



Hanseatic League of Science
Interconnecting infrastructures
for life science research and innovation



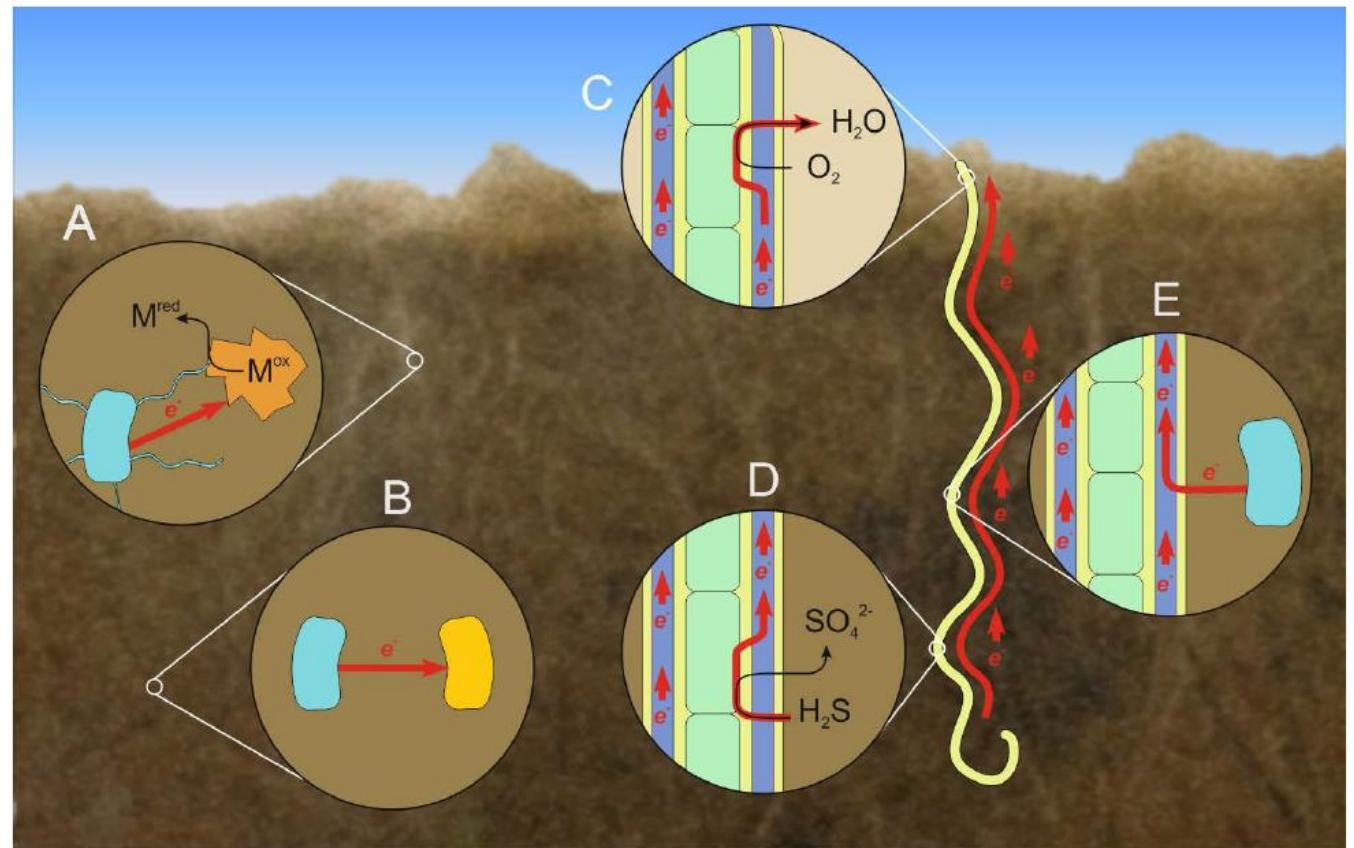
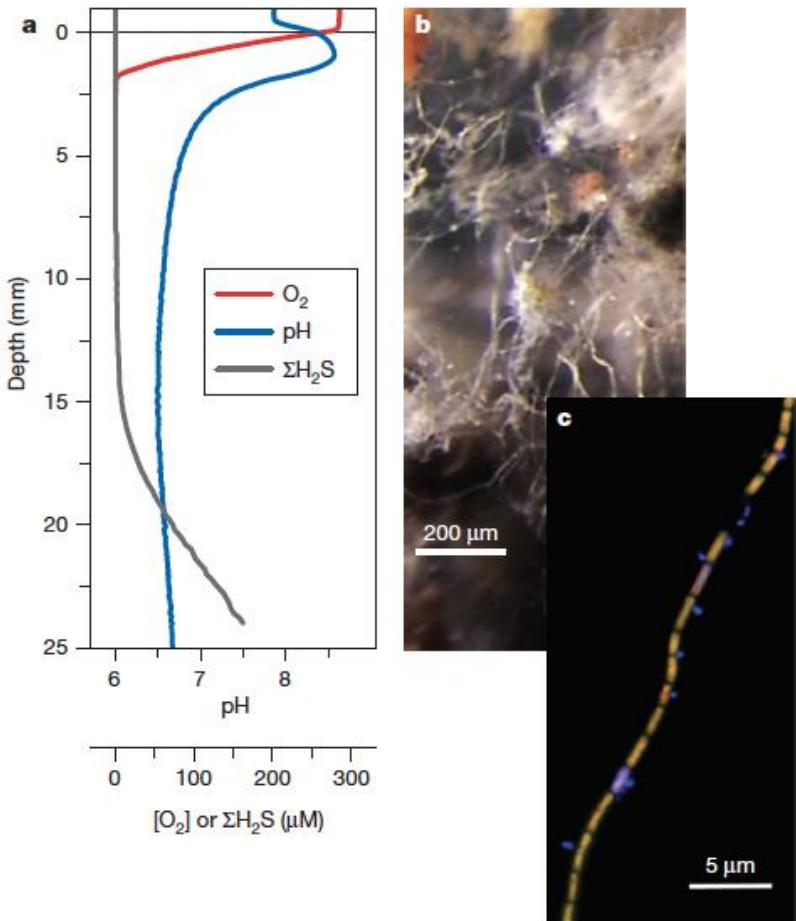
Wired life

