



THE EVOLUTION OF LIFE

The coevolution of the geosphere and biosphere

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Rutgers University

Sagan Workshop - Caltech



e·nig·ma

Evolution of Nanomachines In Geospheres and Microbial Ancestors

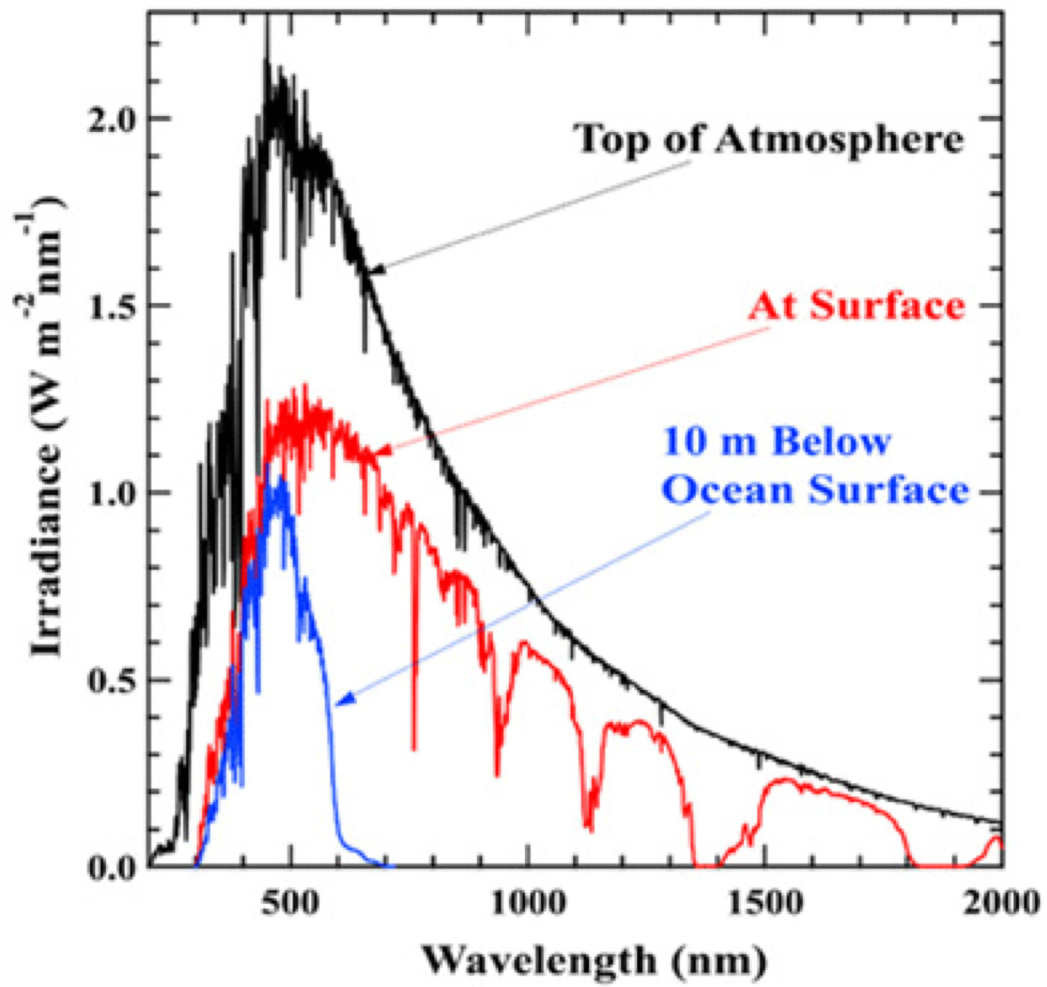
<https://enigma.rutgers.edu>

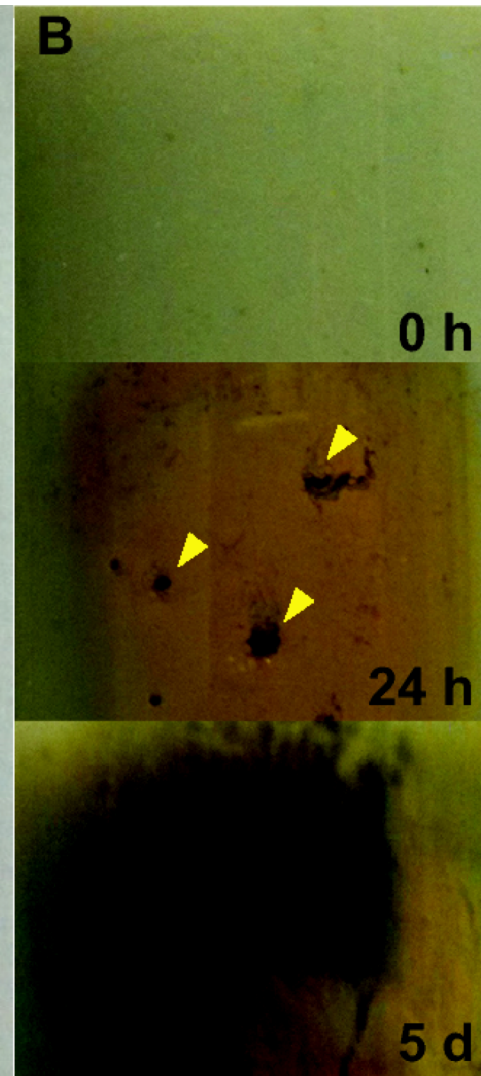
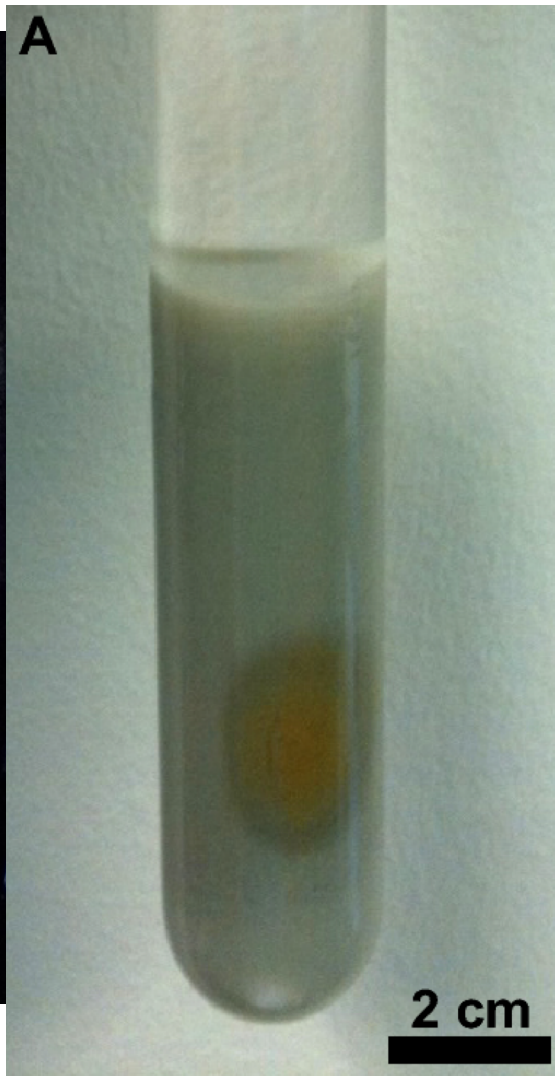
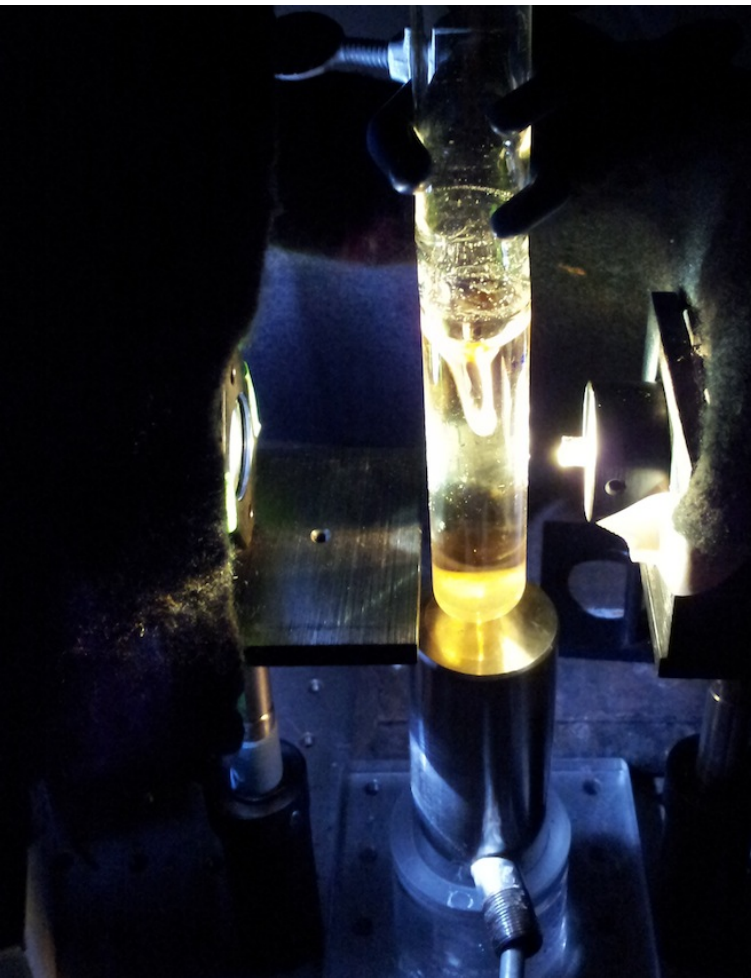


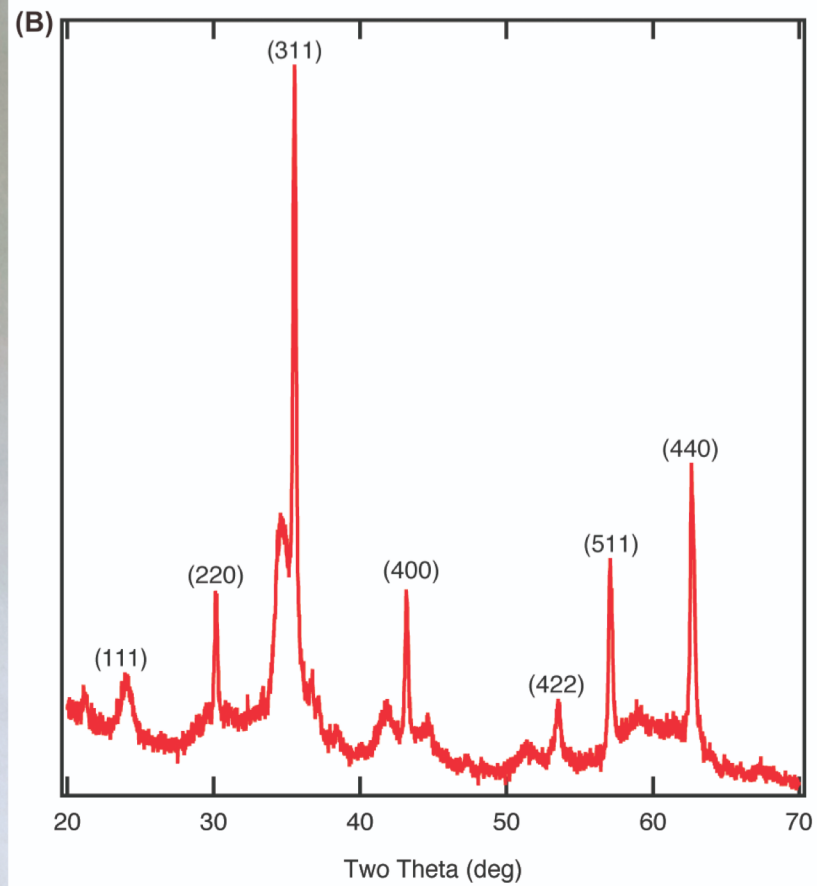
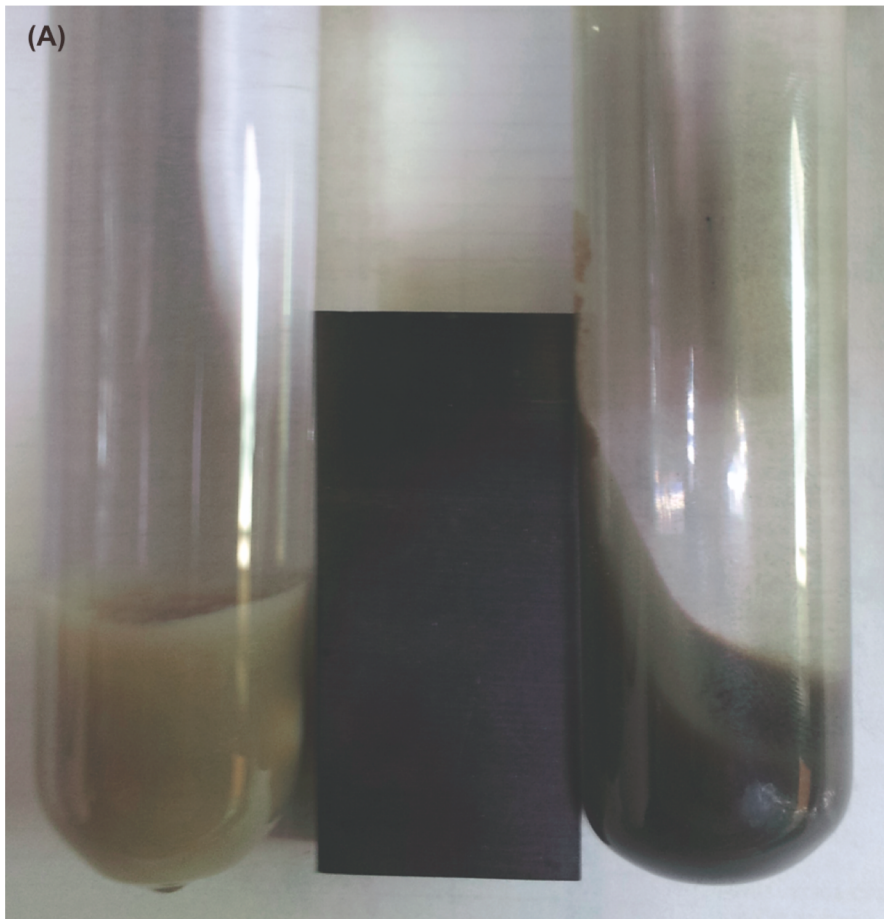
The Enigma research team is part of
the NASA Astrobiology Institute (NAI)











