO₃ under 2% H₂ at pH 7. A-E) Pseudo-Voigt peak-fitting analysis of the 1955-1920 cm⁻¹ in IR spectra of *CrHydA1*^{PDT} measured under 2% H₂ at pH 7. Data in panel B is also shown in **Figure 4A (main text).**

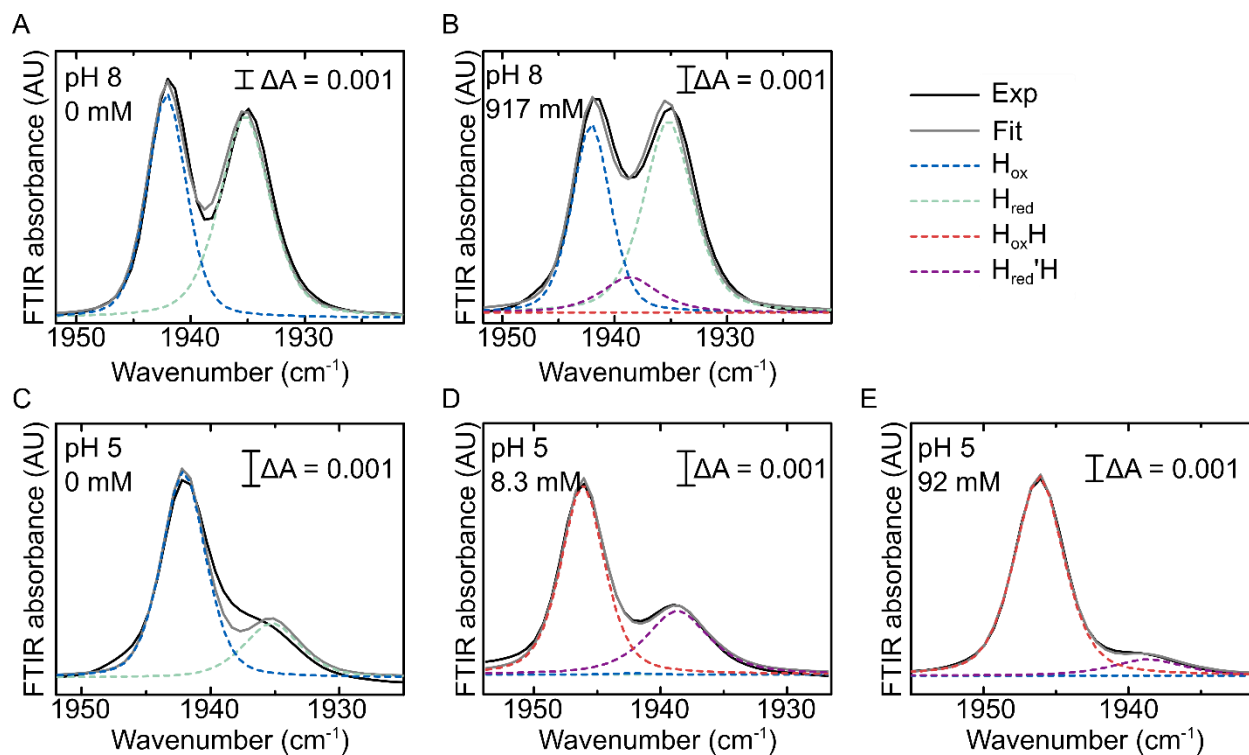


Figure S10 – Full titration of *CrHydA1*^{PDT} with Na₂SO₃ under 2% H₂ at pH 8 and pH 5. Pseudo-Voigt peak-fitting analysis of the 1955-1920 cm⁻¹ in IR spectra of *CrHydA1*^{PDT} measured under 2% H₂ at pH 8 (A, B) and 5 (C, D, E).

