

# tracking the rise of eukaryotes— geochemical proxies & genomic perspectives

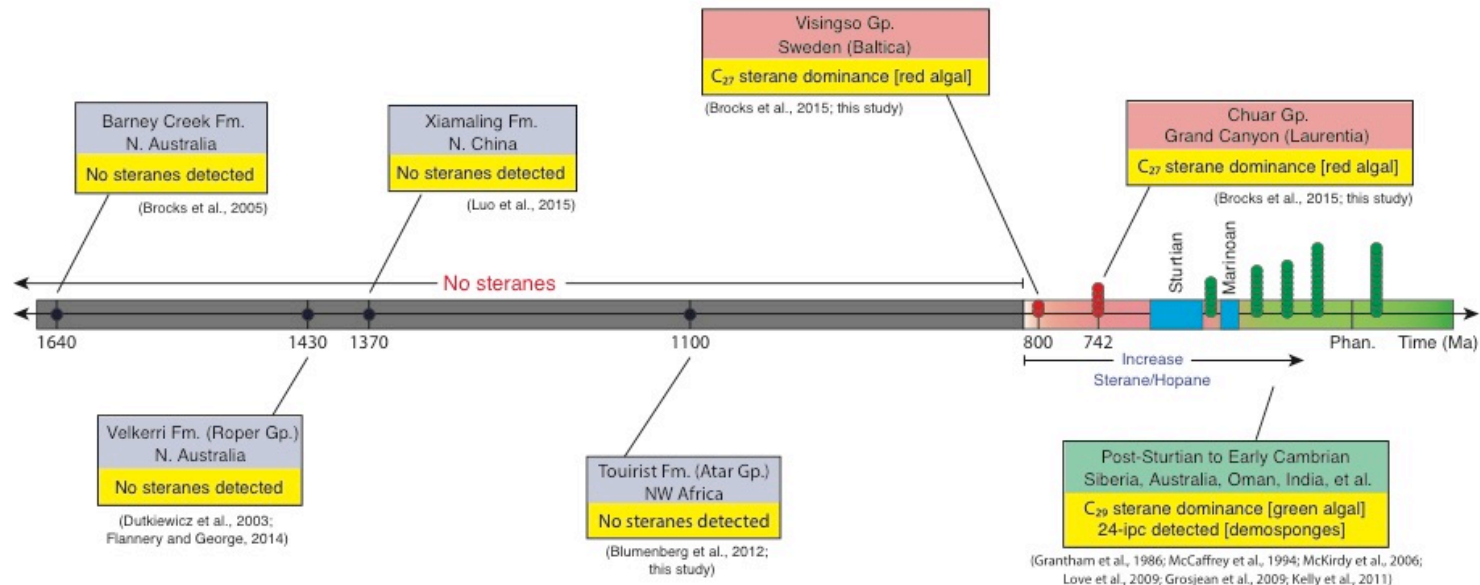
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# tracking the rise of eukaryotes – geochemical proxies

- Limited number of thermal immature Proterozoic unit
- Poor constraints on when the transition to extensive eukaryotic export productivity
- Biomarker records provide local signals

## *Proterozoic Biomarker Record*



## tracking the rise of eukaryotes – geochemical proxies

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Can we use Zn isotopes to track  
Eukaryotes Rise to Prominence?

# biological zinc requirements

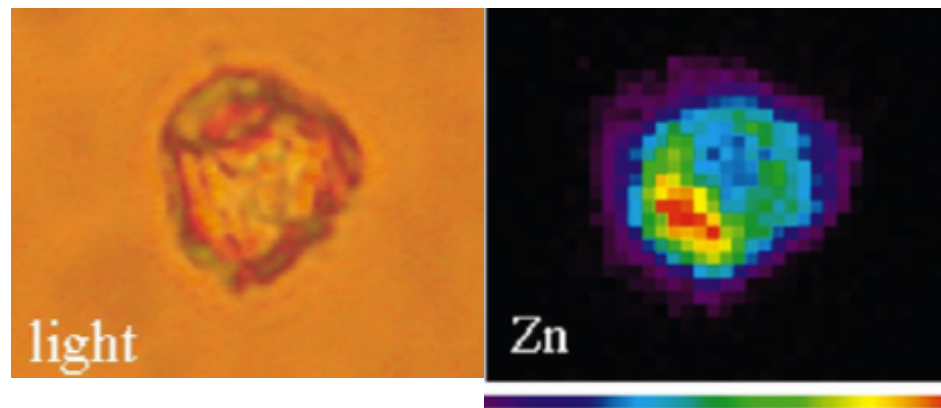
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Modern Phytoplankton Zn : C ( $\mu\text{mol/mol}$ )