insights into unique enzymes from cable bacteria

Thomas Boesen and Kajsa Sigfridsson Clauss

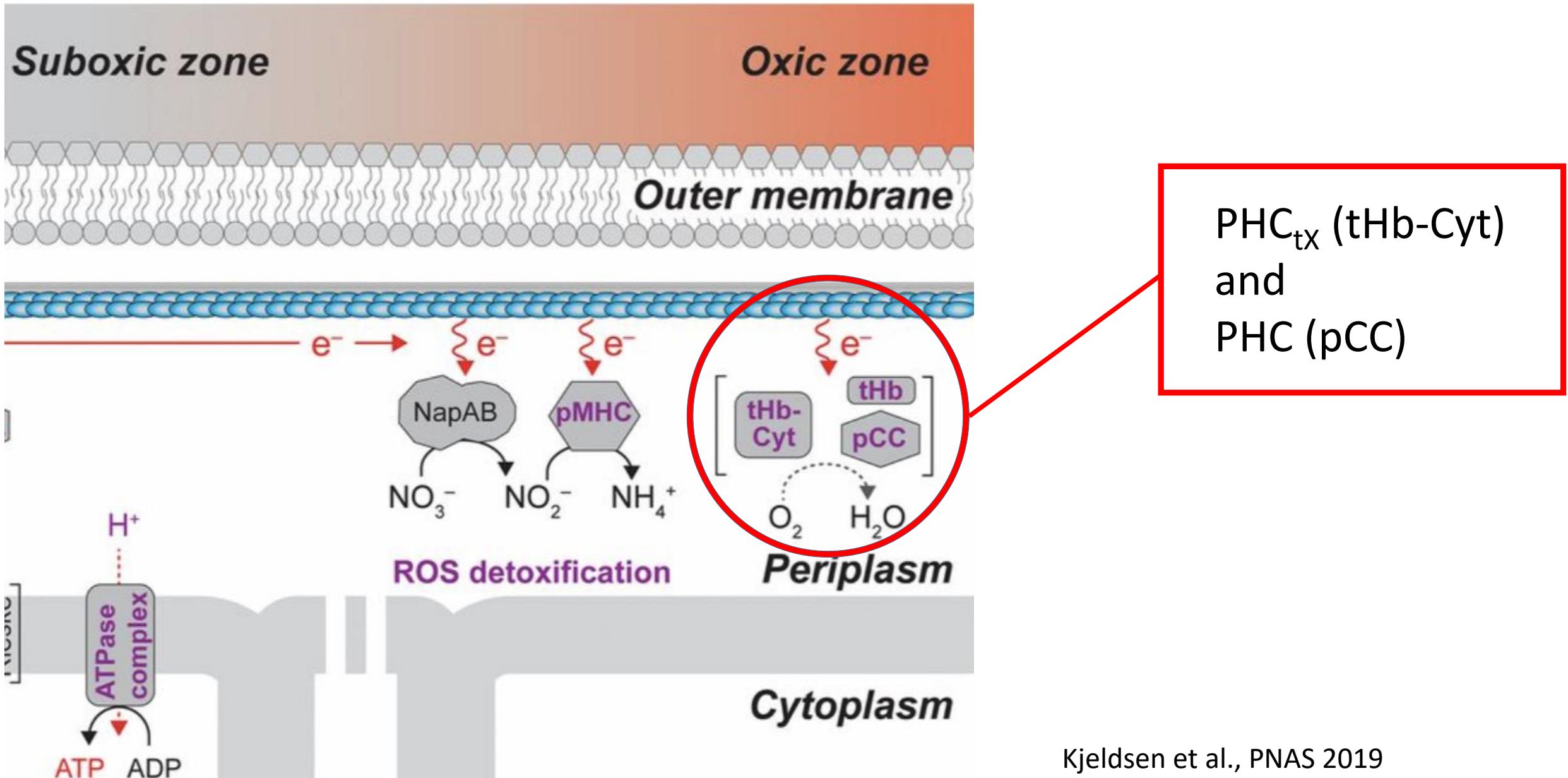


Cable bacteria