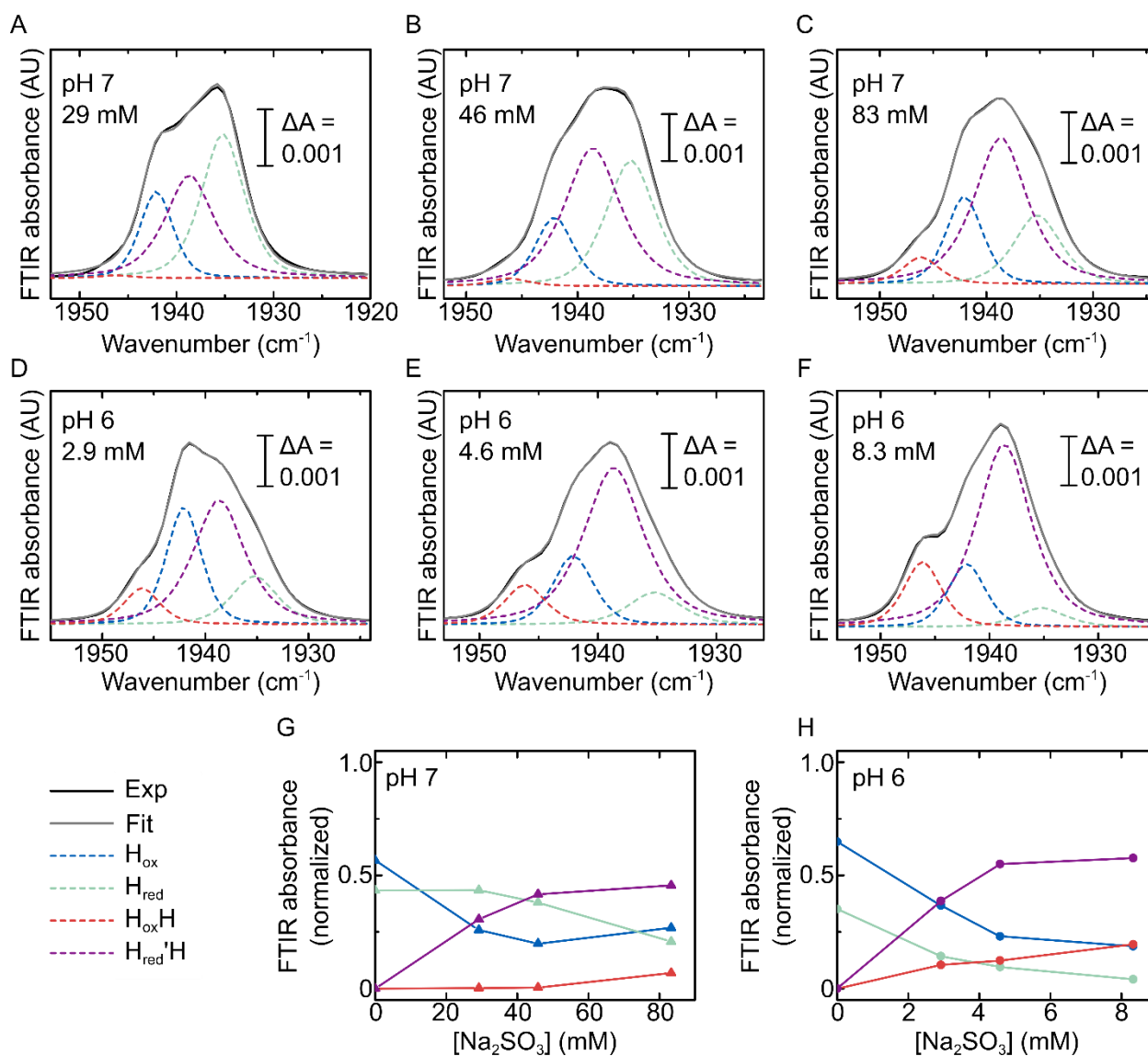


Figure S11 – Calculation of the redox potential of H_{ox}/H_{red} and $H_{ox}H/H_{red}'H$. Pseudo-Voigt peak-fitting analysis of the 1955-1920 cm^{-1} of additional IR spectra of $CrHydA1^{PDT}$ measured under 2% H_2 at pH 7 (A-C) and 6 (D-F) at low concentrations of sulfite. G, H) Variation in the intensity of the 1942 cm^{-1} (H_{ox} , blue), 1935 cm^{-1} (H_{red} , cyan), 1946 cm^{-1} ($H_{ox}H$, red) and 1939 cm^{-1} ($H_{red}'H$, purple) peaks with the Na_2SO_3 concentration at pH 7 (G, data from A-C) and at pH 6 (H, data from D-F). The lines connecting the points in G, H are for visual purposes only. At low Na_2SO_3 concentrations, all four state (H_{ox} , H_{red} , $H_{ox}H$ and $H_{red}'H$) are present at the same time in most cases. At higher concentrations, oxidation of H_2 by the added Na_2SO_3 becomes more prominent, potentially causing an overestimation of the redox potentials. The redox potentials can be calculated for each condition from the H_{ox}/H_{red} and $H_{ox}H/H_{red}'H$ ratios using the Nernst equation and adjusting the $2H^+/H_2$ redox potential according to the pH (7 or 6) – for more details see ref¹⁹. Data in A and B were not used to determine the $H_{ox}H/H_{red}'H$ redox potential, as the $H_{ox}H$ state is not present (see also G). Data from Figure S8B and Figure S9B were included in the calculation of both H_{ox}/H_{red} and $H_{ox}H/H_{red}'H$ redox potentials. The redox potentials were calculated to be $E_m(H_{ox}/H_{red}) = -349 (\pm 17)$ mV and $E_m(H_{ox}H/H_{red}'H) = -293 (\pm 26)$ mV (mean \pm standard deviation, based on the low Na_2SO_3 datapoints at pH 6 and 7).

