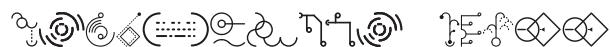


Electron Bifurcation

life's 3rd mechanism of energy conversion

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A brief history of earth and life

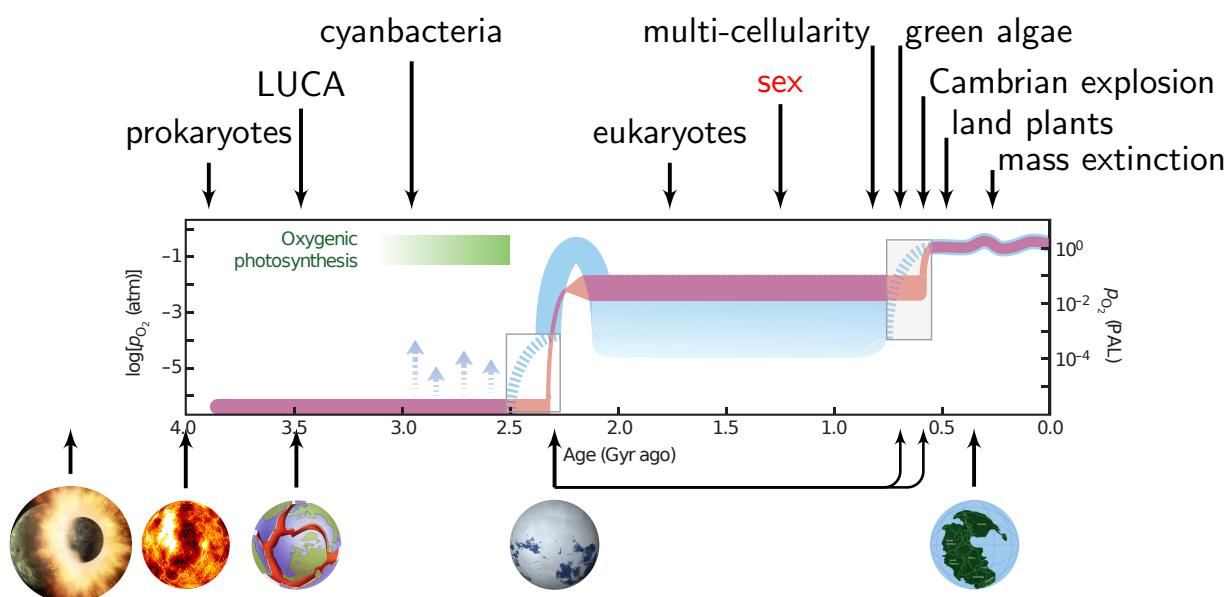
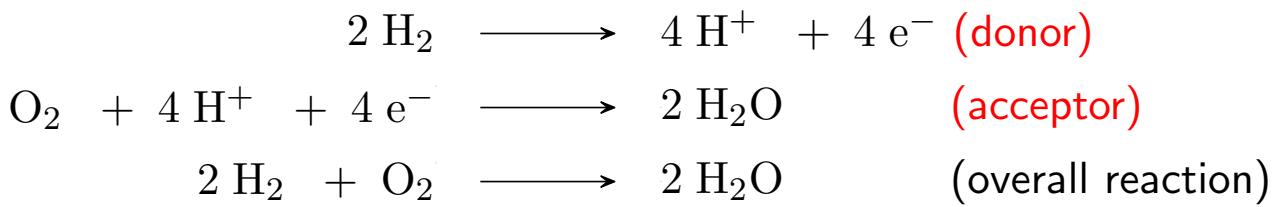


Figure modified from Lyons TW, Reinhard CT & Planavsky NJ (2014), The rise of oxygen in Earth's early ocean and atmosphere, *Nature* 506:307-315 | doi:10.1038/nature13068

Redox reaction



Obviously, not all couples of redox substrates yield the same amount of energy.

ΔG depends on the difference in standard electrochemical potential ΔE_m between e^- donor and acceptor and concentration (Nernst-Equation).

$$E_{h,7} = E_{m,7} + \frac{1}{n} \cdot 59 \text{ mV} \cdot \log([ox]^a/[red]^b)$$