Production of H₂ from the irradiation of MnCO₃ with UV light

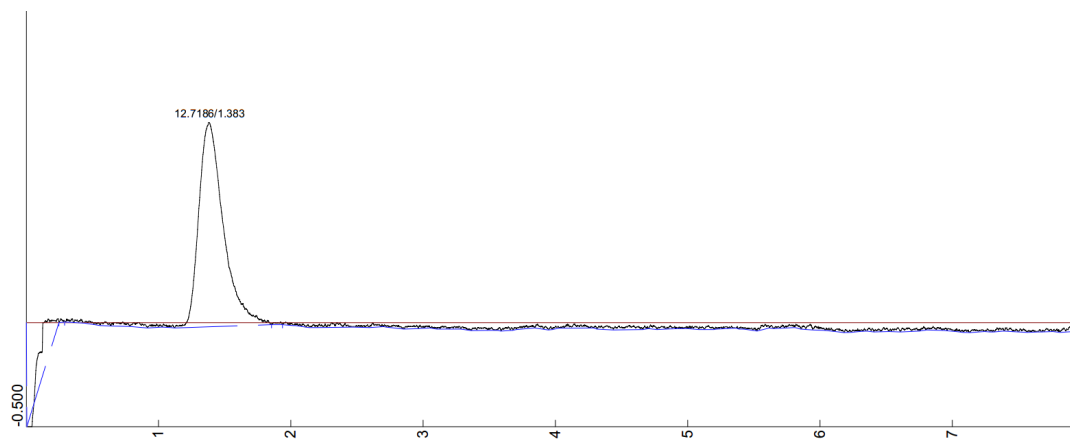
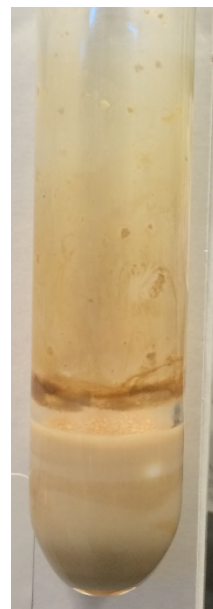
0 hr

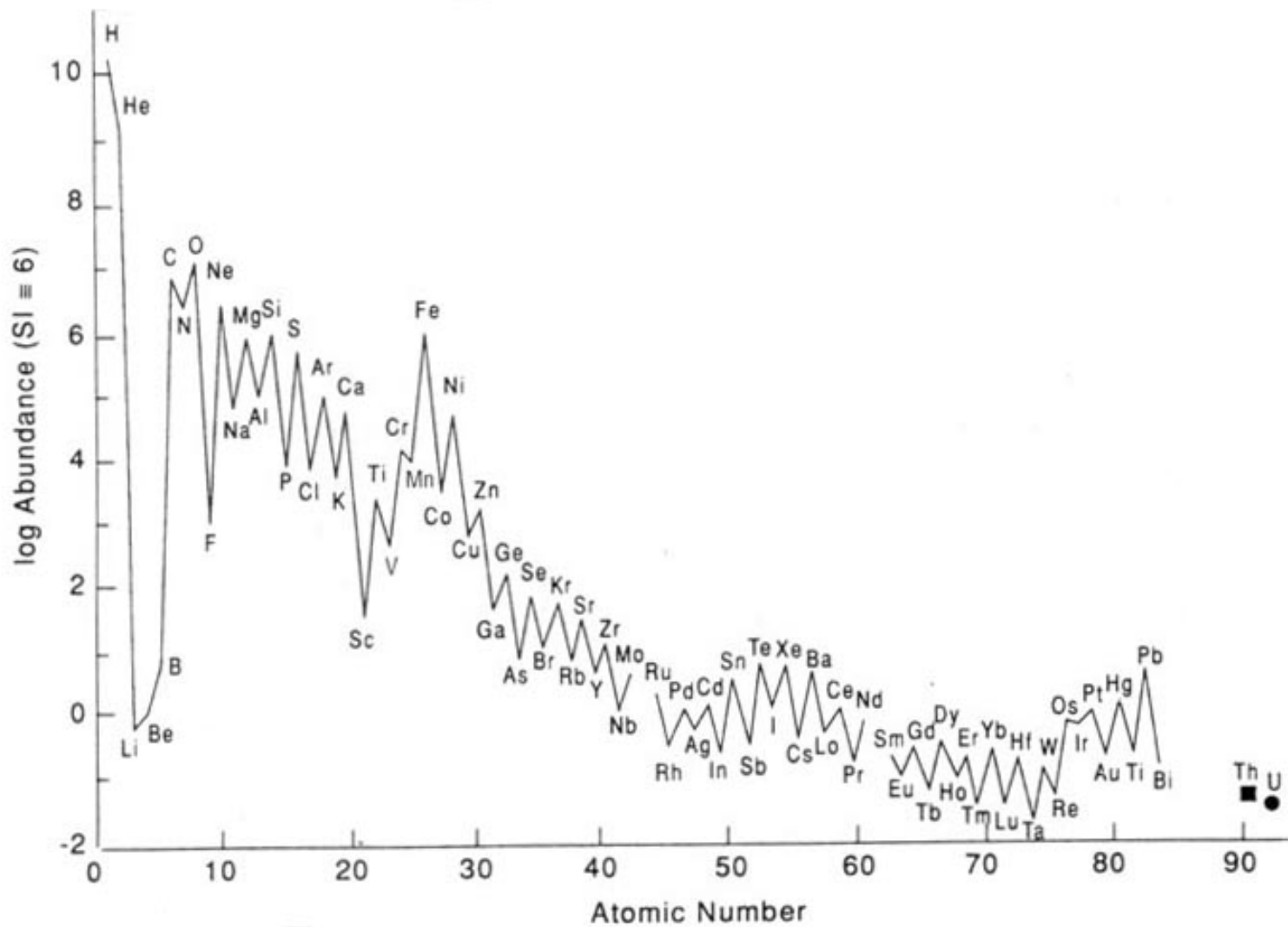


48 hr



380 hr





The origin of life is the invention of
non-equilibrium redox chemistry that
involves five of the
The “Big Six”

H, C, N, O, P and S

**And at least 54 other “trace
elements”**

Life is Electric

- **All organisms derive energy for growth and maintenance by moving electrons from a substrate to a product.**
- **All substrates and products must ultimately be cycled.**
- **Biological processes are paired (e.g., photosynthesis and respiration)**

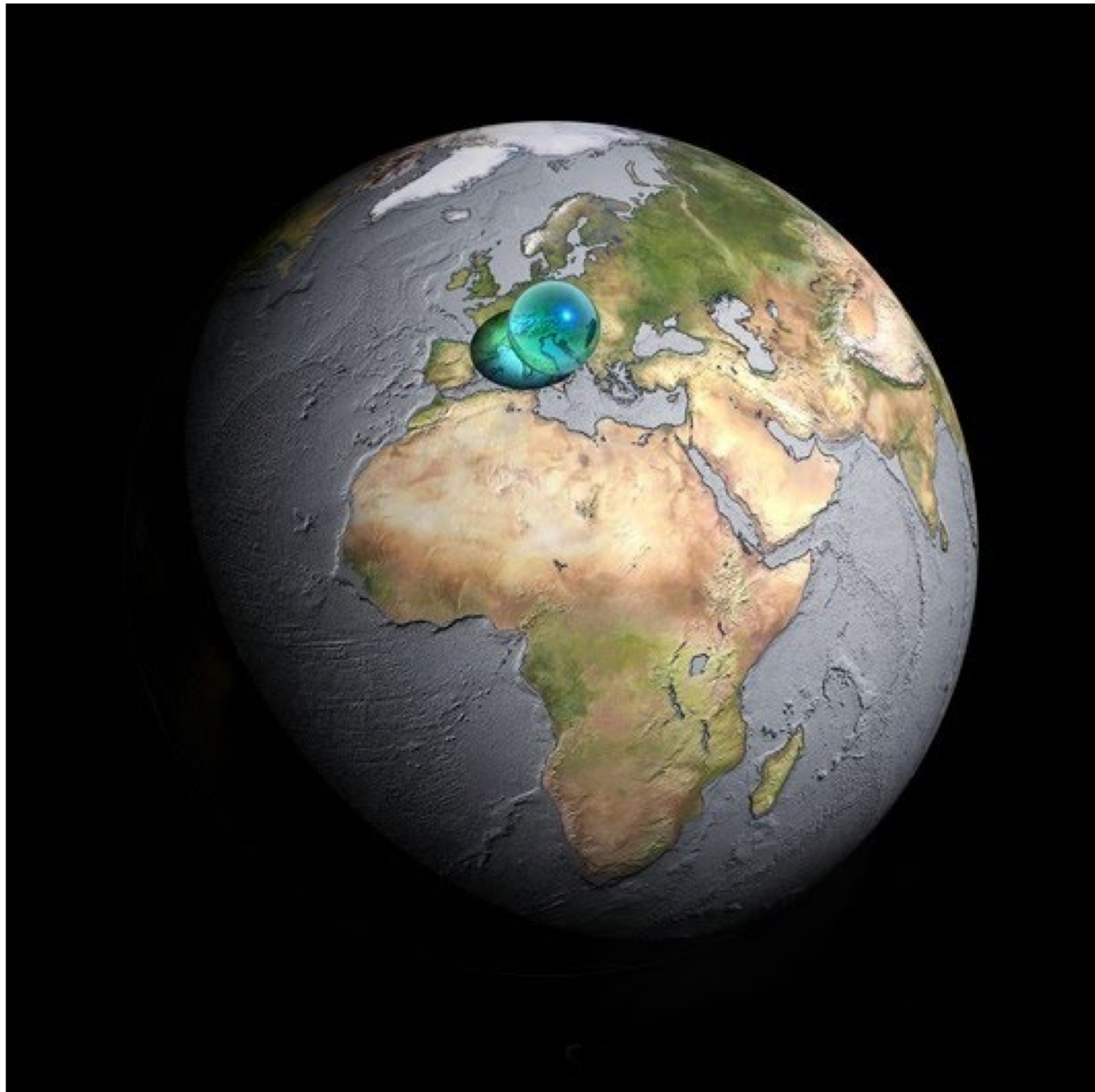
In the Archean Oceans the
primary sources of electrons
were...

H_2 , Fe(II), H_2S and CH_2O

**The major source of electrons
today is**

**LIQUID WATER
(H₂O)**





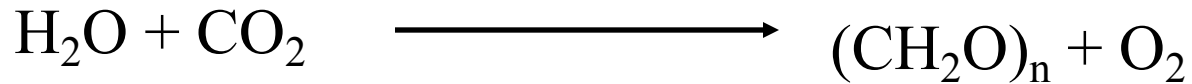
as per Vernadsky

All living organisms on Earth exchange a gas with their environment – via redox reactions

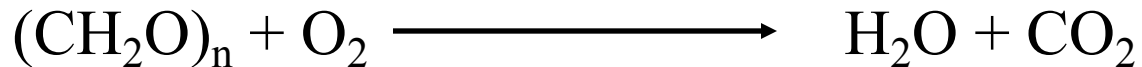
Many of the core metabolic processes are related to gas exchange processes.