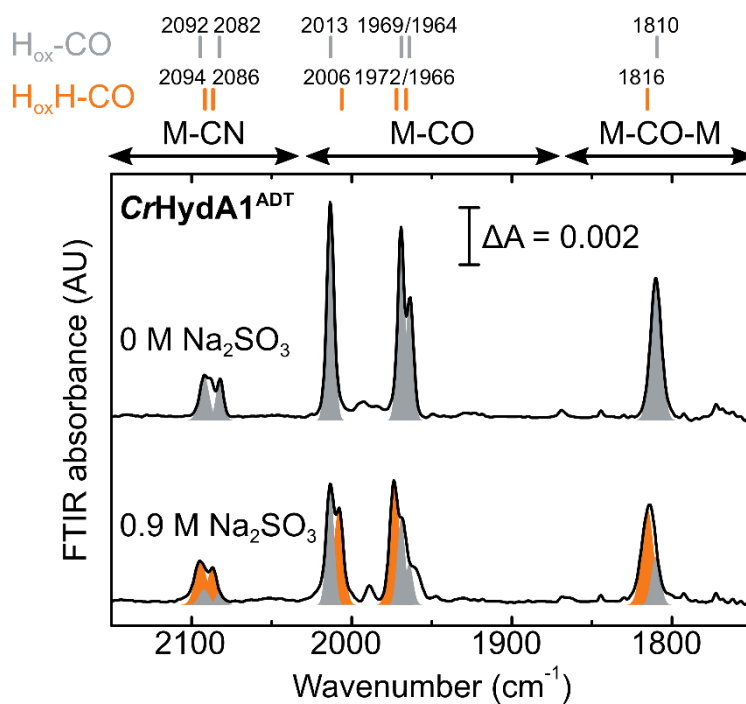


Figure S12 – Formation of the $H_{ox}H-CO$ state. IR spectra of $CrHydA1^{ADT}$ ($\approx 360 \mu M$) diluted in 20 mM mixed buffer at pH 5 (top) or in 0.92 M Na_2SO_3 at pH 5 (bottom) and flushed with 100% CO gas for 10 min before measuring the spectra.

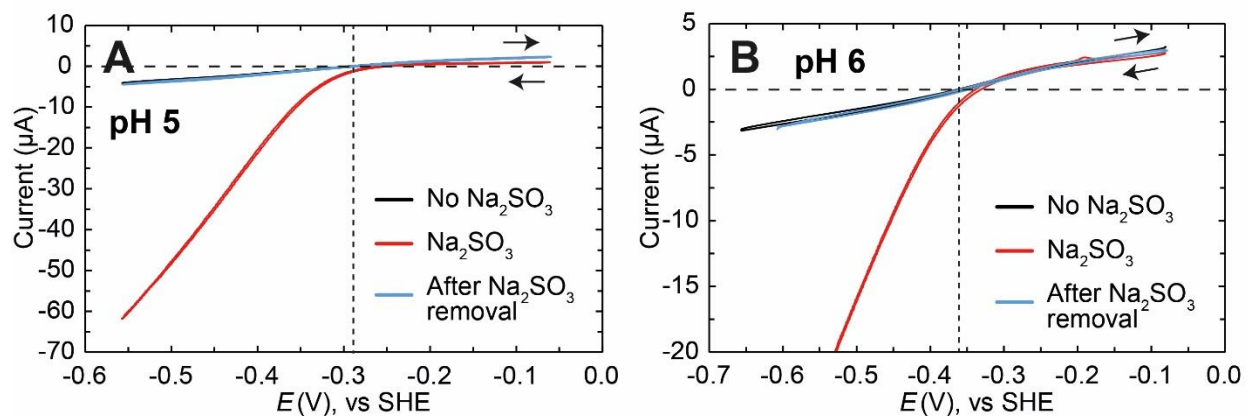


Figure S13 – Cyclic voltammetry of *DdHydAB* in the presence of Na₂SO₃. Enlarged versions of the CVs reported in **Figure 6A** (A, pH 5) and **Figure 6B** (B, pH 6) in the main text. The black arrows indicate the scan direction of the CV. The dashed horizontal line shows the zero current position and the dashed vertical line shows the equilibrium 2H⁺/H₂ potential at each pH value.