$E_{m,7}$... redox midpoint potential at pH 7

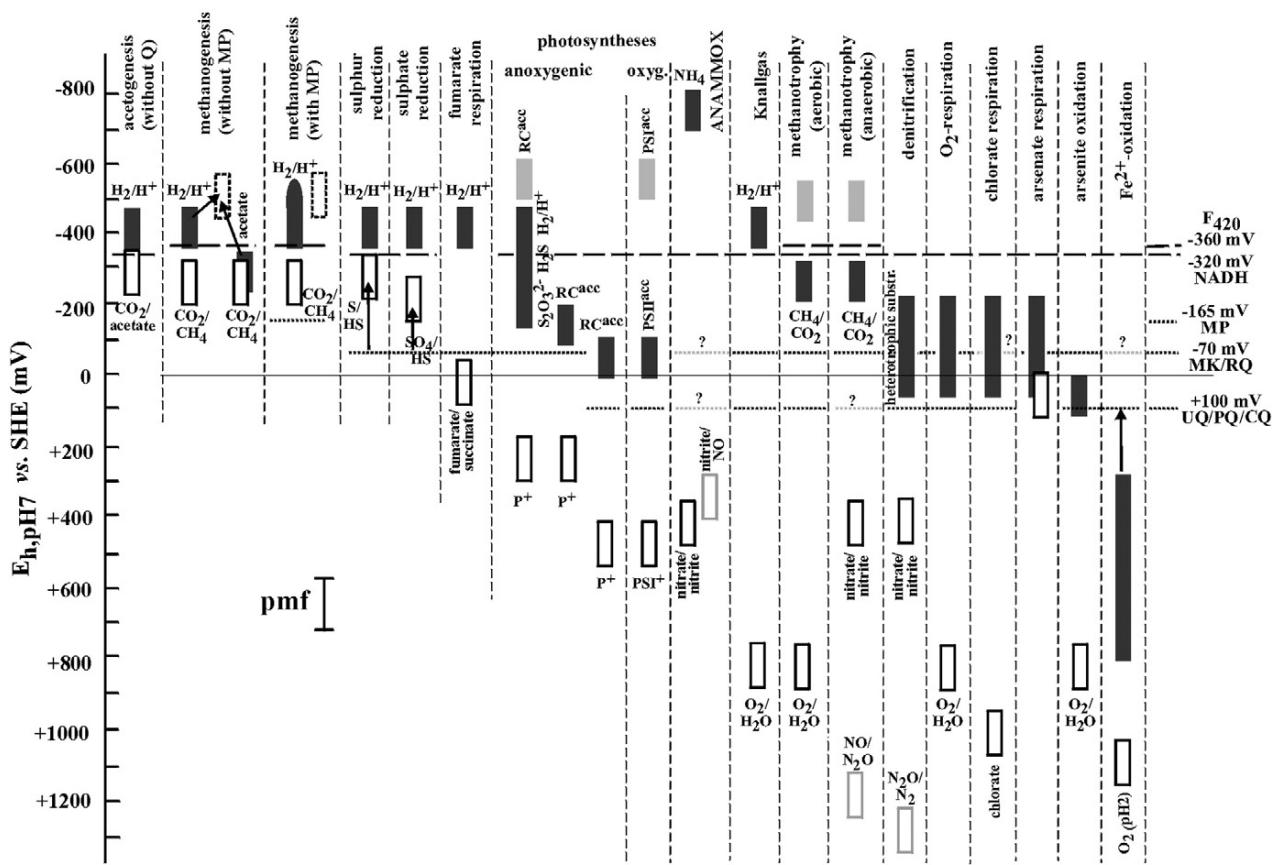
n ... number of transferred electrons.

a, b ... stoichiometric coefficients of oxidized donor / reduced acceptor

Electrochemical driving force for redox equilibrations between a donor (D) and an acceptor (A) molecule: $\Delta E_h = E_h(A) - E_h(D)$; Gibbs free energy of redox reaction: $\Delta G = nF\Delta E_h$ with n , number of transferred electrons and F , Faraday constant

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Life on the redox-scale



black/white boxes are electron donors/acceptors; proton motive force (pmf)

energetically favorable is from more negative to more positive electrochemical potential.

The core of bioenergetics

Life's ability to create order from randomness rests on **disequilibria**:

- ① chemical (ATP/ADP+P_i).
- ② electrochemical (NAD(P)H/oxidizing environment).

$$-\frac{\Delta G}{kT} = \ln \left(\frac{[\text{ATP}]/[\text{ADP}][\text{P}_i]}{[\text{ATP}]_{\text{eq}}/[\text{ADP}]_{\text{eq}}[\text{P}_i]_{\text{eq}}} \right) \approx 20 - 24$$

$\Delta G_{\text{ATP}} \sim 50 - 60 \text{ kJ/mol}$; $kT \sim 3 \text{ kJ/mol}$; unit activities are assumed