The ensemble of redox reactions are coupled on a Planetary SCALE

Oxygenic Photosynthesis

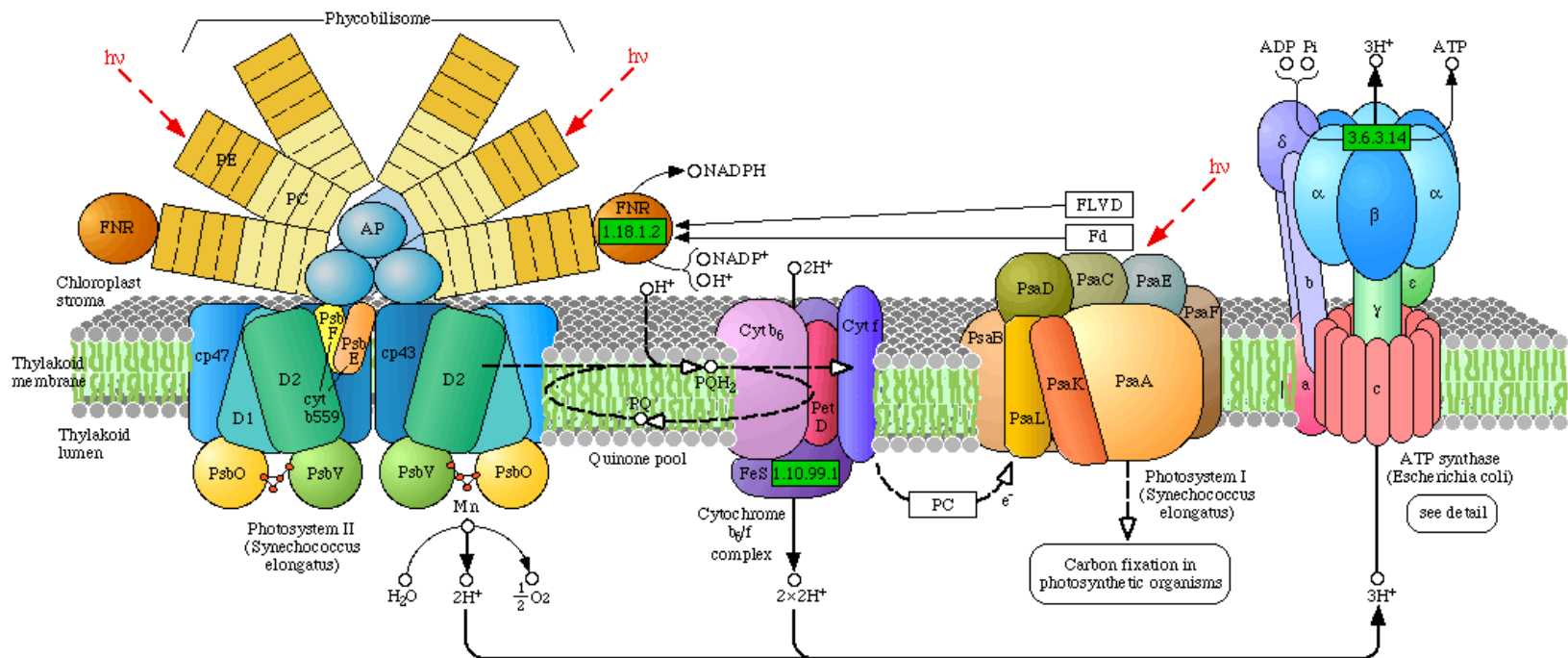


Aerobic Respiration:



The evolution of nannomachines — selection of biological catalysis

PHOTOSYNTHESIS



Photosystem II

| D1 | D2 | cp43 | cp47 | cyt b559 | |
|------|------|------|------|----------|------|
| PsbA | PsbD | PsbC | PsbE | PsbE | PsbF |

| | | | | | | | |
|------|------|------|------|------|------|------|------|
| PsbL | PsbJ | PsbK | PsbM | PsbN | PsbH | PsbT | PsbI |
| PsbO | PsbP | PsbU | PsbV | PsbW | PsbX | PsbY | PsbZ |

Photosystem I

| | | | | | | | |
|------|------|------|------|------|------|------|------|
| PsaA | PsaB | PsaC | PsaD | PsaE | PsaF | PsaH | PsaI |
| PsaJ | PsaK | PsaL | PsaM | PsaN | PsaX | | |

Cytochrome b6/f complex

| | | | | | | | | | | |
|------|------|------|------|------|------|------|------|------|------|------|
| PetB | PetD | PetA | PetC | PetL | PetM | PetN | PetG | PetE | PetF | PetI |
|------|------|------|------|------|------|------|------|------|------|------|

Allophycocyanin (AP)

| | | | | | |
|------|------|------|------|------|------|
| ApcA | ApcB | ApcC | ApcD | ApcE | ApcF |
|------|------|------|------|------|------|

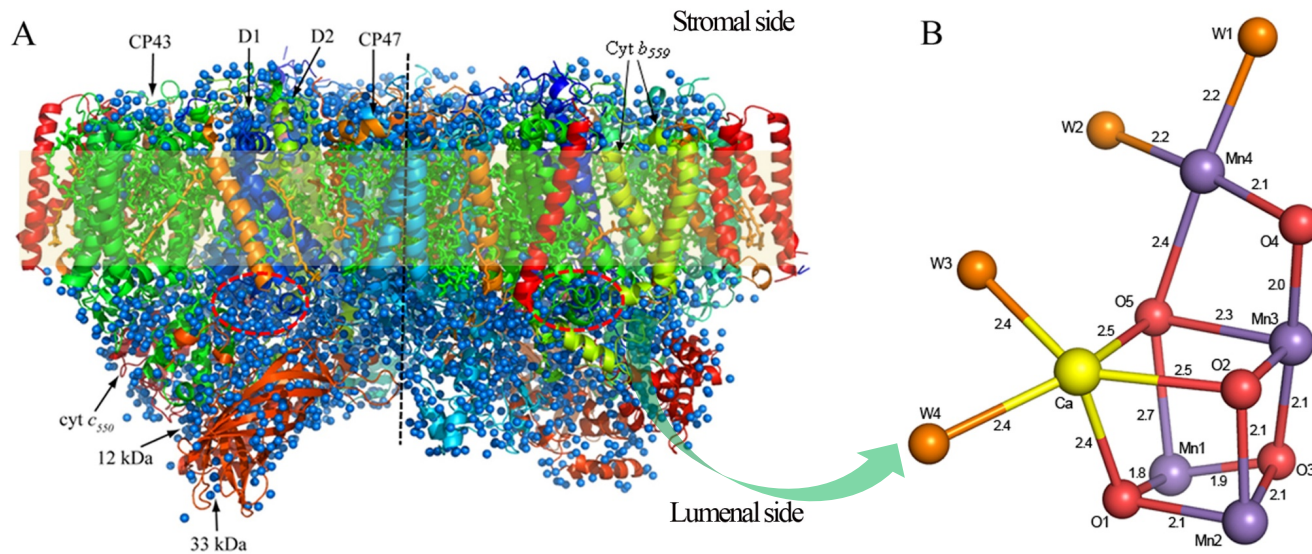
Phycocyanine (PC)

| | | | | | | |
|------|------|------|------|------|------|------|
| CpcA | CpcB | CpcC | CpcD | CpcE | CpcF | CpcG |
|------|------|------|------|------|------|------|

Phycoerythrin (PE)

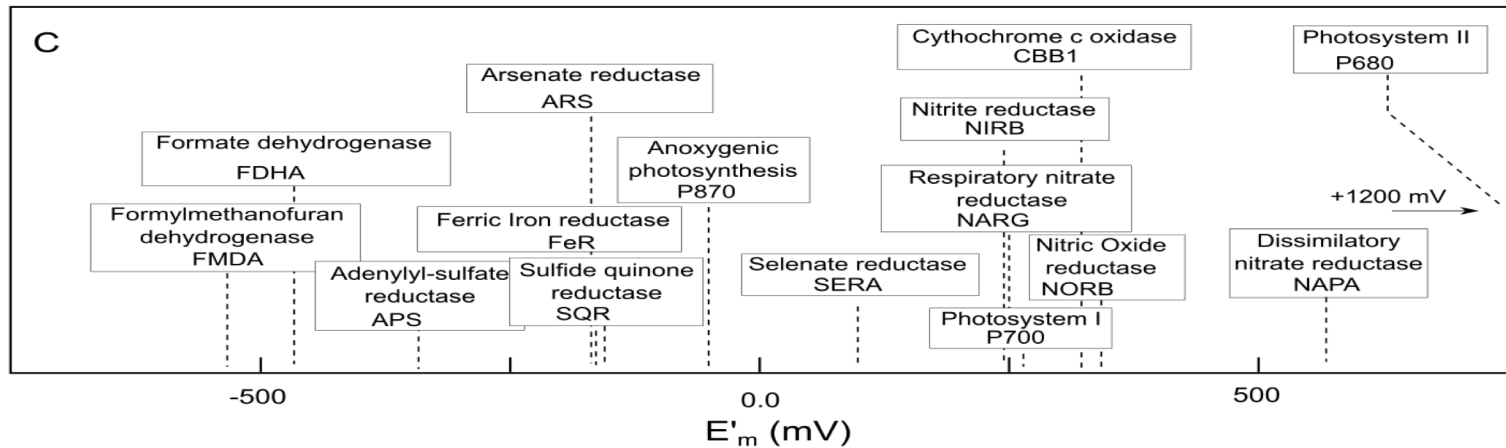
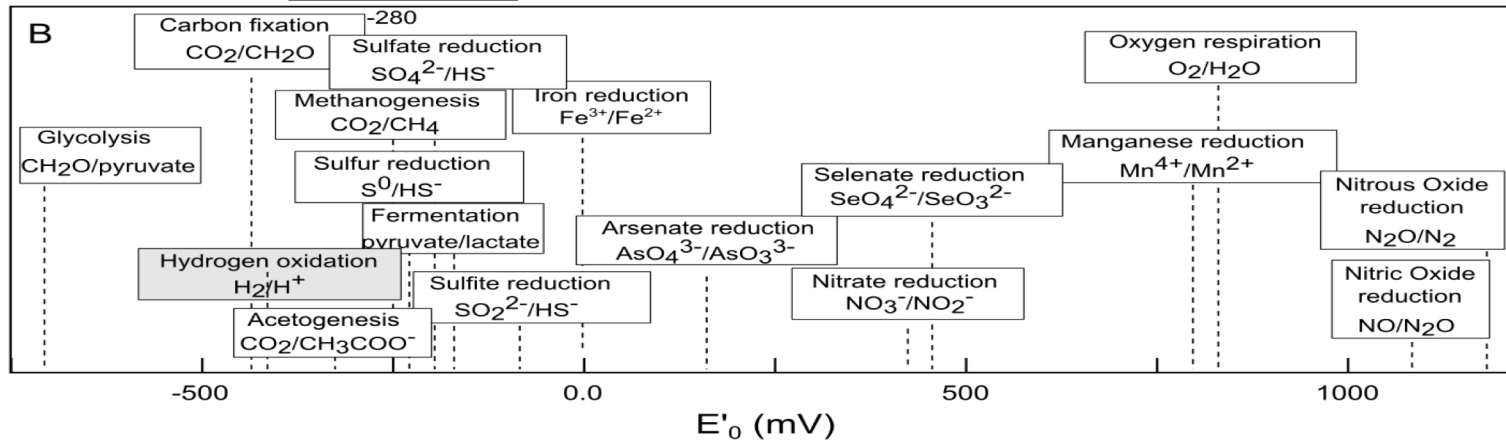
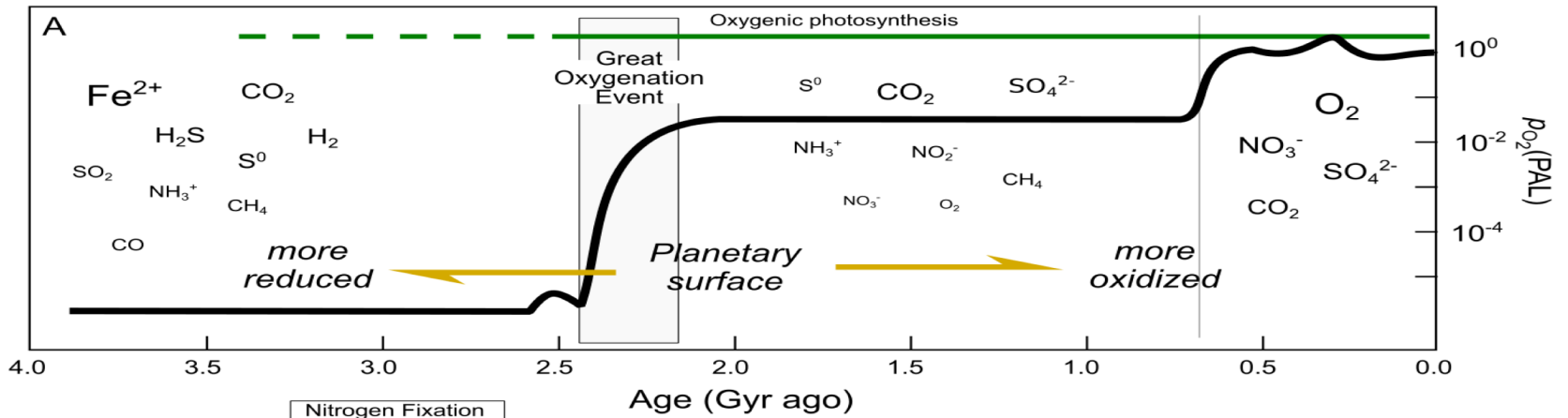
| | | | | |
|------|------|------|------|------|
| PecA | PecB | PecC | PecE | PecF |
|------|------|------|------|------|