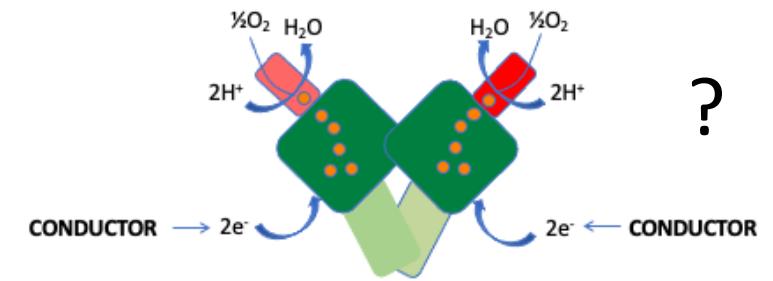
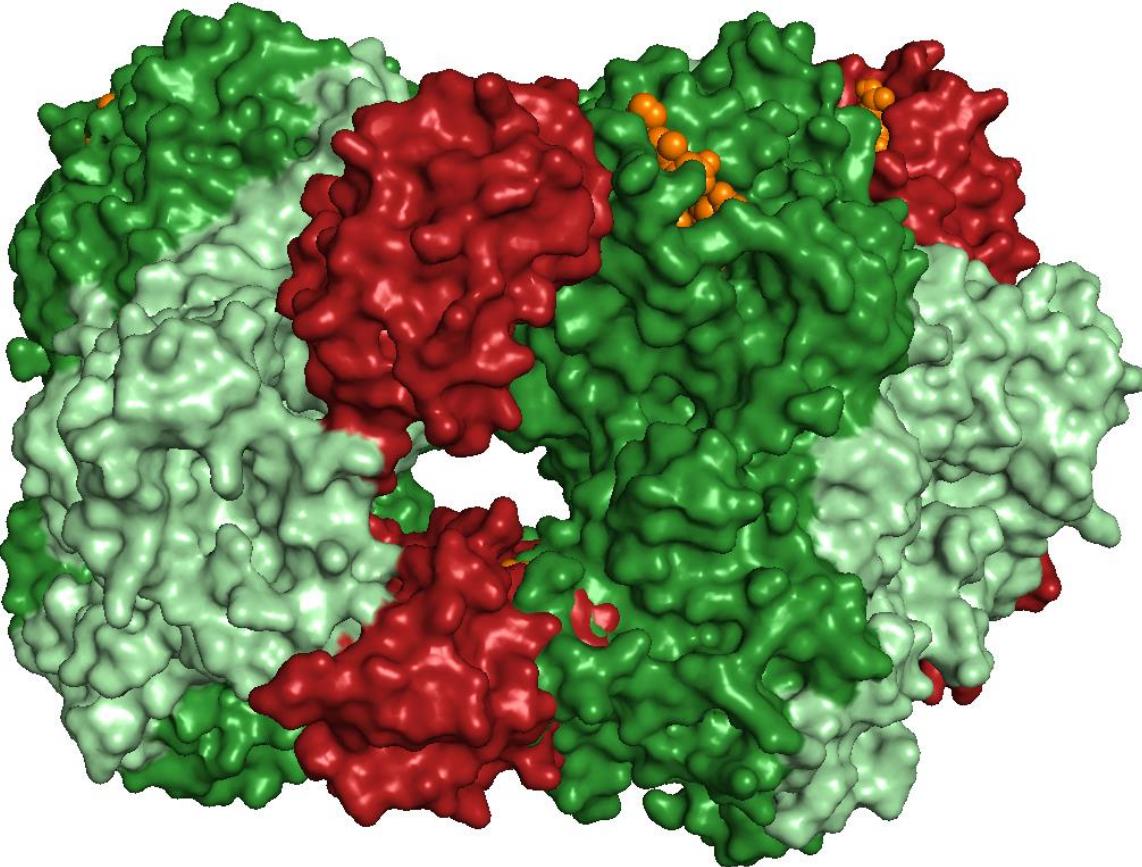
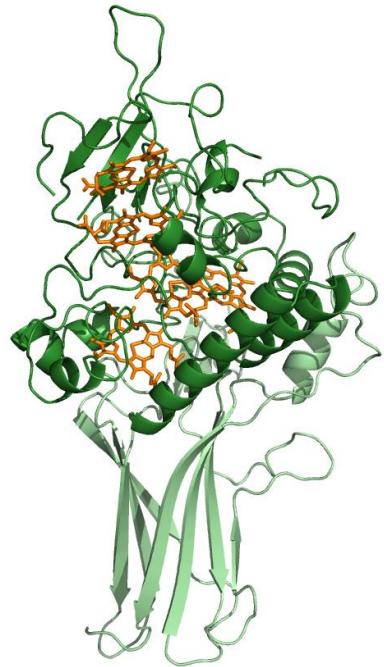
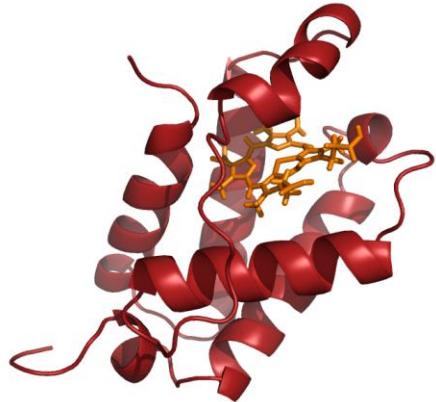