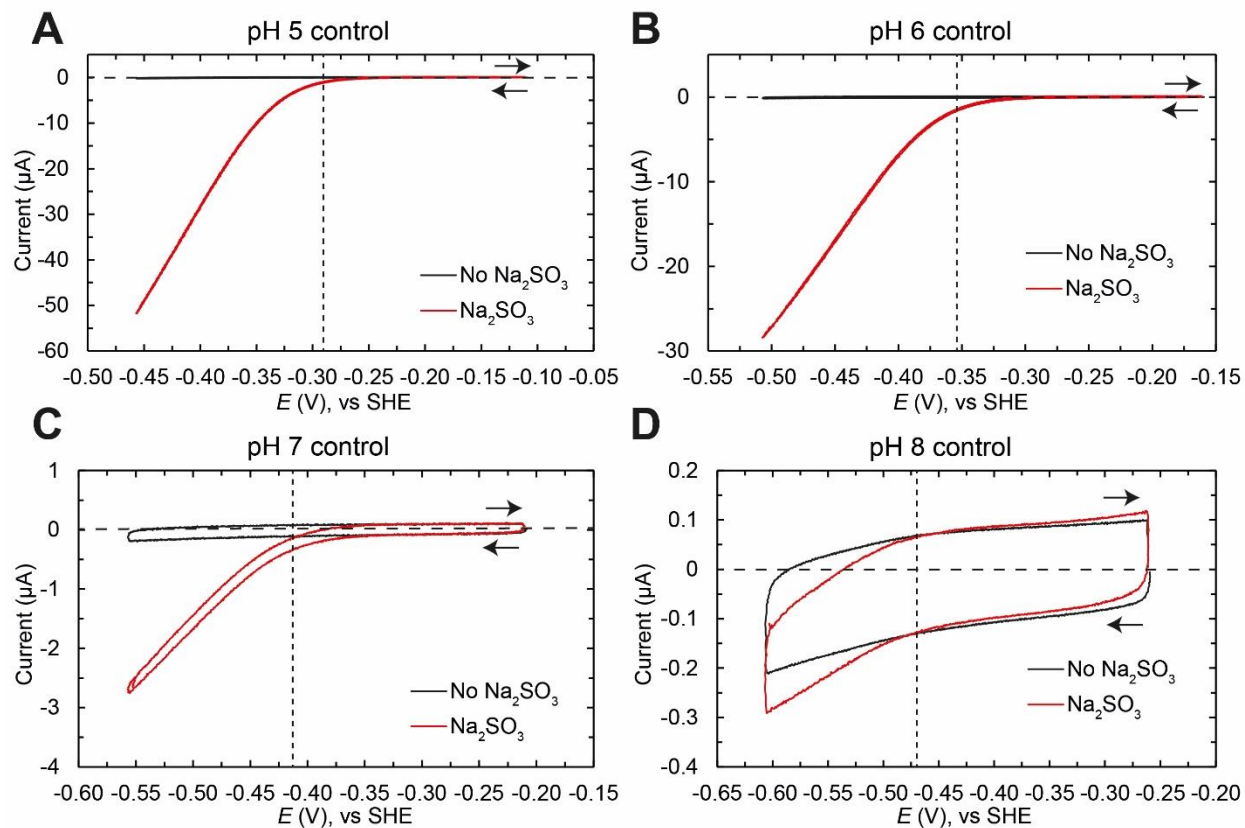


Figure S14 – Cyclic voltammetry of Na_2SO_3 at various pH values. A blank PGE electrode was used for cyclic voltammetry in 20 mM mixed buffer containing 100 mM NaCl at pH 5 (A), 6 (B), 7 (C) and 8 (D) in the absence (black curve) and presence (red curve) of 40 mM Na_2SO_3 . Cyclic voltammetry was measured at 25 °C with 100% H_2 flow and 2000 rpm rotation rate. The black arrows indicate the scan direction of the CV. The dashed horizontal line shows the zero current position and the dashed vertical line shows the equilibrium $2\text{H}^+/\text{H}_2$ potential at each pH value.

