

Stepwise assembly of the [NiFe]-hydrogenase active site

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[NiFe]-hydrogenases catalyze the reversible reaction $\text{H}_2 \rightleftharpoons 2\text{H}^+ + 2\text{e}^-$ under ambient conditions. Their basic module consists of a large subunit, hosting the heterobimetallic $\text{NiFe}(\text{CN})_2(\text{CO})$ cofactor, and a small subunit that carries electron-transferring Fe-S clusters. We have recently shown that the large subunits of the O_2 -tolerant regulatory (RH) and membrane-bound (MBH) hydrogenases of *R. eutropha* (HoxC and HoxG, respectively), when separated from the corresponding small subunits (HoxB and HoxK), exhibit catalytic and spectroscopic properties that differ markedly from those of native enzymes.^{1,2,3} The de-assembly process is reversible, as the RH subunits can be reassembled *in vitro* to yield a fully active enzyme. This provides a new tool for the isolation of native-like [NiFe]-hydrogenases equipped with ⁵⁷Fe exclusively at their catalytic site.⁴

Here, we have gone a step further and elucidated the stepwise assembly of the $\text{NiFe}(\text{CN})_2(\text{CO})$ cofactor in the large subunit prior to small subunit attachment by isolating key maturation intermediates of HoxG. These included the cofactor-free apo-HoxG, a nickel-free version carrying only the $\text{Fe}(\text{CN})_2(\text{CO})$ fragment, a precursor containing all cofactor components but redox-inactive, and the fully mature HoxG. Using biochemical analyses in combination with comprehensive spectroscopic studies, we present detailed insights into the sophisticated maturation process of [NiFe]-hydrogenase.

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