Nielsen *et al*, Nature 2010, Pheffer *et al*, Nature 2012

A metabolic model for electron flow



PHC_{tx}

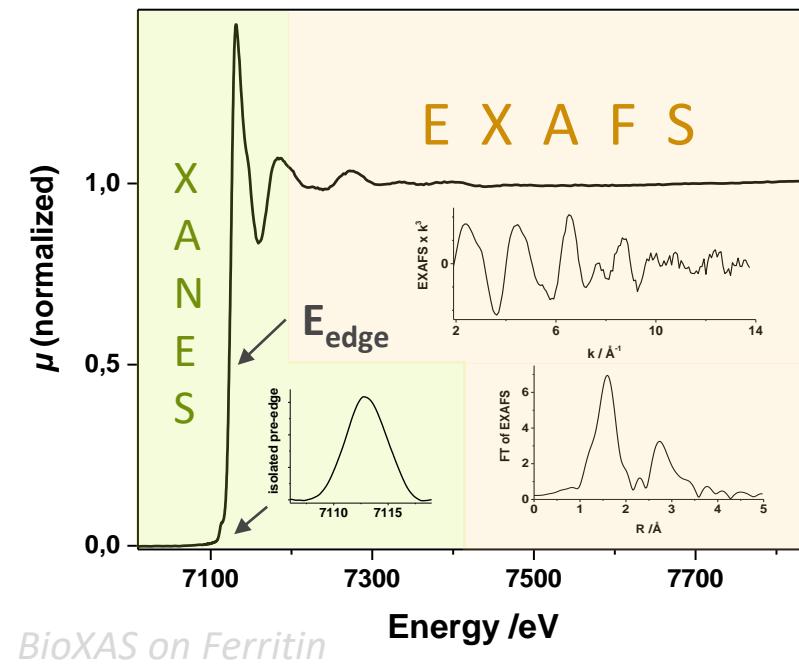


PHC_{tNC}

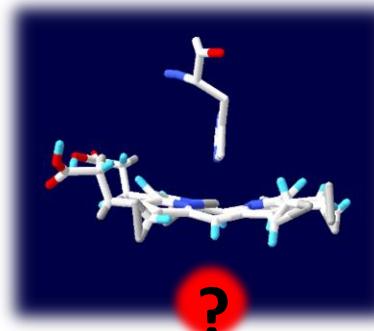
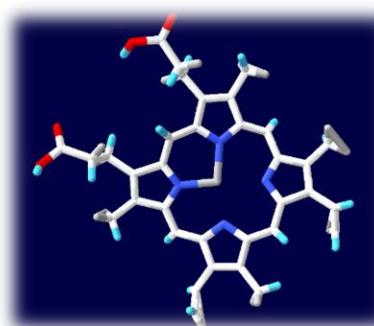
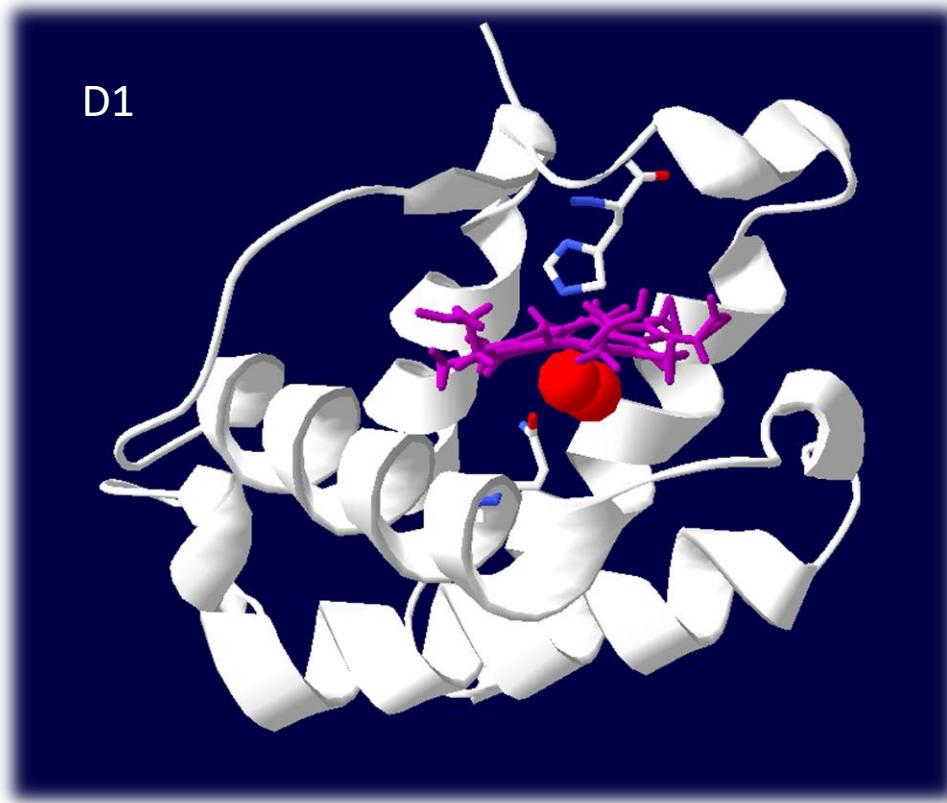
MAX IV /Balder /X-ray Absorption Spectroscopy



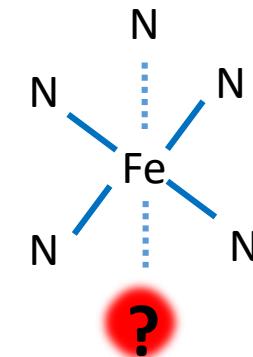
XAS = HEAVY METAL



Metal site structure of hemoglobin domain



O₂, H₂S, H₂O, ?