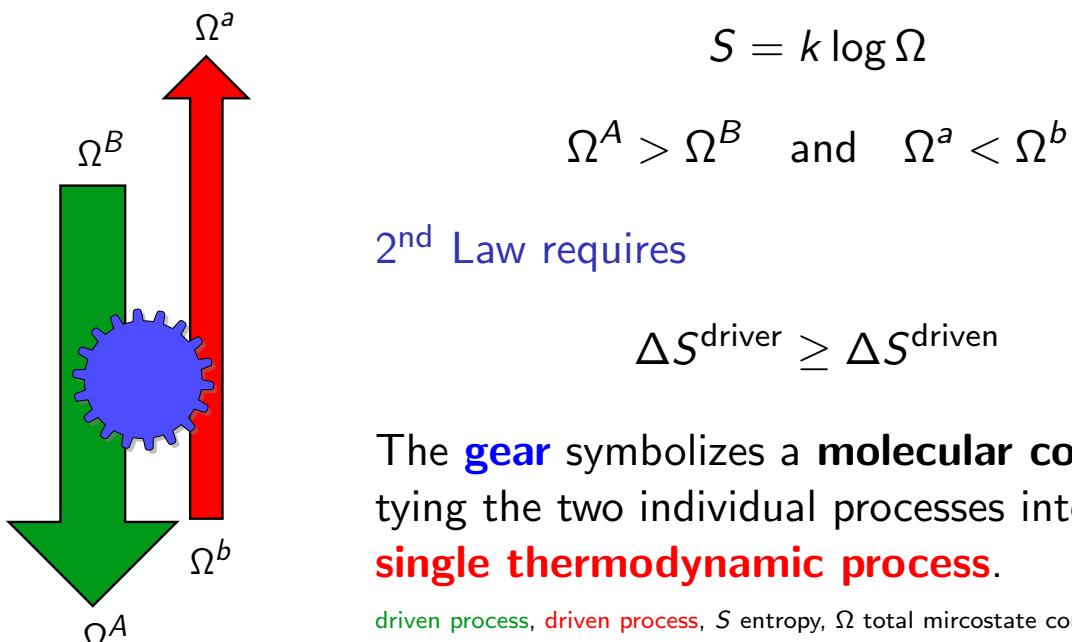
Implying that the **ratio** of concentrations of ATP to its hydrolysis products ATP+P_i **is in the order of $10^9 - 10^{11}$!!!**

- ▶ ATP is **actively driven** out of equilibrium with respect to ADP+P_i.
- ▶ This **astronomical disequilibrium** provides the “driving force”.
- ▶ The disequilibrium is **constently dissipated** to maintain structure.

Branscomb E & Russell MJ (2017), Escapement mechanisms and the conversion of disequilibria; the engines of creation, *Phys Rep* 677:1-60 | doi:10.1016/j.physrep.2017.02.001

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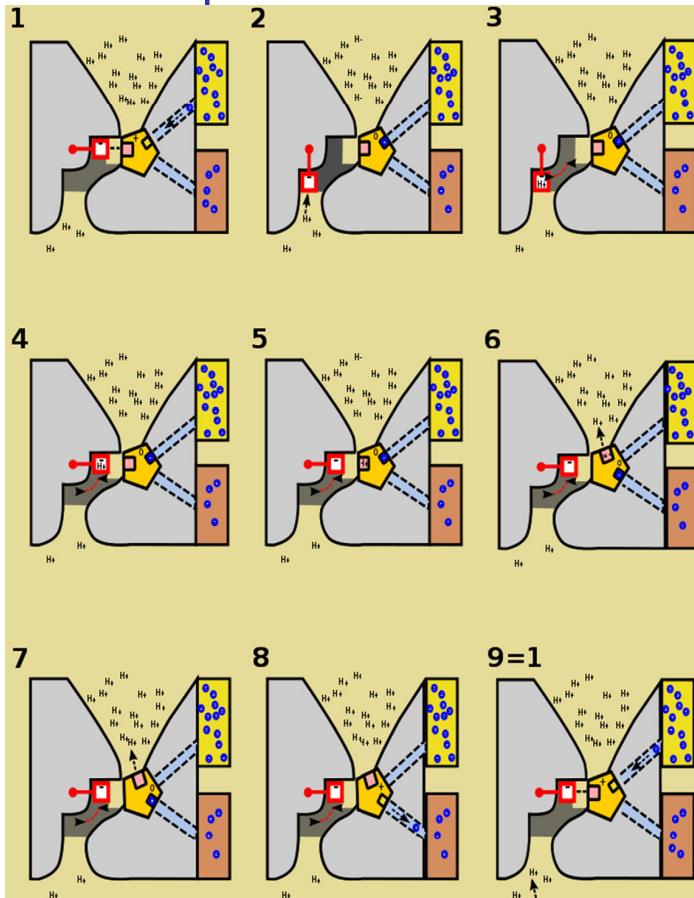
What is energy conversion?