Eukaryotes :  $102 \pm 20$  (Ave) - (Twining et al., 2003; 2004)

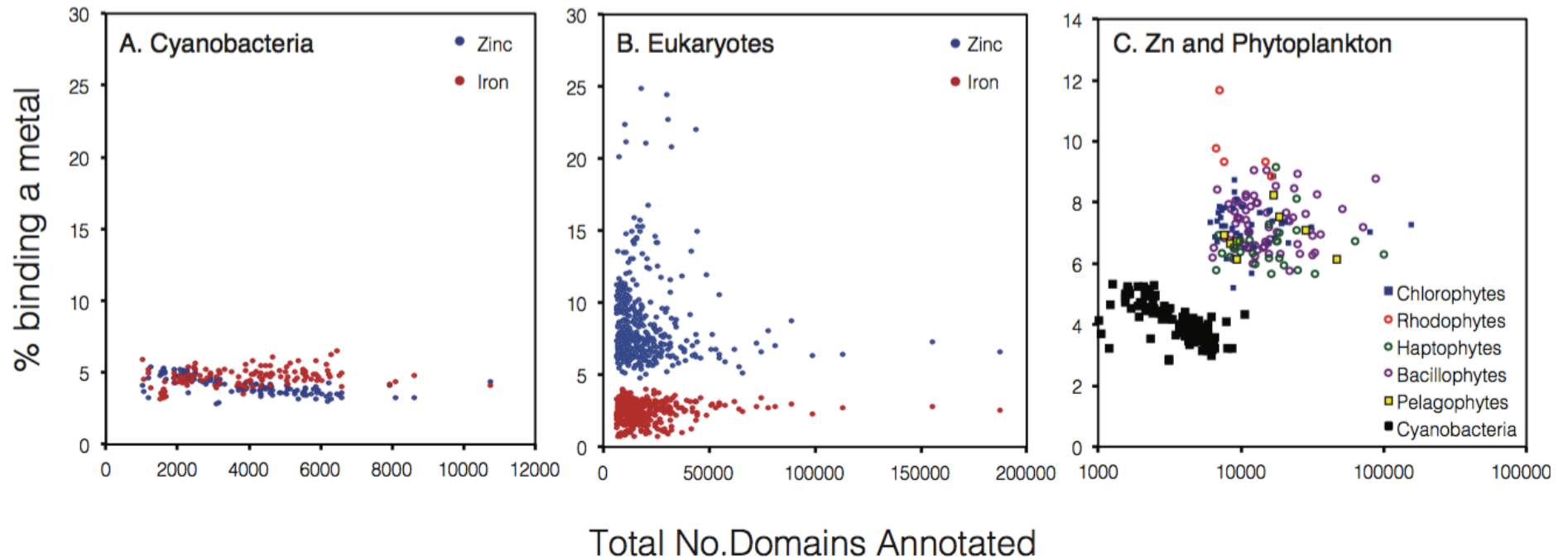
Cyanobacteria :  $17 \pm 14$  (Ave.) - (Nuester et al., 2012)

Much of this difference comes from proteins (e.g Zn Fingers)  
found within the nucleus