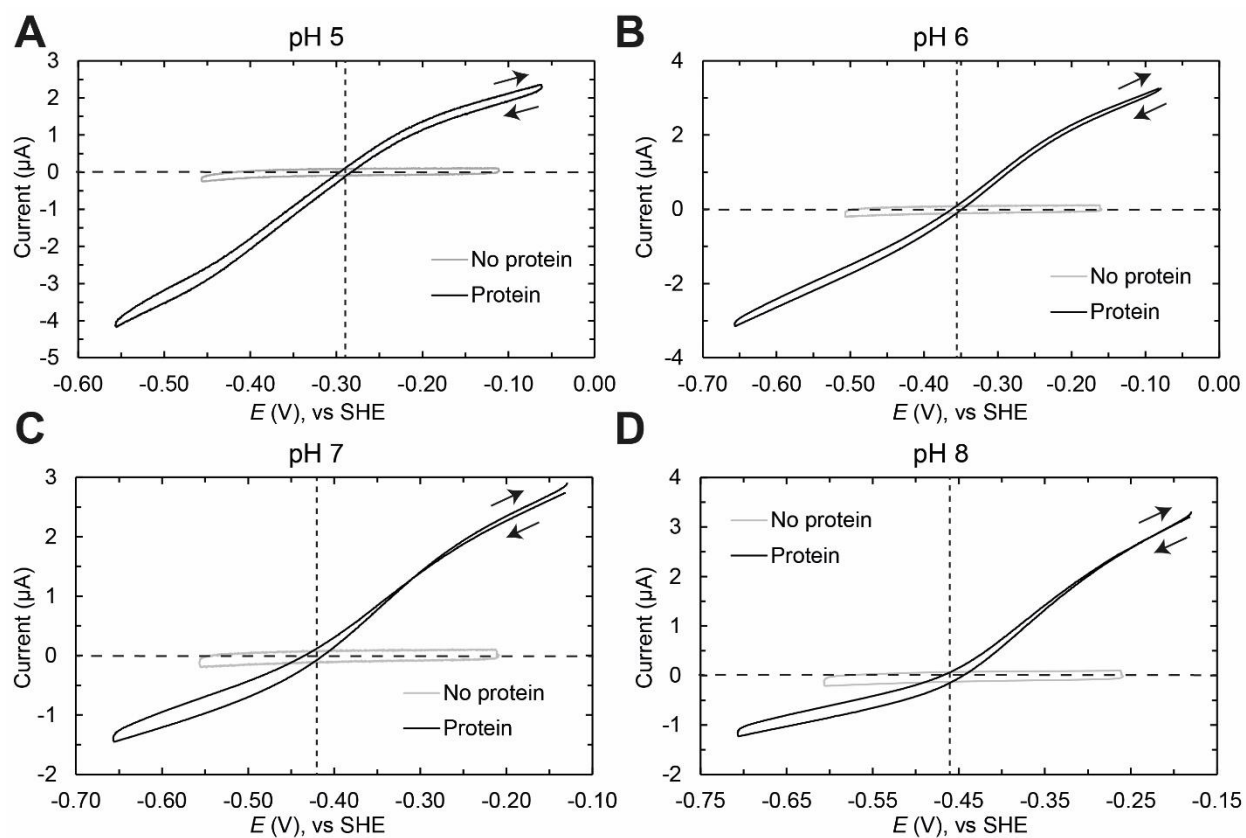


Figure S15 – Cyclic voltammetry of *DdHydAB* at various pH values. A blank (gray curves) PGE electrode and a *DdHydAB* modified PGE electrode (black curves) were used for cyclic voltammetry in 20 mM mixed buffer containing 100 mM KCl at pH 5 (**A**), 6, (**B**), 7 (**C**) and 8 (**D**). Cyclic voltammetry was measured at 25 °C with 100% H_2 flow and 2000 rpm rotation rate. The black arrows indicate the scan direction of the CV. The dashed horizontal line shows the zero current position and the dashed vertical line shows the equilibrium $2H^+/H_2$ potential at each pH value.

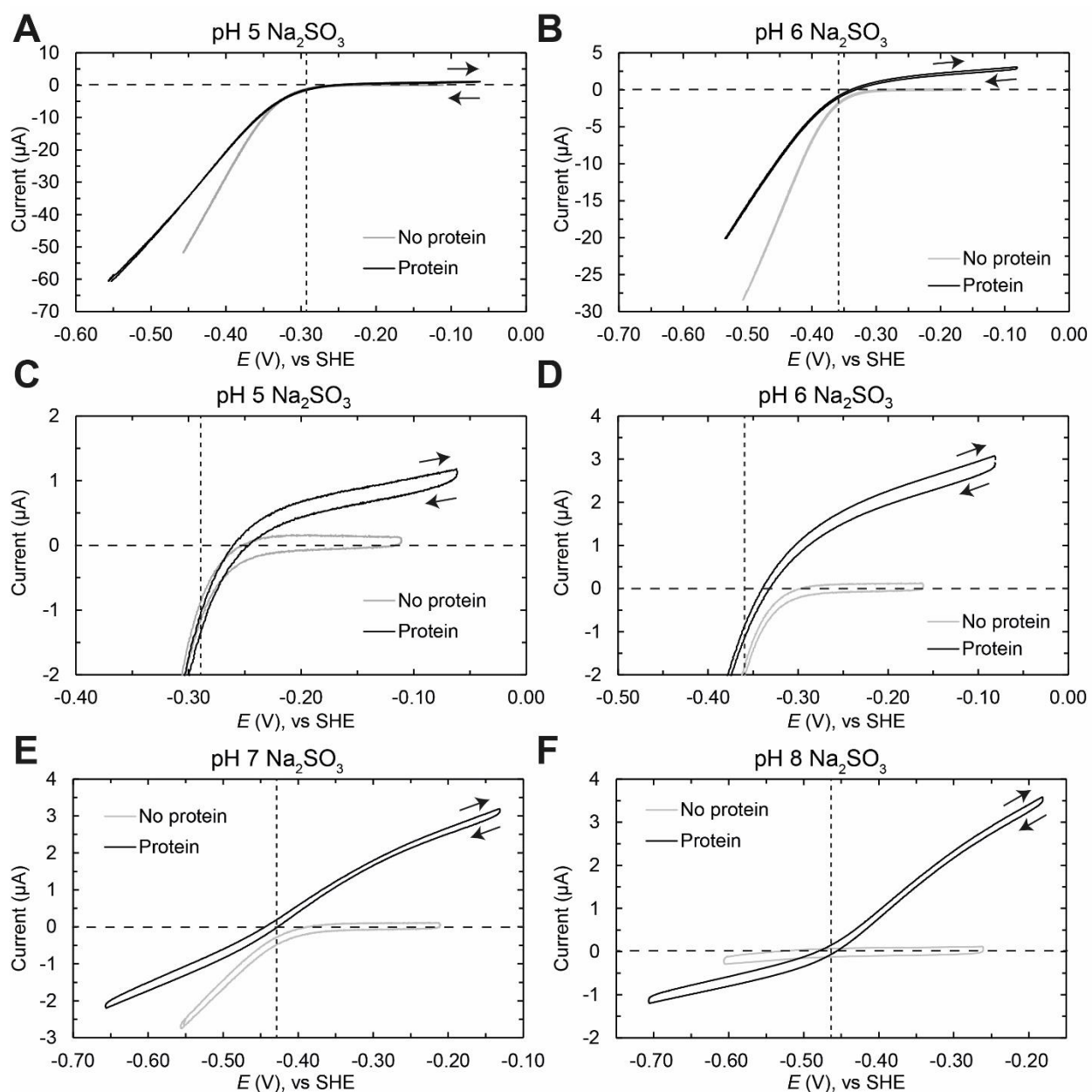


Figure S16 – Cyclic voltammetry of *DdHydAB* at various pH values in the presence of Na_2SO_3 . A blank (gray curves) PGE electrode and a *DdHydAB* modified PGE electrode (black curves) were used for cyclic voltammetry in 20 mM mixed buffer containing 100 mM KCl and 40 mM Na_2SO_3 at pH 5 (A), 6 (B), 7 (E) and 8 (F). Enlarged versions of the H_2 oxidation region of A and B are shown in C and D, respectively. Cyclic voltammetry was measured at 25 °C with 100% H_2 flow and 2000 rpm rotation rate. The black arrows indicate the scan direction of the CV. The dashed horizontal line shows the zero current position and the dashed vertical line shows the equilibrium $2\text{H}^+/\text{H}_2$ potential at each pH value.