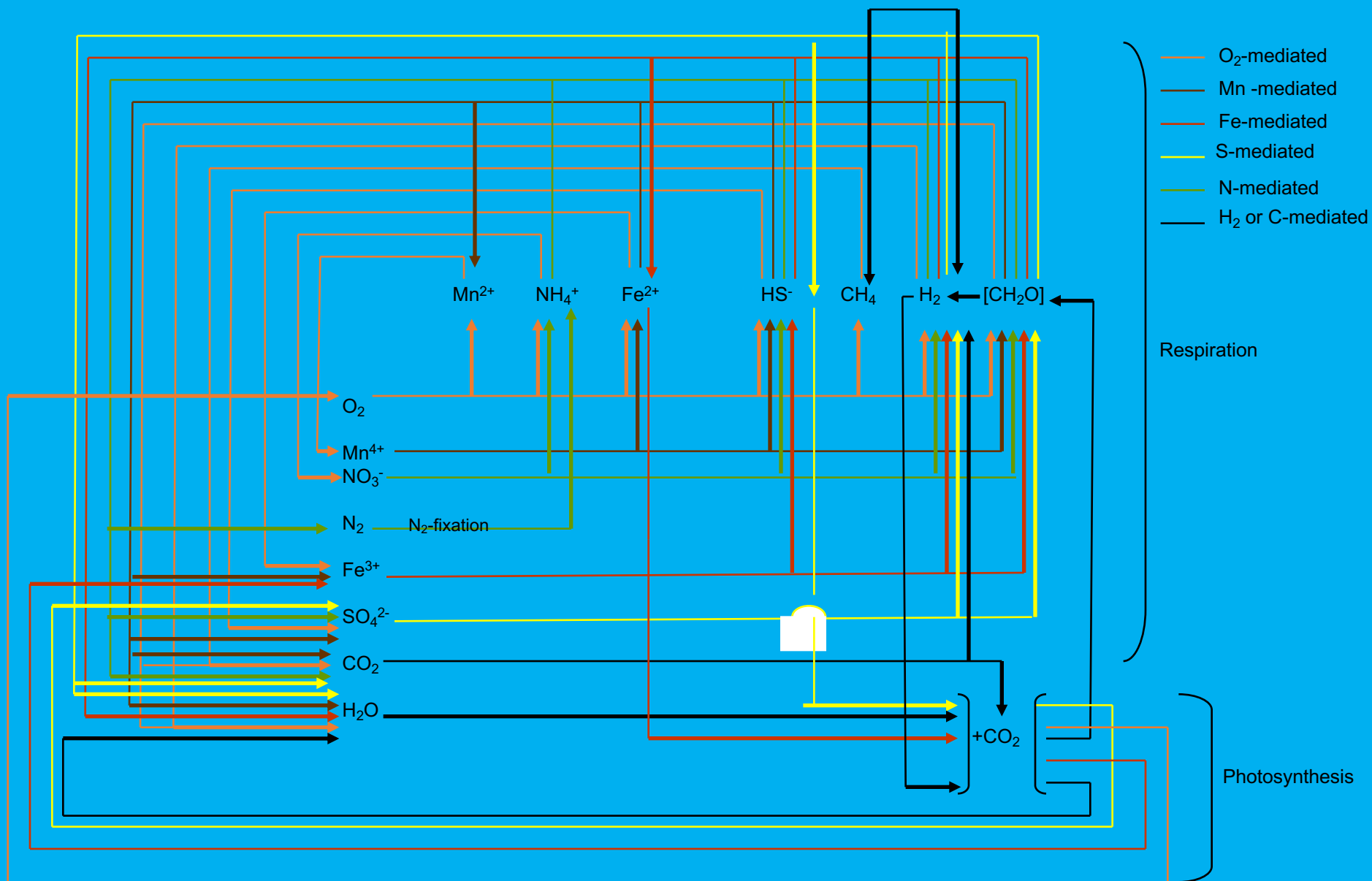
1.9 Å resolution structure of PSII dimer



Nature (2011, 2015, 2017)

Total subunits : 40
Overall MW: 700 kDa
The largest membrane protein complex whose structure has been solved beyond 2.0 Å





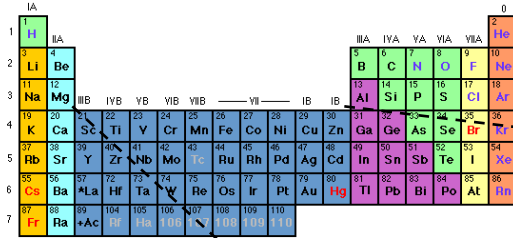
Falkowski, Fenchel and Delong, Science, 2008

This annotation of this metabolic map reveals there are only about **400** core genes responsible for all electron transport reactions on the planet!

How are the electrons connected?

“Periodic Table” of Transition Metal Binding Proteins

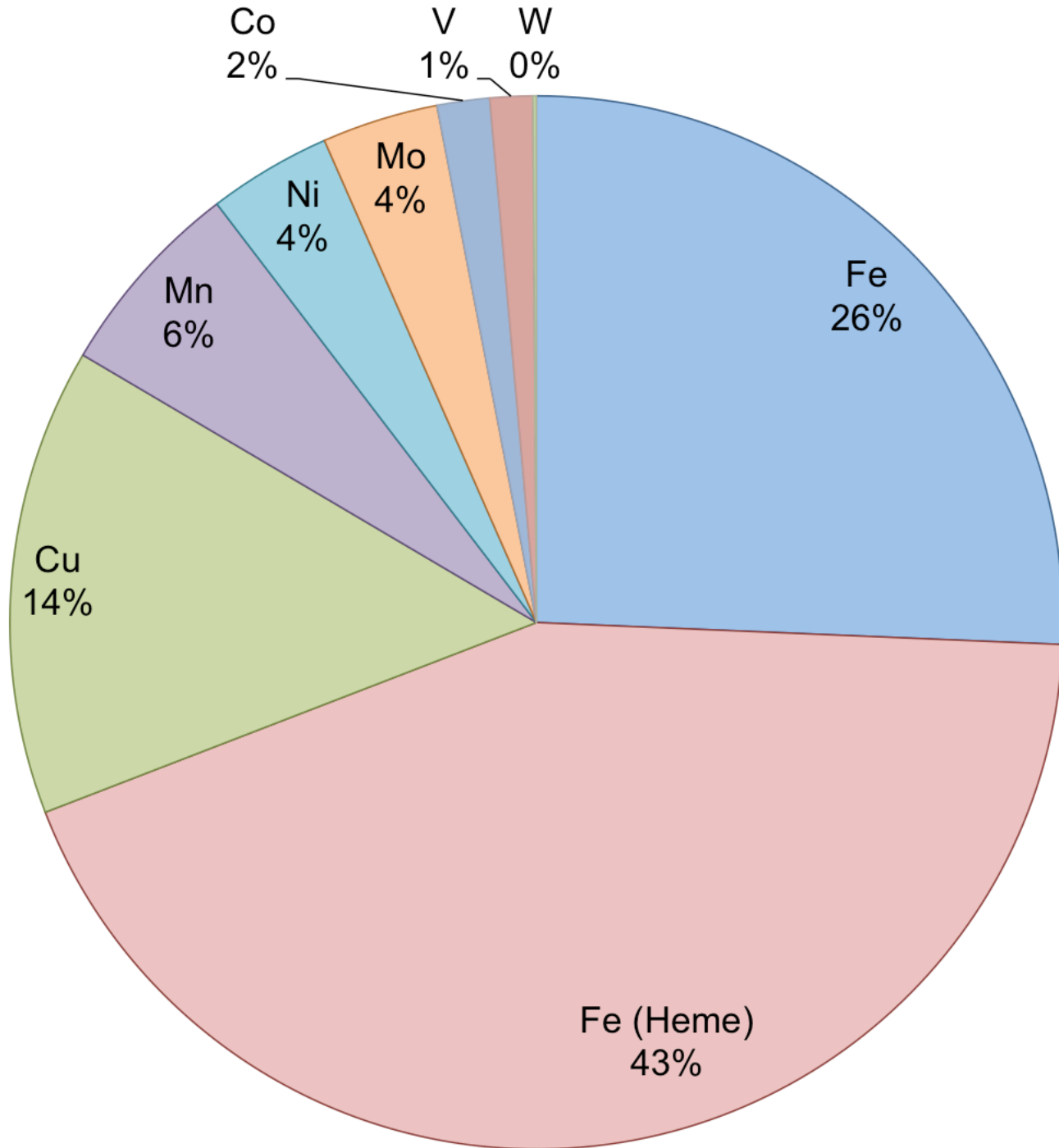
Swiss-Prot+Gene Ontology



| | | | | | | | | | |
|--|--|---|---|---------------------------------------|---|---|--|---|--|
| 21 Sc Scandium 44.955910 | 22 Ti Titanium 47.867 | 7 <input type="text"/> 1 V 7 | 24 Cr Chromium 51.9961 | 647 23 Mn 7039 | 8248 <input type="text"/> 360 Fe 25006 | 86 <input type="text"/> Co 914 | 247 75 Ni 2829 | 1299 <input type="text"/> 47 Cu 2730 | 2090 <input type="text"/> 77 Zn 37263 |
| 39 Y Yttrium 88.90585 | 40 Zr Zirconium 91.224 | 41 Nb Niobium 92.90638 | 42 452 16 Mo 554 | 43 Tc Technetium (98) | 44 Ru Ruthenium 101.07 | 45 Rh Rhodium 102.90550 | 46 Pd Palladium 106.42 | 47 Ag Silver 107.8682 | 4 Cd 39 |
| 57 La Lanthanum 138.9055 | 72 Hf Hafnium 178.49 | 73 Ta Tantalum 180.9479 | 1 <input type="text"/> 1 W 1 | 75 Re Rhenium 186.207 | 76 Os Osmium 190.23 | 77 Ir Iridium 192.217 | 78 Pt Platinum 195.078 | 79 Au Gold 196.96655 | 26 <input type="text"/> 1 Hg 78 |
| 89 Ac Actinium (227) | 104 Rf Rutherfordium (261) | 105 Db Dubnium (262) | 106 Sg Seaborgium (263) | 107 Bh Bohrium (262) | 108 Hs Hassium (265) | 109 Mt Meitnerium (266) | 110 (269) | 111 (272) | 112 (277) |

