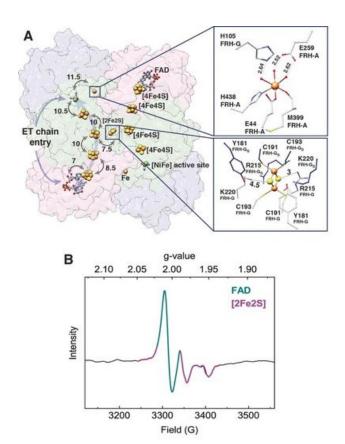


Where hydrogen goes

Start Time: Tuesday, October 29, 2019

End Time:



The UniSysCat groups of <u>Holger Dobbek</u> and <u>Ingo Zebger</u> present in cooperation with Marius Horch from the University of York and Seigo Shima from the MPI in Marburg crystallographic and vibrational-spectroscopic insights into the unexplored structure of the H₂-binding [NiFe] intermediate.

Using an F420-reducing [NiFe]-hydrogenase from Methanosarcina barkeri as a model enzyme, they show that the protein backbone provides a strained chelating scaffold which tunes the [NiFe] active site for efficient H_2 binding and conversion.

The protein matrix also directs H_2 diffusion to the [NiFe] site via two gas channels and allows the distribution of electrons between functional protomers through a subunit-bridging FeS cluster.

Their findings emphasize the relevance of an atypical Ni coordination, thereby providing a



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blueprint for the design of bio-inspired $\rm H_2\mathchar`-conversion$ catalysts.

To find out more <u>click here</u> (Wiley Online Library).