Branscomb E & Russell MJ (2013), Turnstiles and bifurcators: The disequilibrium converting engines that put metabolism on the road, *Biochim Biophys Acta* 1827:62-78 | doi:10.1016/j.bbabi.2012.10.003

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Example: Ferredoxin I from *Azotobacter vinelandii*



Moving parts:

- **box** (CO₂⁻-moiety).
- **pentagon** (3Fe4S-cluster).

dark gray channel blocks H⁺.

yellow box redox donor.

brown box redox acceptor.

Figure modified from Branscomb & Russell 2013;

Chen K et al (2000), Atomically defined mechanism for proton transfer to a buried redox centre in a protein, *Science* 405:814-817 | doi:10.1126/science.289.5478.814

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The electron transfer puzzle

Hydrogenotrophic methanogenesis



Thermodynamic conundrum:

Crucial electron transfer steps **go steeply uphill** against a substantial unfavorable redox midpoint potential difference.

Solution:

Energy conserving electron bifurcation[†].

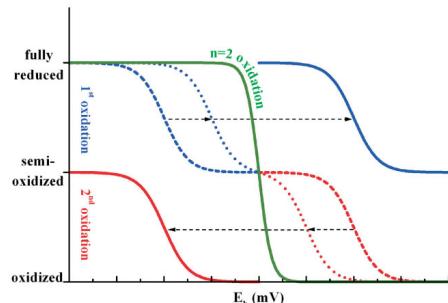
Mechanism:

The highly exergonic electron transfer to the high potential acceptor compensates the endergonic reduction of an acceptor substantially more negative than the flavin.

[†] The hypothesis put forward by Buckel, Thauer, and colleagues in essence stipulates that the 2-electron carrier, flavin, bifurcates its two electrons towards acceptors with much lower and much higher redox potentials than that of the average (2-electron) transition of flavin.

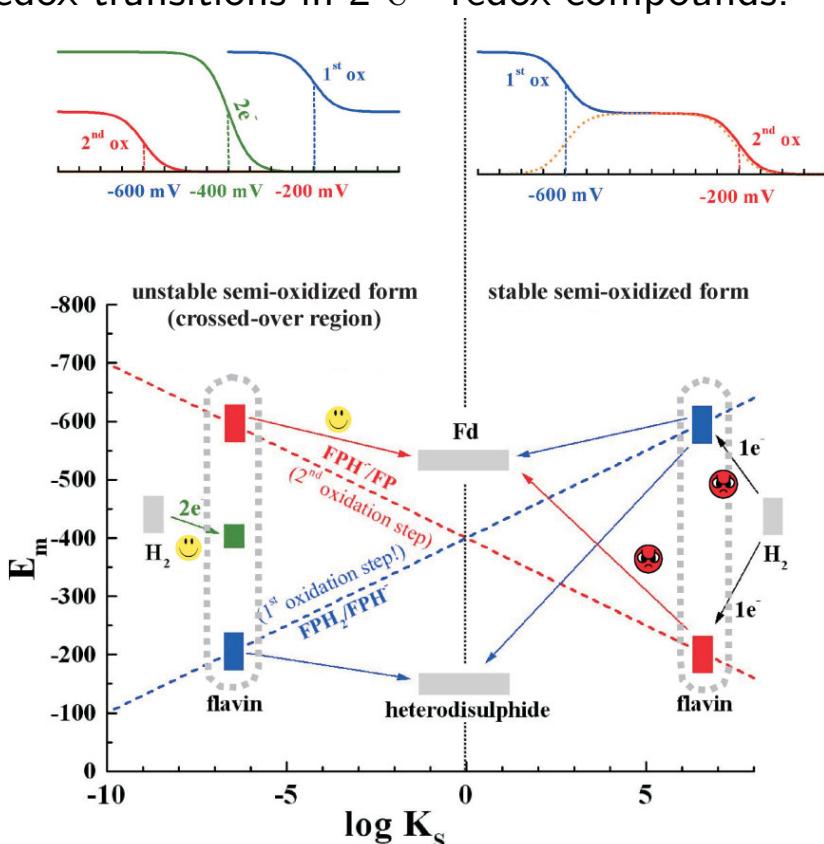
Mechanism of redox-bifurcation

Cross-over of individual redox transitions in 2-e⁻ redox compounds.