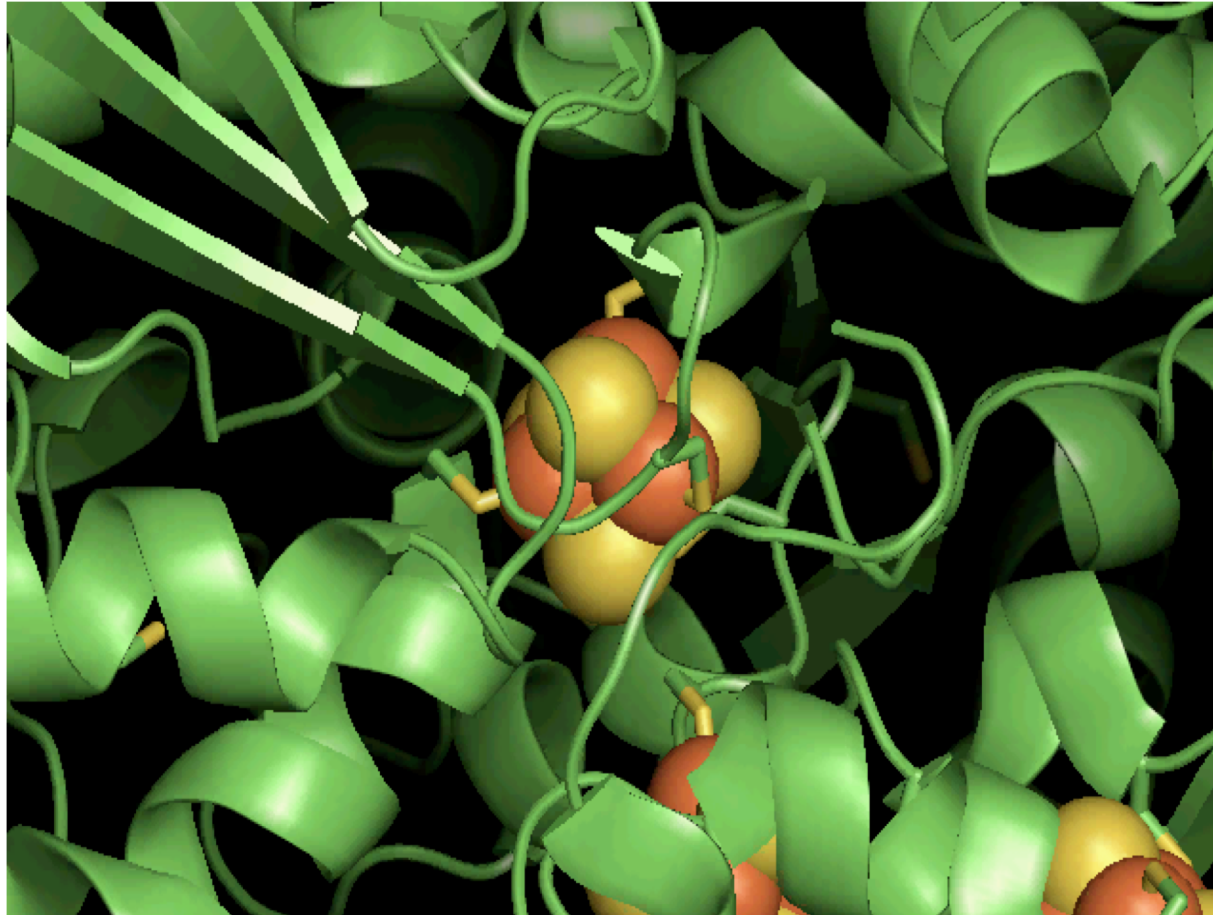
Number of SwissProt proteins associated with transition metals: **All**, **EC1**, and **experimentally validated EC1**





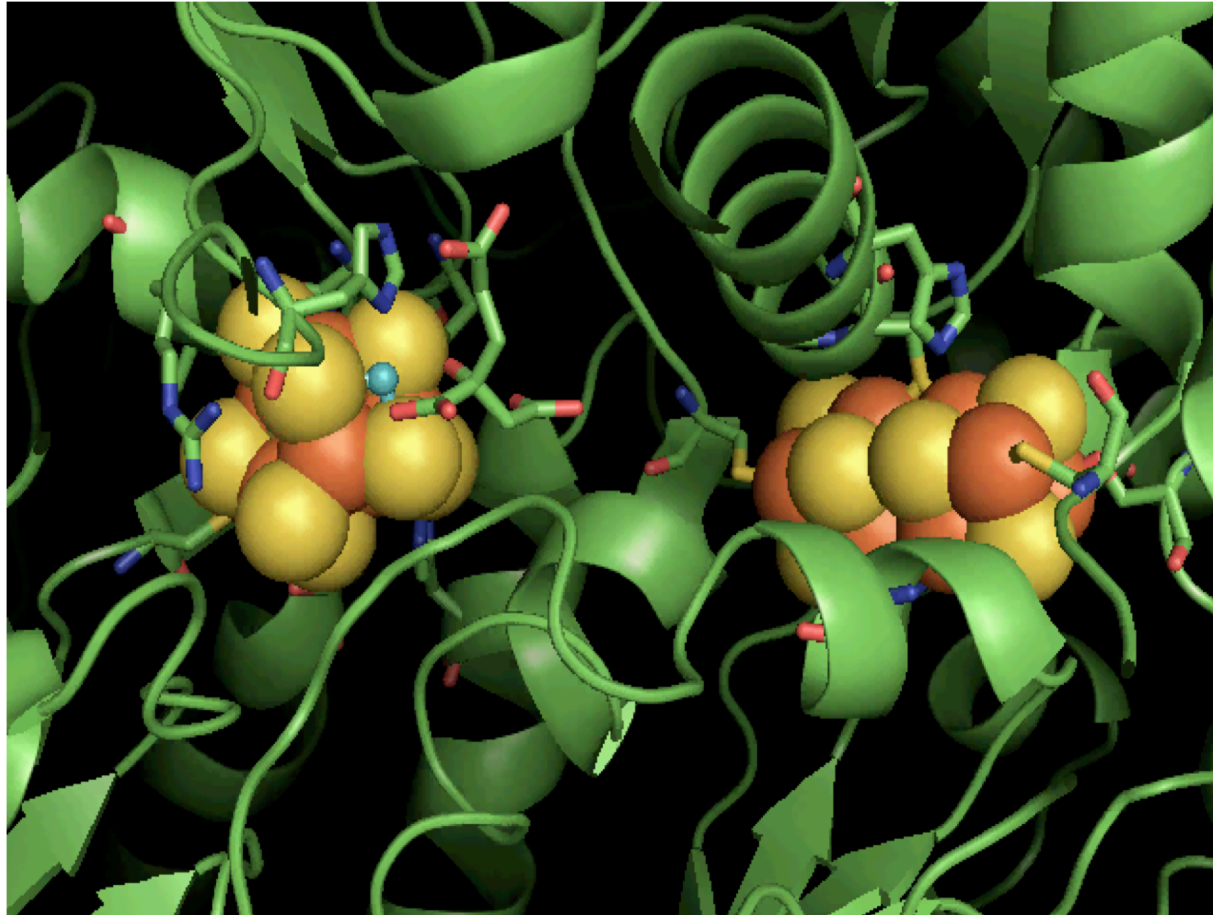
ferredoxin

| | | | | | | | | | |
|---------------------------------------|--|---------------------------------------|--|--|--|--------------------------------------|--|-------------------------------------|--------------------------------------|
| scandium 21 Sc 44.956 | titanium 22 Ti 47.867 | vanadium 23 V 50.942 | chromium 24 Cr 51.996 | manganese 25 Mn 54.938 | iron 26 Fe 55.845 | cobalt 27 Co 58.933 | nickel 28 Ni 58.693 | copper 29 Cu 63.546 | zinc 30 Zn 65.39 |
| yttrium 39 Y 88.906 | zirconium 40 Zr 91.224 | niobium 41 Nb 92.906 | molybdenum 42 Mo 95.94 | technetium 43 Tc [98] | ruthenium 44 Ru 101.07 | rhodium 45 Rh 102.91 | palladium 46 Pd 106.42 | silver 47 Ag 107.87 | cadmium 48 Cd 112.41 |
| lutetium 71 Lu 174.97 | hafnium 72 Hf 178.49 | tantalum 73 Ta 180.95 | tungsten 74 W 183.84 | rhenium 75 Re 186.21 | osmium 76 Os 190.23 | iridium 77 Ir 192.22 | platinum 78 Pt 195.08 | gold 79 Au 196.97 | mercury 80 Hg 200.59 |



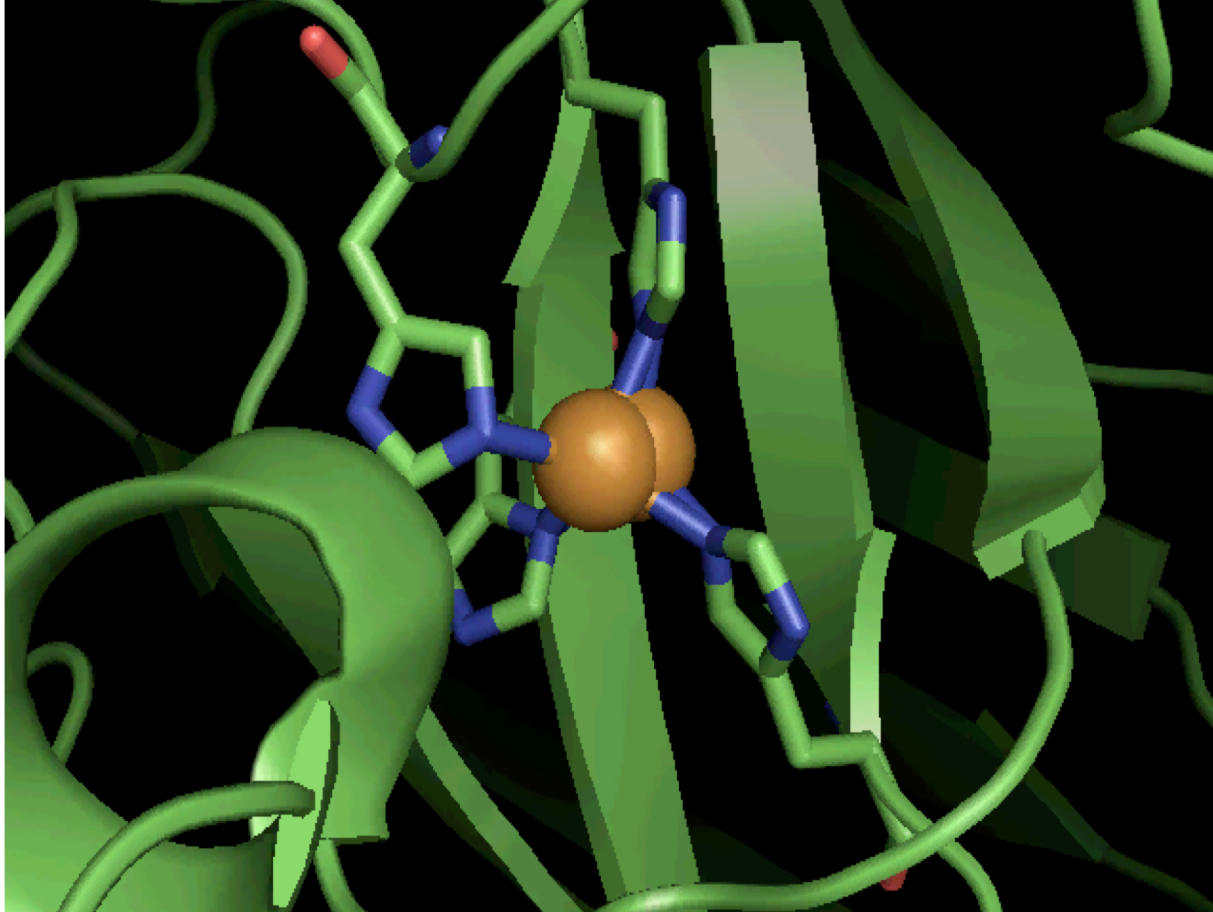
nitrogenase

| | | | | | | | | | |
|---------------------------------------|--|---------------------------------------|--|--|--|--------------------------------------|--|-------------------------------------|--------------------------------------|
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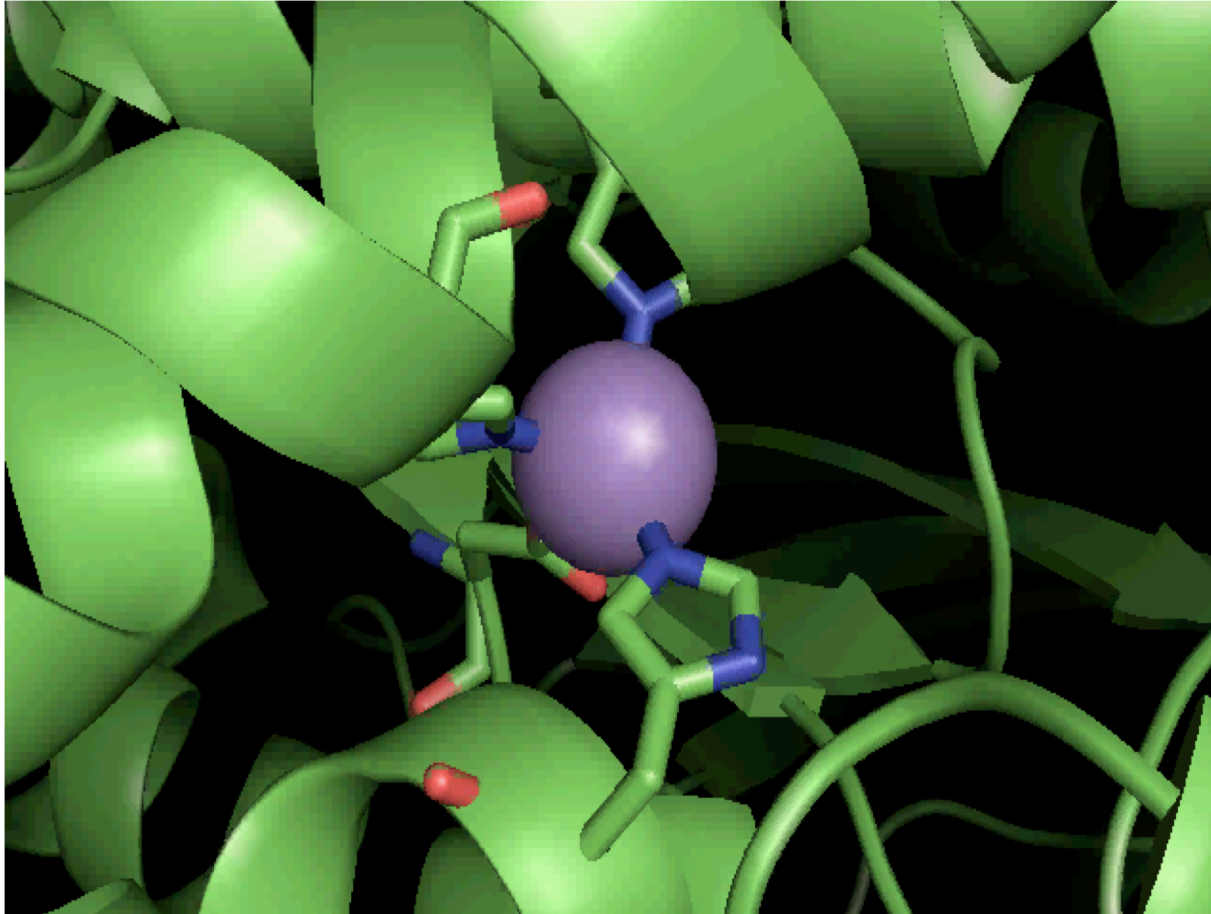
superoxide dismutase