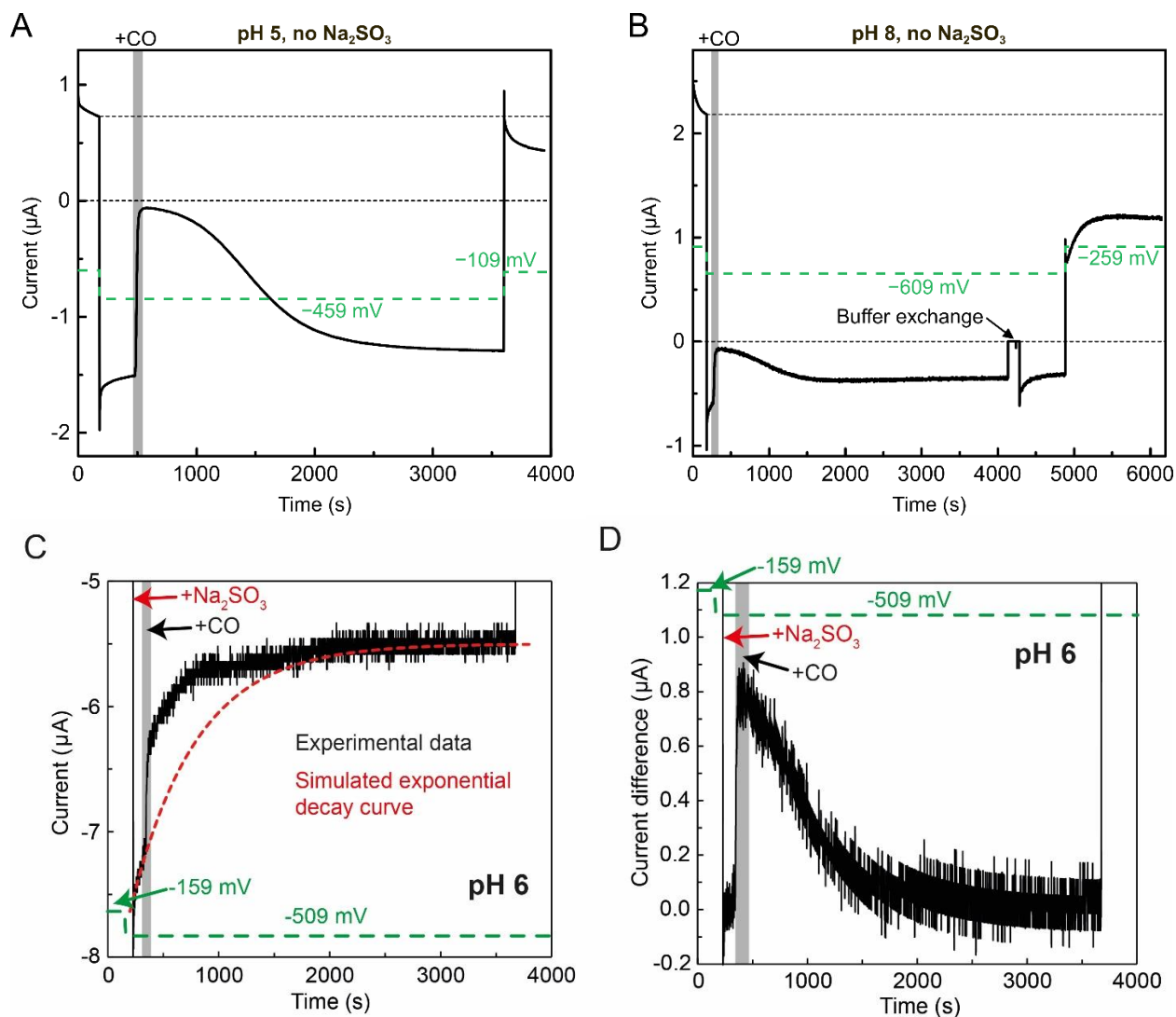


Figure S17 – Chronoamperometry experiments of *DdHydAB* in the absence of Na₂SO₃. A *DdHydAB* modified PGE electrode was used for chronoamperometry experiments in 20 mM mixed buffer containing 100 mM NaCl at pH 5 (**A**) and 8 (**B**) under 1 bar 90 % H₂ in N₂ (1000 ml/min), at 25 °C and with 2000 rpm rotation. The potential was varied as indicated by the green profile, where all potentials are reported vs SHE. At pH 5 (**A**), the potential was set initially to -109 mV, then to -459 mV and finally to -109 mV. At pH 8 (**B**) it was set initially to -259 mV, then to -609 mV and finally to -259 mV. Addition of 10% CO to the gas mixture (replacing 10% N₂) is indicated by the shaded area. In **B** after 4000 s the buffer was exchanged to fresh buffer. (**C**) Expanded region of Figure 7B (pH 6 with Na₂SO₃) from the main text with a simulated exponential decay curve (red dashed line) overlaying the experimental data. (**D**) The current difference between the experimental data and the simulated exponential decay curve from **C** is plotted showing a positive increase in the current after addition of CO, followed by a decay of the difference back to 0. This behavior is analogous to that observed for the experiments in the absence of Na₂SO₃ (**A** and **B**).