“Magic” Cofactors:

- ▶ flavins
- ▶ quinones



Figures modified from Nitschke W & Russell MJ (2011), Redox bifurcations: Mechanisms and importance to life now, and at its origin, *Bioessays* 34:106-109 | doi:10.1002/bies.201100134

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Example: Methanogenic heterodisulfide reductase complex

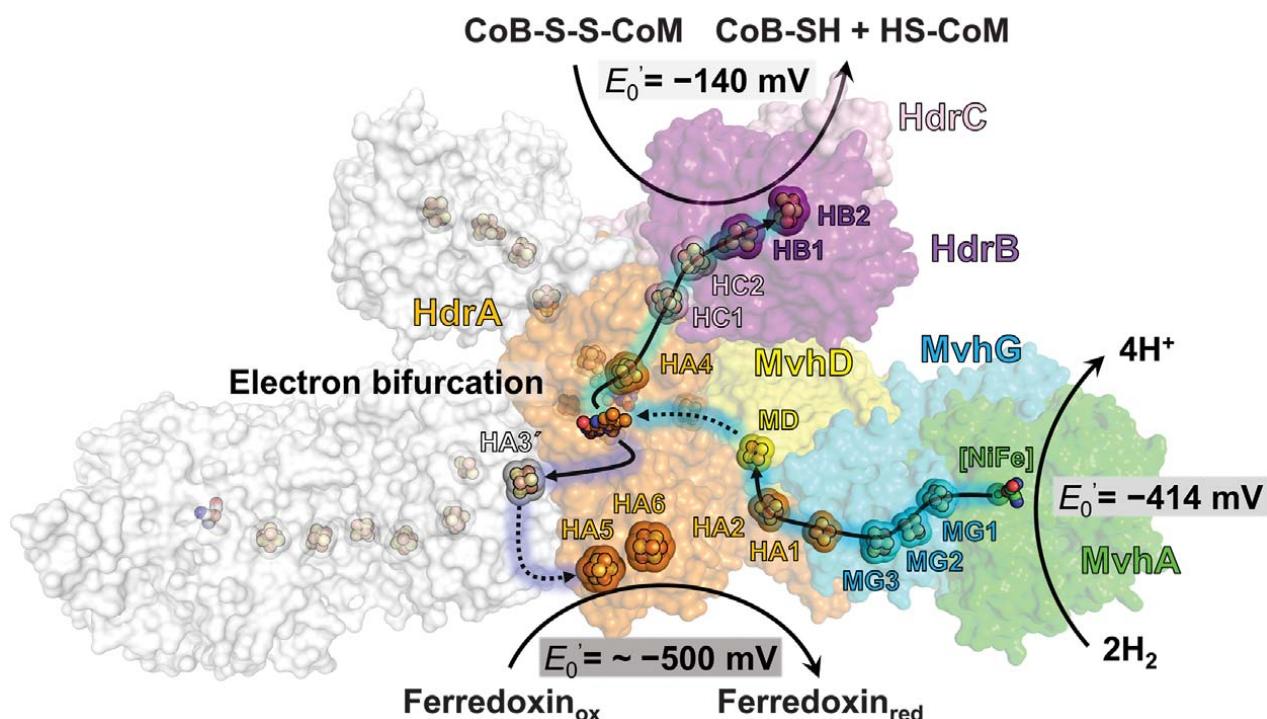


Figure modified from Wagner T et al (2017), Methanogenic heterodisulfide reductase (HdrABC-MvhAGD) uses two noncubane [4Fe-4S] clusters for reduction, *Science* 357:1-4 | doi:10.1126/science.aan0425

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Further Reading

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-  Domagal-Goldman SD, Wright KE, Adamala K, Arina de la Rubia L, Bond J, Dartnell LR, Goldman AD, Lynch K, Naud ME, Paulino-Lima IG, Singer K, Walter-Antonio M, Abrevaya XC, Anderson R, Arney G, Atri D, Azúa-Bustos A, Bowman JS, Brazelton WJ, Brennecke GA, Carns R, Chopra A, Colangelo-Lillis J, Crockett CJ, DeMarines J, Frank EA, Frantz C, de la Fuente E, Galante D, Glass J, Gleeson D, Glein CR, Goldblatt C, Horak R, Horodyskyj L, Kaçar B, Kereszturi A, Knowles E, Mayeur P, McGlynn S, Miguel Y, Montgomery M, Neish C, Noack L, Rugheimer S, Stüeken EE, Tamez-Hidalgo P, Imari Walker S & Wong T. **The Astrobiology Primer v2.0.** *Astrobiol* (2016), 16:561-653 | doi:10.1089/ast.2015.1460
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