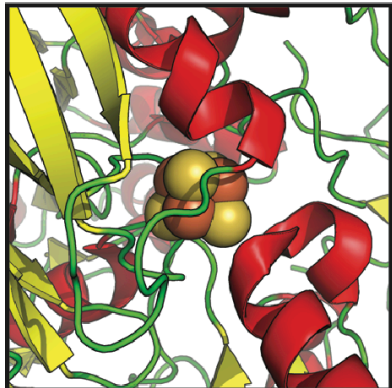
| | | | | | | | | | |
|---------------------------------------|--|---------------------------------------|--|--|--|--------------------------------------|--|--|--------------------------------------|
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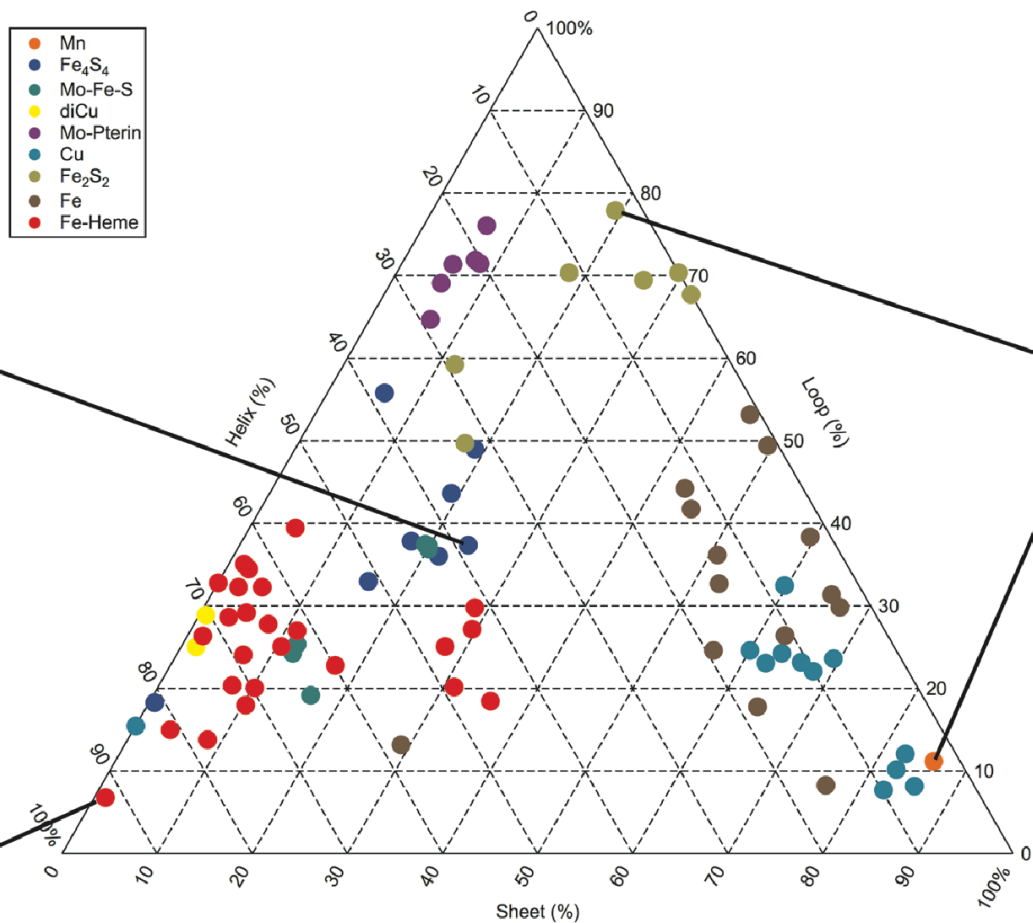
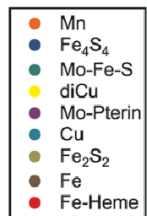
superoxide dismutase

| | | | | | | | | | |
|---------------------------------------|--|---------------------------------------|--|--|--|--------------------------------------|--|-------------------------------------|--------------------------------------|
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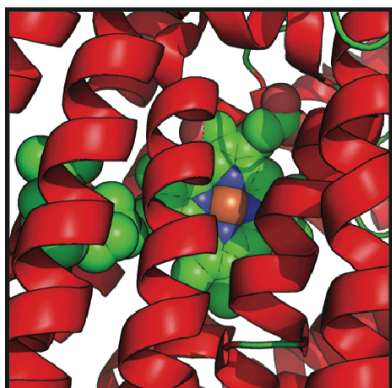




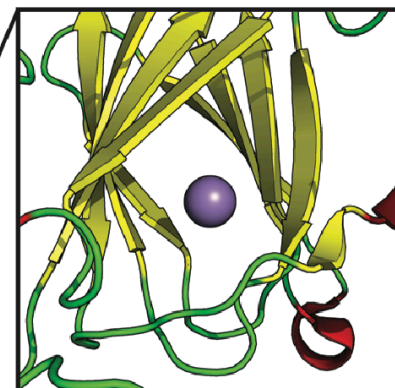
formate dehydrogenase (PDB: 2iv2)



toluene 2,3-dioxygenase (PDB: 3en1)



cytochrome c oxidase (PDB: 2gsm)

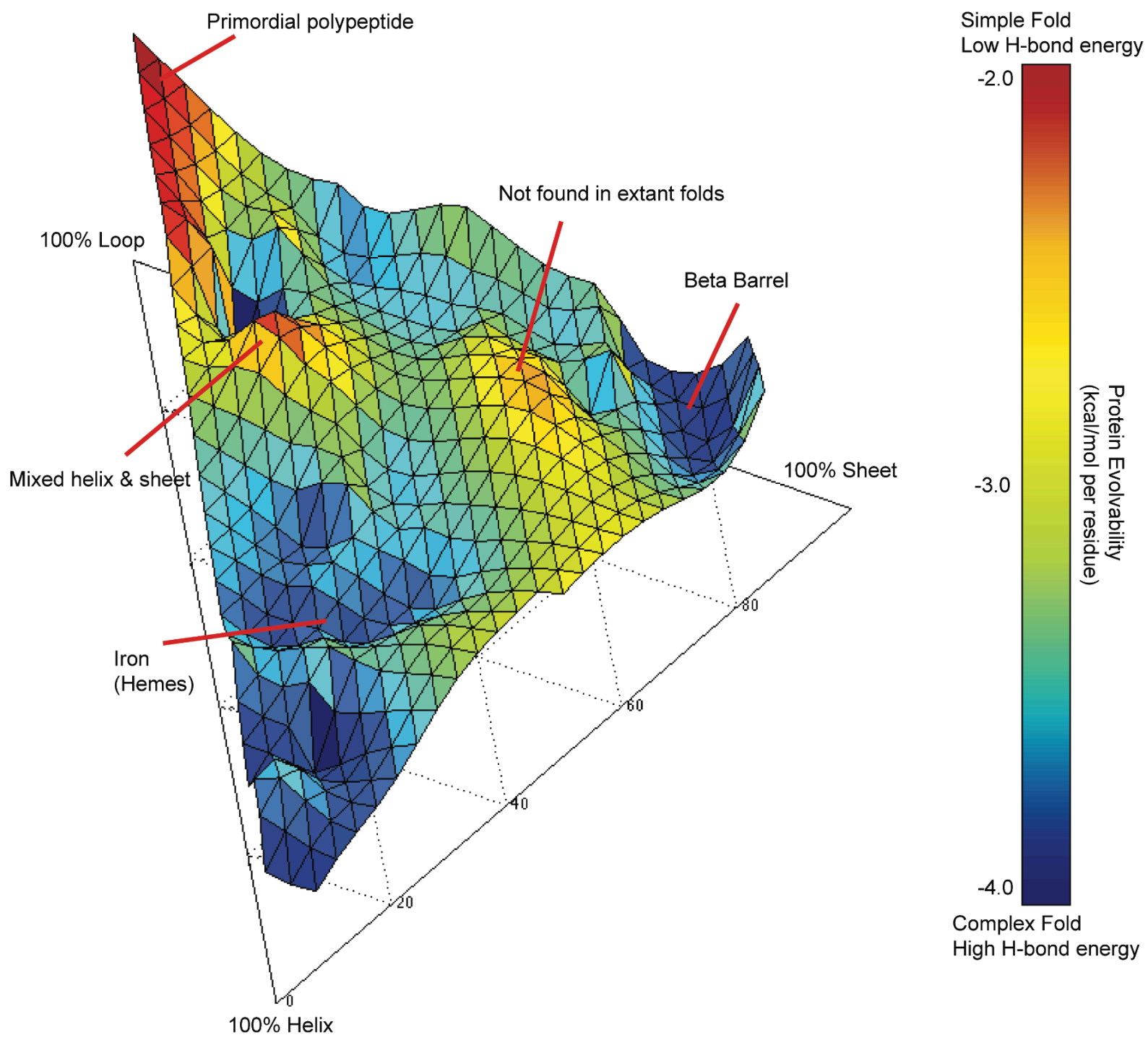


oxalate oxidase (PDB: 2et1)

Sheet (%)

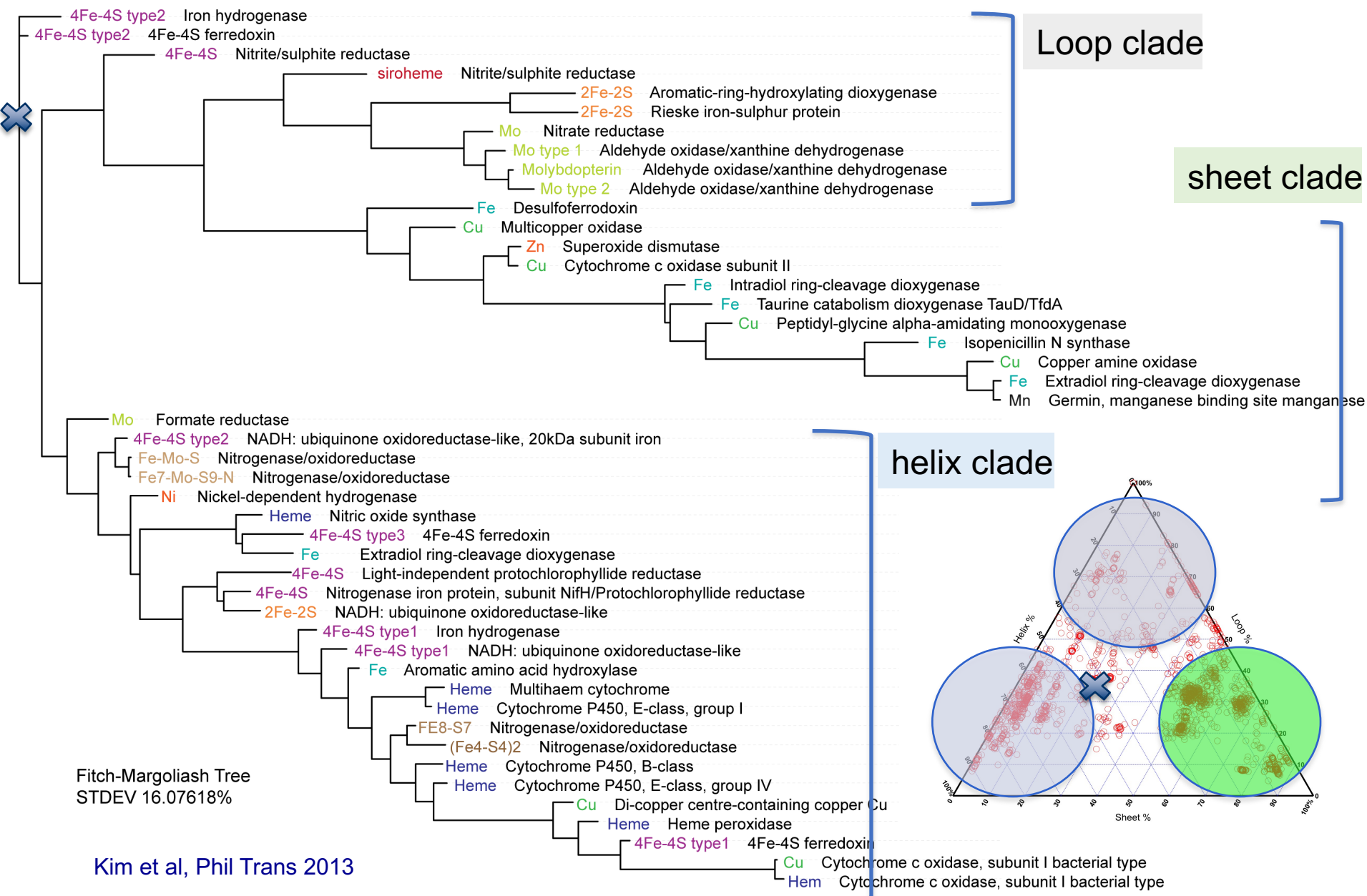
Helix (%)

