

The Mononuclear Iron Active Site of the Iron-Sulfur-Cluster-Free Hydrogenase (Hmd)

Rudolf K. Thauer

Department of Biochemistry, Max Planck Institute for Terrestrial Microbiology, Marburg, Germany

Abstract: The iron-sulfur-cluster free-hydrogenase (Hmd) from methanogenic archaea catalyzes the reversible reduction of methenyltetrahydromethanopterin (H_4MPT^+) with H_2 to methylenetetrahydromethanopterin (methylene- H_4MPT) in a stereospecific hydride transfer reaction. The homodimeric enzyme is reversibly inhibited by CO and by cyanide. Hmd contains per 38 kDa subunit one mol of iron, which is associated with a light sensitive cofactor that can be extracted. Active enzyme can be reconstituted from apoprotein and the iron-containing cofactor. We report here on IR-, Mössbauer- and XA spectra of the holoenzyme and of the isolated cofactor. The results indicate that the active site of Hmd harbors a mononuclear Fe center in a low spin and low oxidation state. The iron is complexed by 2 CO, 1 S and 1 N/O and has an open coordination site for the binding of extrinsic CO or of cyanide. Interaction of H_2 with the iron center is dependent on the presence of methenyl- H_4MPT^+ as revealed by IR spectroscopy, which is consistent with the finding that Hmd catalyzes an H_2/H^+ exchange or the conversion of *para* H_2 into *ortho* H_2 only in the presence of the carbocation.

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