Supplementary Discussion: pH dependent redox potentials

As mentioned in the main text, Senger *et al.* observed a pH dependent redox potential for both the H_{ox}/H_{red}' and $H_{ox}H/H_{red}'H$ (their nomenclature) transitions in $CrHydA1^{PDT}$, as well as a pH dependent formation of $H_{ox}H$ from H_{ox} .¹⁰ They attributed this to protonation of the $[4Fe-4S]_H$ subcluster of the H-cluster at two distinct sites – one site was responsible for the pH dependence of the H_{ox}/H_{red} and $H_{ox}/H_{ox}H$ transitions, and the other site was responsible for the $H_{ox}H/H_{red}'H$ and $H_{red}/H_{red}'H$ transitions. However, as we mentioned in a previous report,¹⁹ simply having a proton-dependent step linking H_{ox} and $H_{ox}H$ as well as H_{red}' and $H_{red}'H$ gives rise to apparent pH dependencies for all of the above-mentioned transitions. As such, two protonation sites are not required. We also demonstrated that, in the absence of sodium dithionite, neither $H_{ox}H$ nor $H_{red}'H$ are observed.¹⁹ This behavior is not in agreement with a simple protonation of the $[4Fe-4S]$ cluster.

In the current work, we have shown that the formation of $H_{ox}H$ and $H_{red}'H$ are strictly dependent on Na_2SO_3 , an oxidation product of sodium dithionite. Moreover, these two states only form with Na_2SO_3 at low pH implicating SO_2 as the species that interacts with the H-cluster to form $H_{ox}H$ and $H_{red}'H$. Accordingly, we have renamed these states as H_{ox-DT_i} and H_{red-DT_i} (for dithionite inhibited). With this knowledge at hand we can try and explain the redox titrations published by Senger *et al.* In Figure S18, theoretical redox potential titrations of $CrHydA1^{PDT}$ can be observed for pH 7.5 (Figure S18A), pH 6.5 (Figure S18B) and pH 5.5 (Figure S18C), reflecting the titrations presented in Figure 3 of Senger *et al.*¹⁰ These figures were generated using the model shown in Figure S18D in which H_{ox} and H_{ox-DT_i} undergo a simple one-electron reduction to give H_{red} and H_{red-DT_i} , respectively. Meanwhile, H_{ox} and H_{red} convert to H_{ox-DT_i} and H_{red-DT_i} by simple binding of SO_2 . As the concentration of SO_2 depends on pH, the SO_2 bound states are more abundant in the pH 5.5 titration than in the pH 6.5 and pH 7.5 titrations. Furthermore, as the redox potential for the H_{ox-DT_i}/H_{red-DT_i} transition is more positive than the H_{ox}/H_{red} transition, the apparent redox midpoint potentials of all the titration curves shift with pH, having a value of ≈ -350 mV at pH 7.5, ≈ -320 mV at pH 6.5, and ≈ -300 mV at pH 5.5. Importantly, while the redox potentials for the H_{ox}/H_{red} and H_{ox-DT_i}/H_{red-DT_i} transitions were explicitly defined in the model to be pH-independent, the apparent redox potential for the same transitions appear to be pH-dependent in Figures S18A-C. This is simply the consequence of the coupled equilibria (Figure S18D).

A further curiosity of the results from Senger *et al.* is that, as they titrate the hydrogenase from high potential to low potential at pH 6.5 and pH 5.5, they change from a mixture of H_{ox}/H_{ox-DT_i} to mostly H_{red-DT_i} and in a further step to H_{red} (see Figure 3 in ref.¹⁰), suggesting that H_{red-DT_i} conversion to H_{red} requires reduction. It is important to note that in this experiment the researchers apply an electrochemical potential while flushing gas that has been bubbled through a solution containing NaDT. Our explanation of these results is that at high potentials, NaDT will be oxidized by the electrode generating SO_2 , which can bind to H_{ox} forming H_{ox-DT_i} . As the potential becomes more negative, H_{ox-DT_i} is converted to H_{red-DT_i} , but at the same time NaDT stops being oxidized to SO_2 and the SO_2 that was produced at high potentials is purged from the protein film because of the constant gas flow. This decreases the SO_2 concentration to zero, allowing H_{red-DT_i} to convert back to H_{red} . This behavior is not easily described by a simple thermodynamic model and as such is not observed in Figure S18. A final phenomenon from Senger *et al.* that we cannot yet explain is that they measure the redox potential for H_{ox}/H_{red} at pH 7.5 to be ≈ -450 mV. Multiple publications from our laboratory have estimated this redox potential to be ≈ -350 mV using both spectroelectrochemical IR titrations^{19, 22} and pH titrations under H_2 ¹⁹ (also in this study).