Fe oxidation state
Geometric structure
=XANES

Local atomic structure around Fe
Identity of neighbor atoms
Distance to neighbor atoms
Number of neighbor atoms

= EXAFS

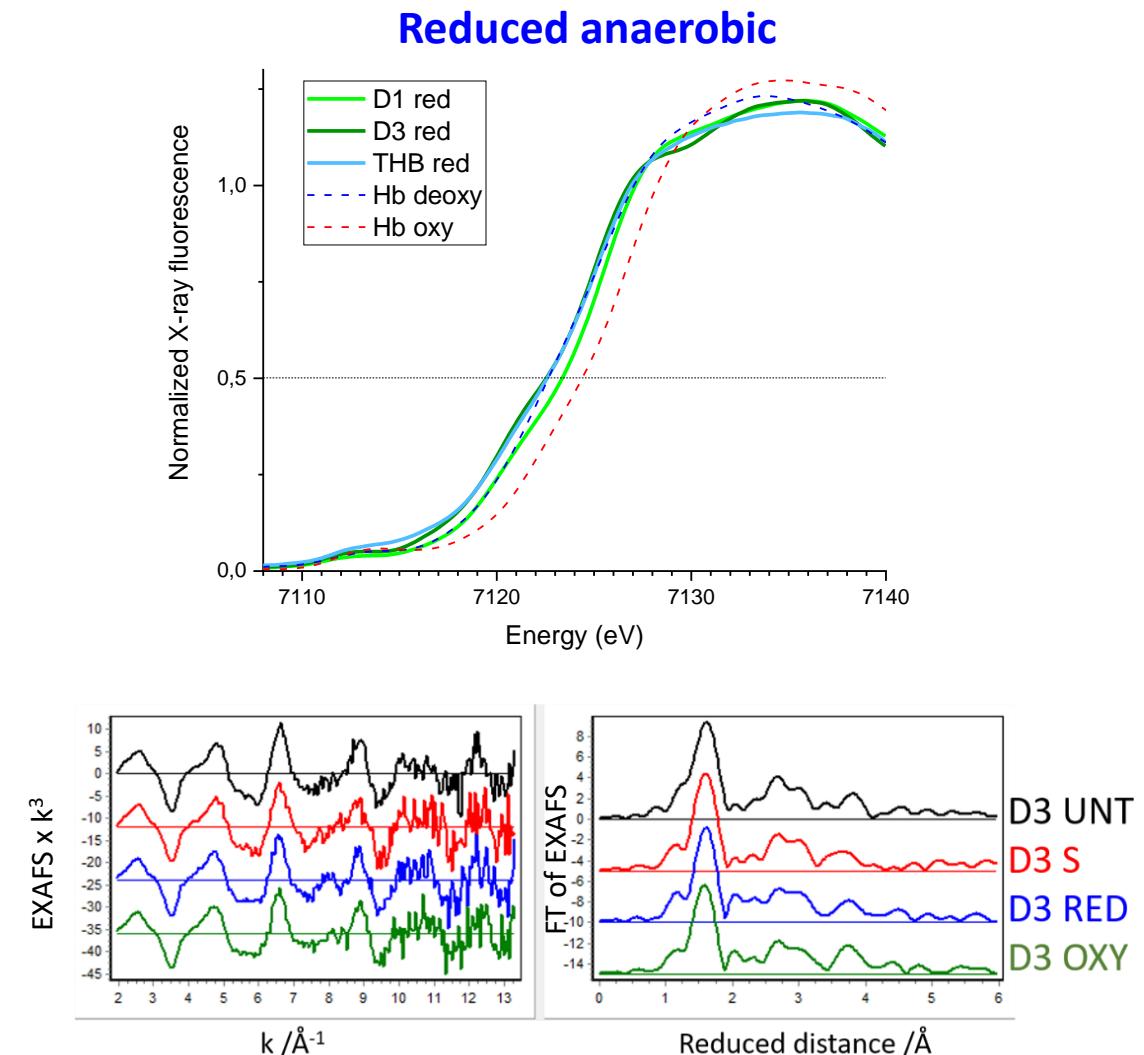
XANES and EXAFS on D1, D3 and THB

Sample environments

- **Cryostat 15 K**
- Microfluidic flowcell
- Electrochemistry flowcell

Different conditions

- Reduced (anaerobic)
- O₂ flushed
- H₂S treated
- Untreated



Next steps

- Next beamtime: pH dependence of the O₂ and H₂S affinity to the hemoglobin domain “the Bohr effect”
- Investigate other heme proteins in the CB electron transfer chain.

Long term goal:

To understand the reaction mechanism from heme to molecule to organism using XAS.