Loop (%)



A secondary structural tree of EC1 proteins

— 0.01



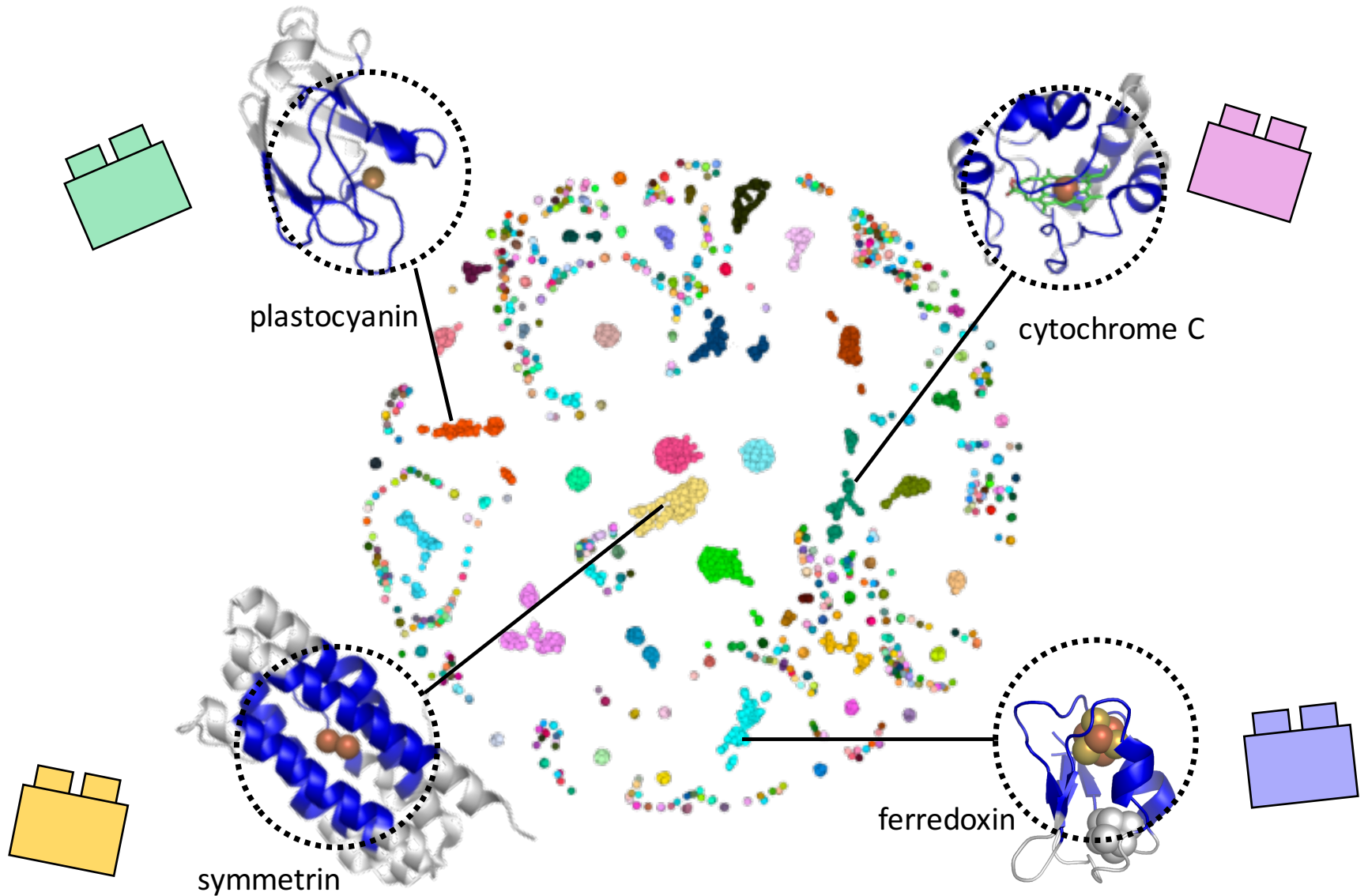
Evolution of Sequences and Folds

- Evolution of core structural motifs
- The paradox of structure/sequence divergence
- ***The processes of natural selection severely inhibit any change a well-adapted system on which several other essential components depend. (Eck RV, Dayhoff MO, 1966)***

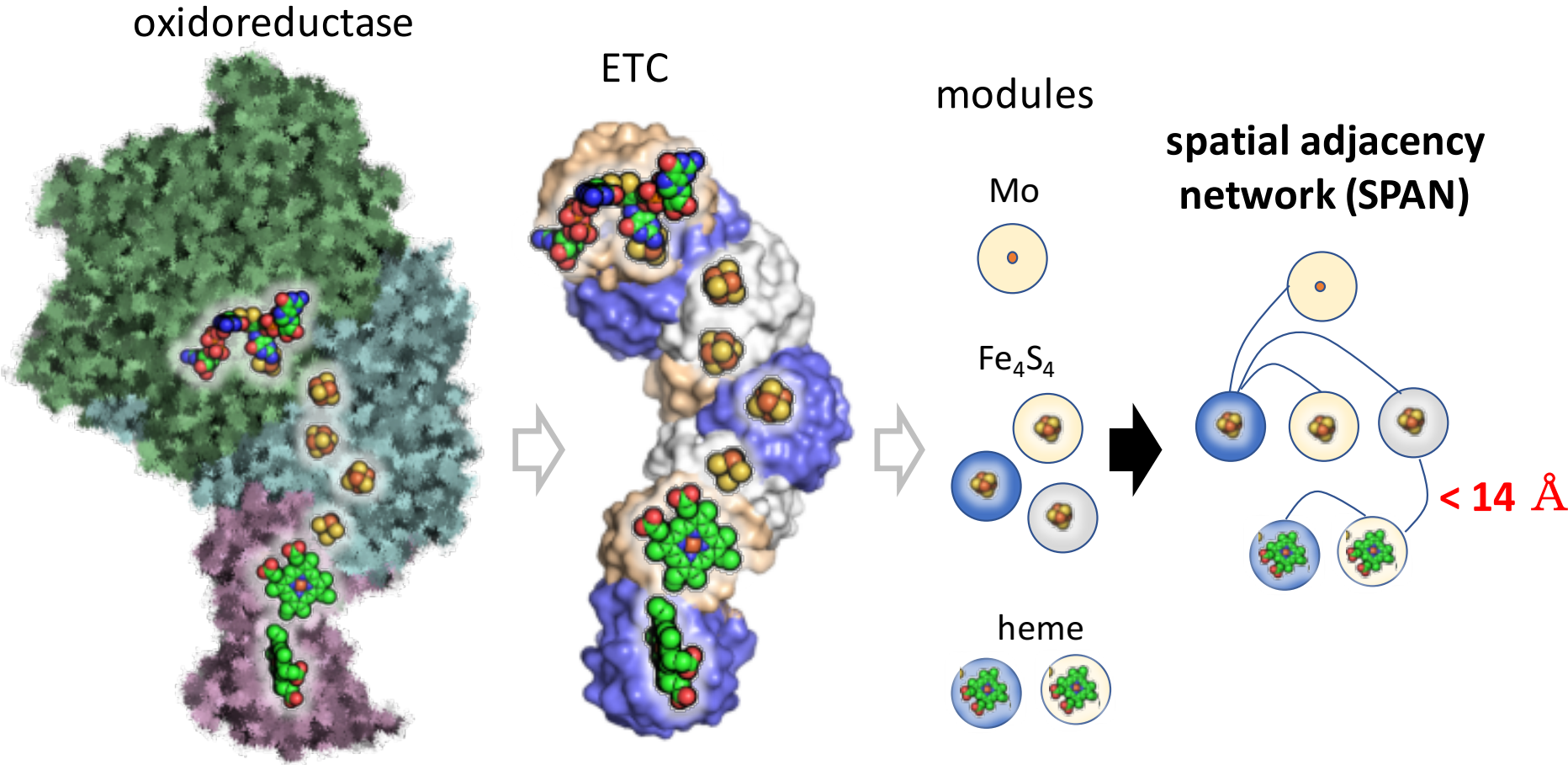
Motifs (i.e., folds)

- There appear to be only **35** motifs in all of the extant electron transfer reactions in nature.
- Deciphering the evolutionary history of these motifs is one of the grand challenges of science.

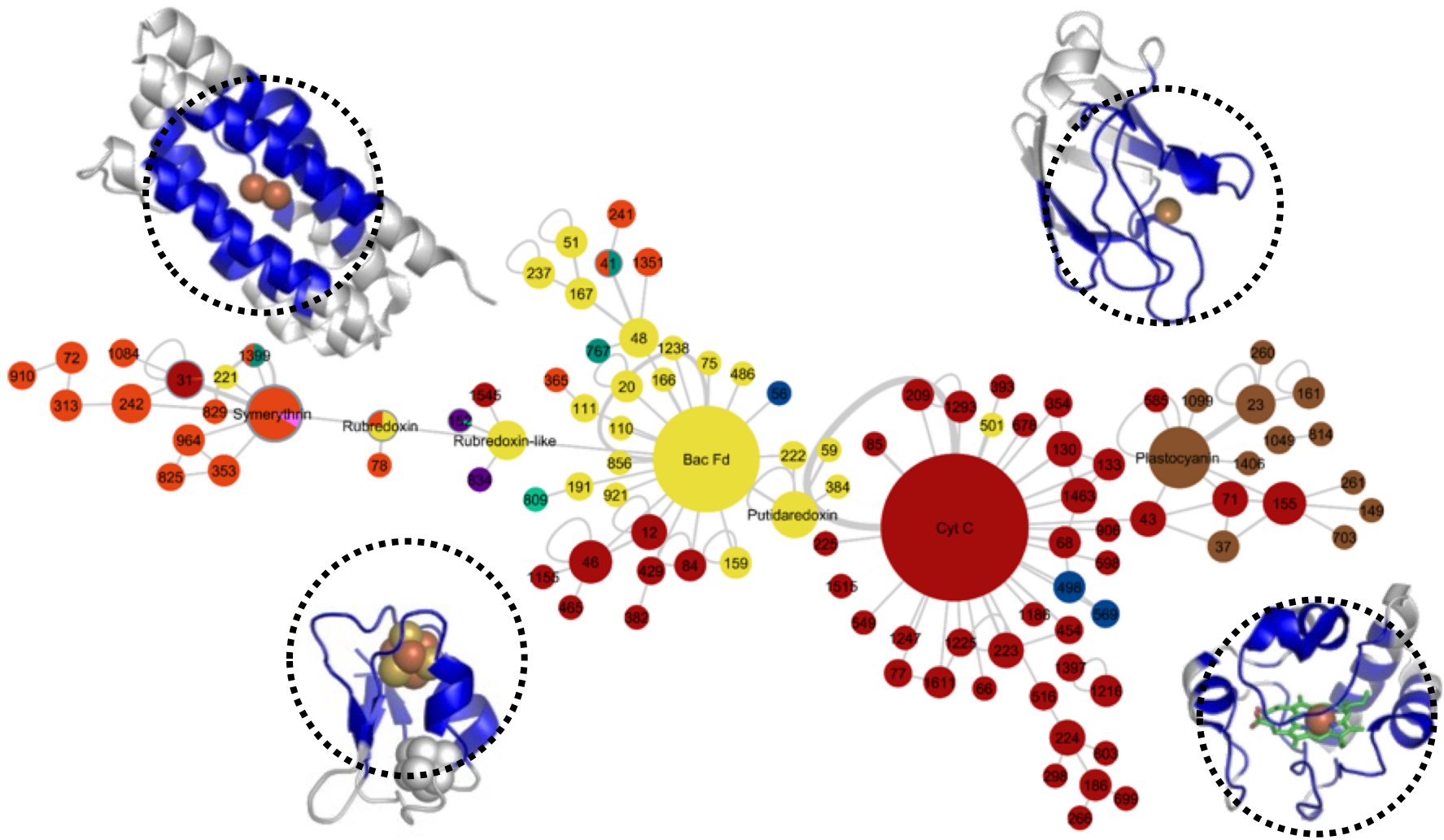
“redox protein construction kit” ~ 1000 domains