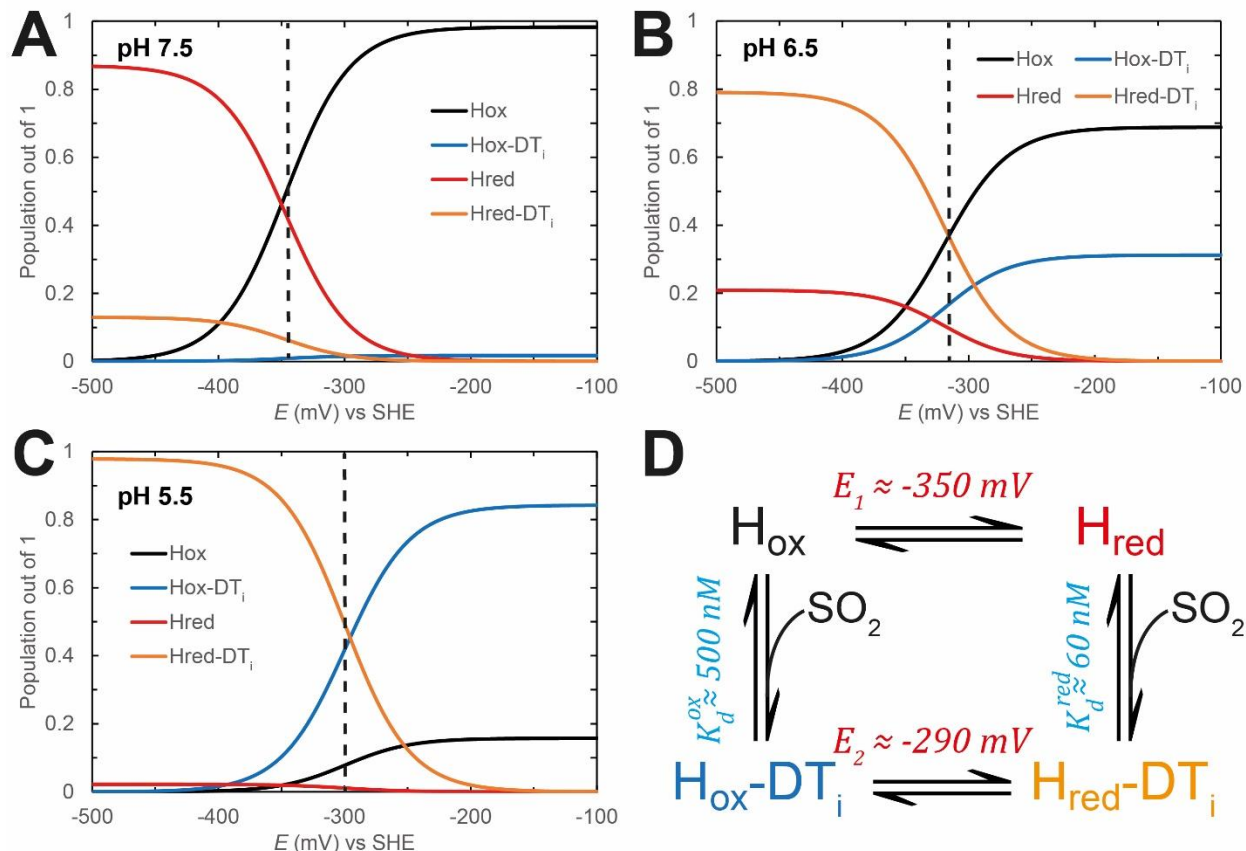


Figure S18 – Model for pH dependent redox potential titrations of $CrHydA1^{PDT}$. The variation in the populations of the H_{ox} , H_{red} , H_{ox-DT_i} and H_{red-DT_i} states (out of a total of 1) with the applied potential at pH 7.5 (**A**), 6.5 (**B**) and 5.5 (**C**) are shown based on equations derived using the model shown in **D**. H_{ox} and H_{red} are connected by a one electron reduction with a redox potential (E_1) of -350 mV, H_{ox-DT_i} and H_{red-DT_i} are connected by a one electron reduction with a redox potential (E_2) of -290 mV, H_{ox} and H_{ox-DT_i} are connected by SO_2 binding with a dissociation constant (K_d^{ox}) of 500 nM, and H_{red} and H_{red-DT_i} are connected by SO_2 binding with a dissociation constant (K_d^{red}) of 60 nM (this work). The SO_2 concentration was calculated using the pH-dependent speciation of Na_2SO_3 based on a pK_a value for $SO_3^{2-} + H^+ \rightleftharpoons HSO_3^-$ of 7.19 and equilibrium constant for $HSO_3^- + H^+ \rightleftharpoons SO_2 + H_2O$ of 1.79,²³ setting the total concentration of $[Na_2SO_3] = [SO_3^{2-}] + [HSO_3^-] + [SO_2] = 10 \text{ mM}$ (in Senger *et al.*¹⁰, 10 mM NaDT was present in the assay). The dotted lines indicate the midpoint potentials of the titration curves (i.e. the position where each state is at 50% of its maximum population) at each pH value.

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