construct a SPAN for oxidoreductases



oxidoreductase SPAN composed of ~100 modules

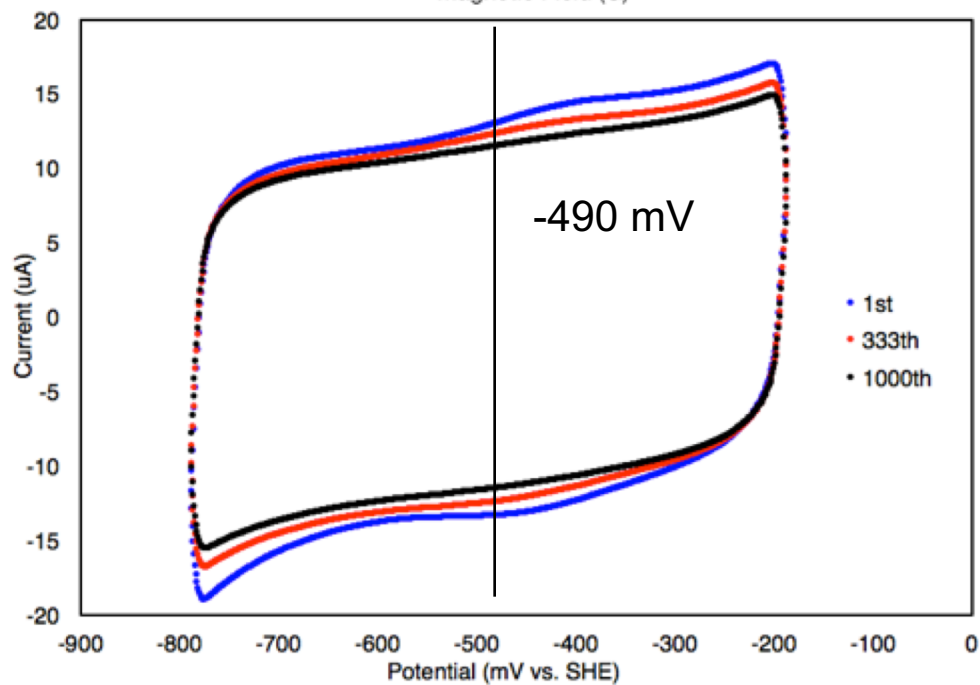
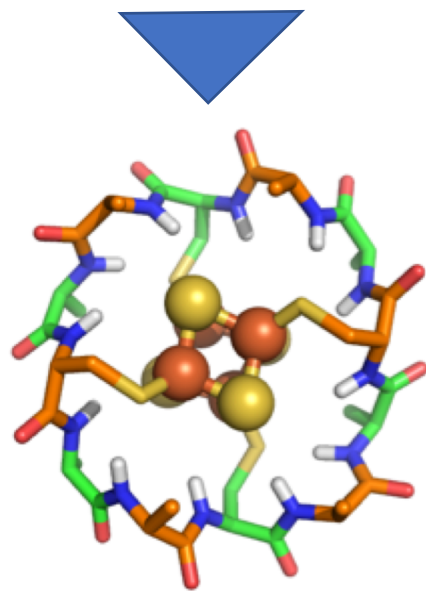
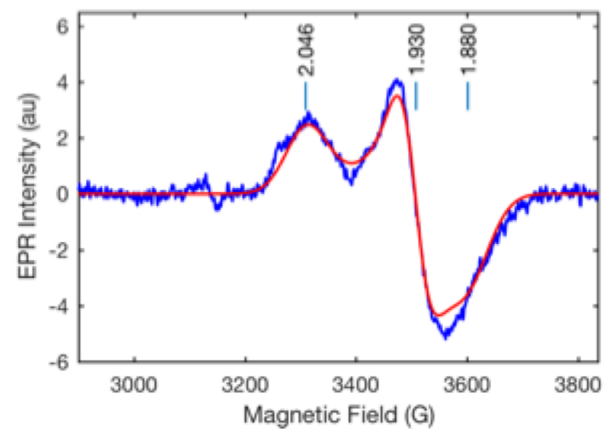
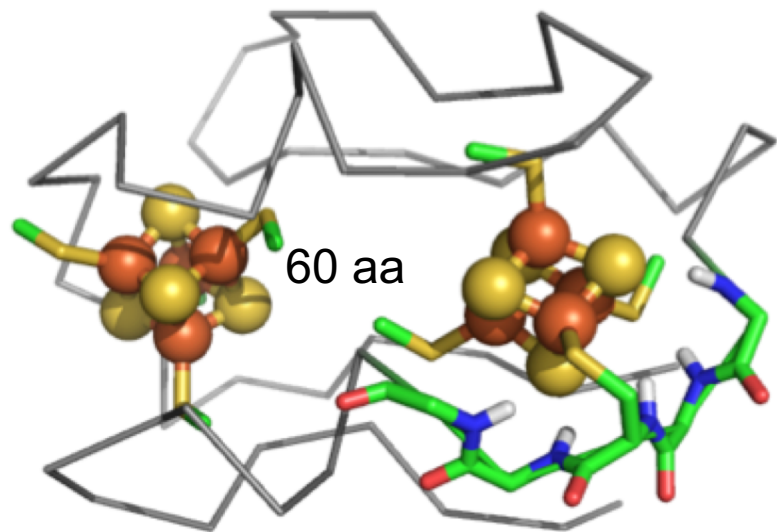


node = module; size = number of edges to other modules

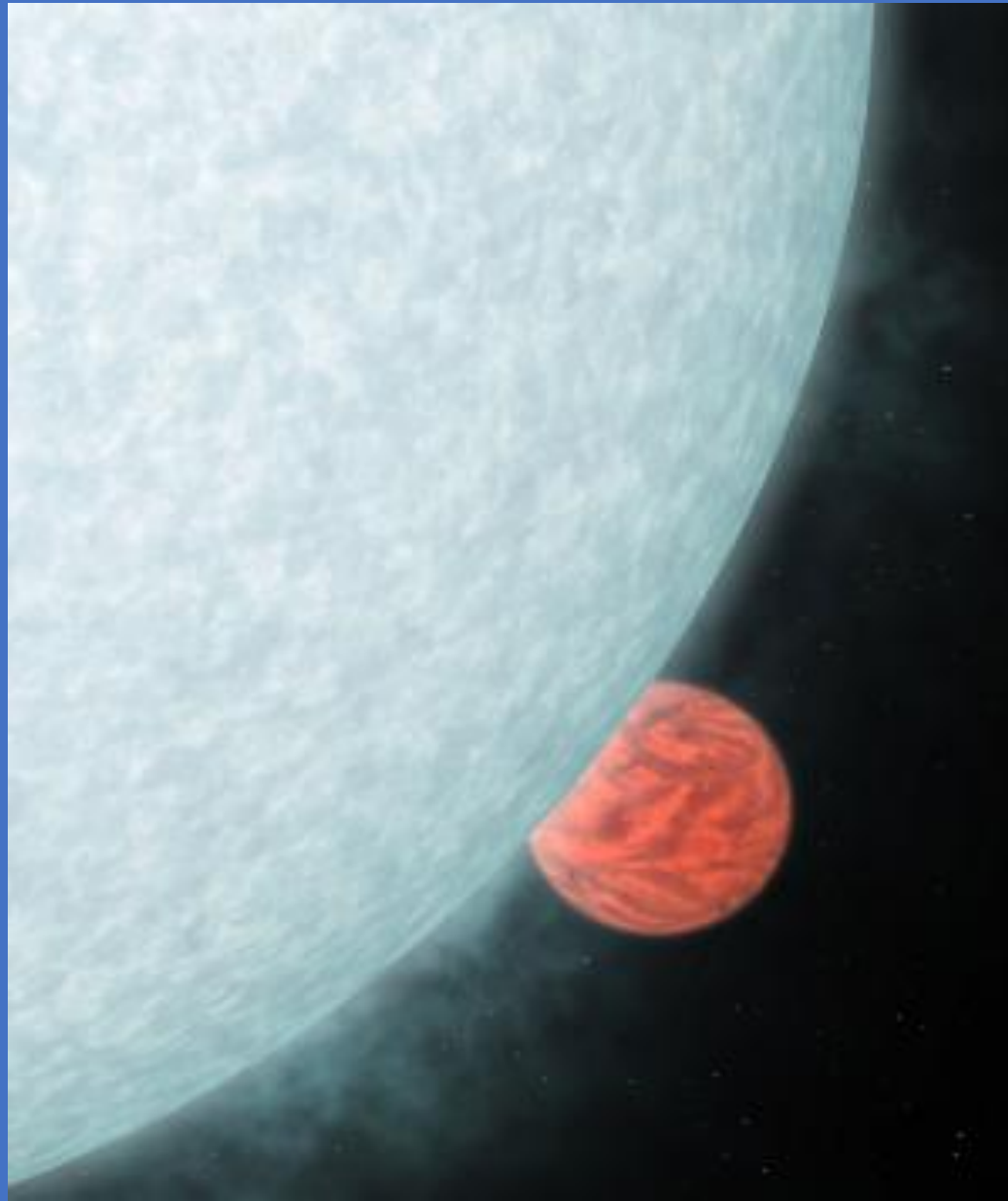
edge = co-occurrence in structure (m-m distance < 14Å);
thickness = number of instances

Ranaan et. al. *PNAS* (2018)

design of a primordial ferredoxin



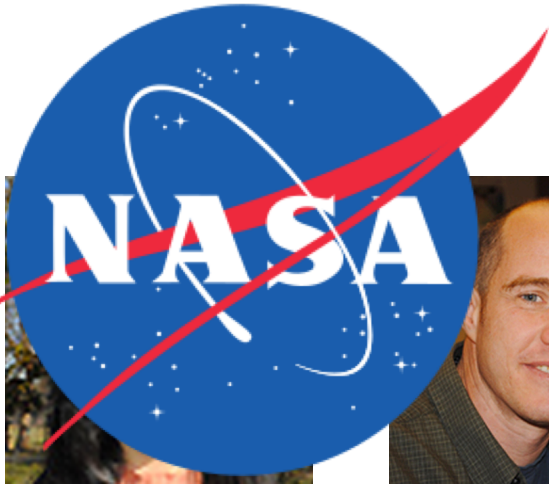
Glow from other planets



Conclusions

- 1. In the first ca. 2.5 Ga of Earth's history, nature invested heavily in R&D from which a “core” set of metabolic machines evolved.**
- 2. All of the key metabolic processes were developed in prokaryotes**
- 3. There are approximately 400 core metabolic genes that make biological electrons flow across the planet world.**
- 4. These metabolic sequences are coupled on local and planetary scales to facilitate an electron market between C, N, O, and S.**
- 5. The electronic potential is driven by light**

THE TEAM



Shu Cheng



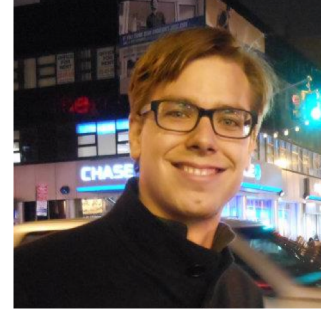
Arye Harel



Ben Jelen



John Kim



Stefan Senn



Debashish
Bhattacharya



Yana Bromberg



David Case



Paul Falkowski



Max Haggblom



Vikas Nanda



Nathan Yee



A photograph of a sunset over a large body of water. The sky is filled with dark, dramatic clouds, with a bright orange and yellow glow from the setting sun breaking through in the center. The water reflects the colors of the sky. In the foreground, there is a dark, silhouetted shoreline with some trees and bushes. The text "Thank you!" is overlaid in a bright yellow, bold font in the center of the image.

Thank you!