SXRF element maps of a centric diatom - *Southern Ocean*

# A Census of Zinc Binding Protein Domains (Structural Bioinformatics & Whole Genome Sequences)

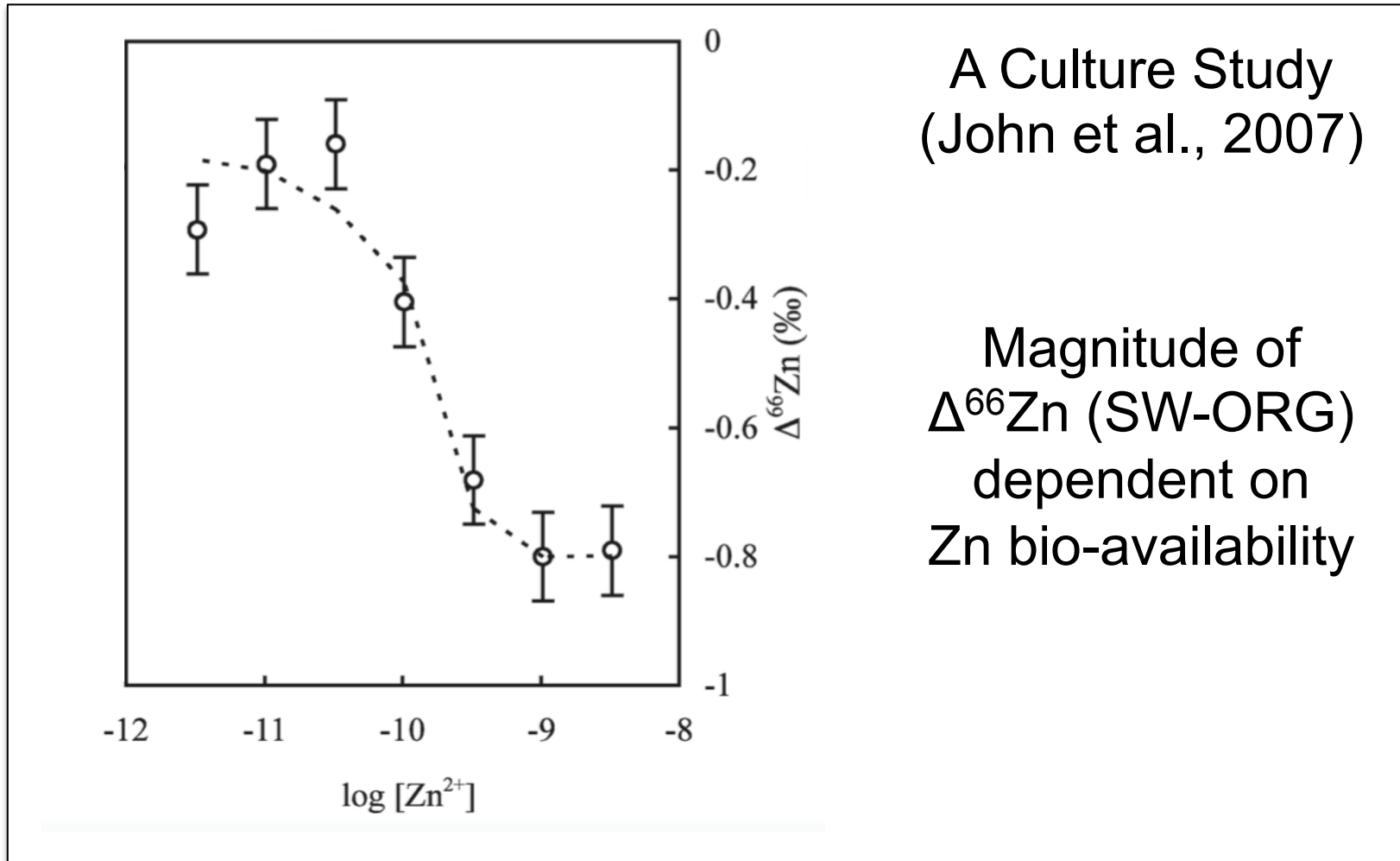


- Database: SCOP; Cyano-GEBA; MMETSP
- Whole genome sequences of known cyanobacteria and eukaryotes
- Determine the number of metal binding domains coded for in each proteome
- Zinc is one of the most abundant inorganic co-factor in eukaryotic enzymes – a fundamental genetic difference from prokaryotes

(Terry Tang et al., – *in review*)

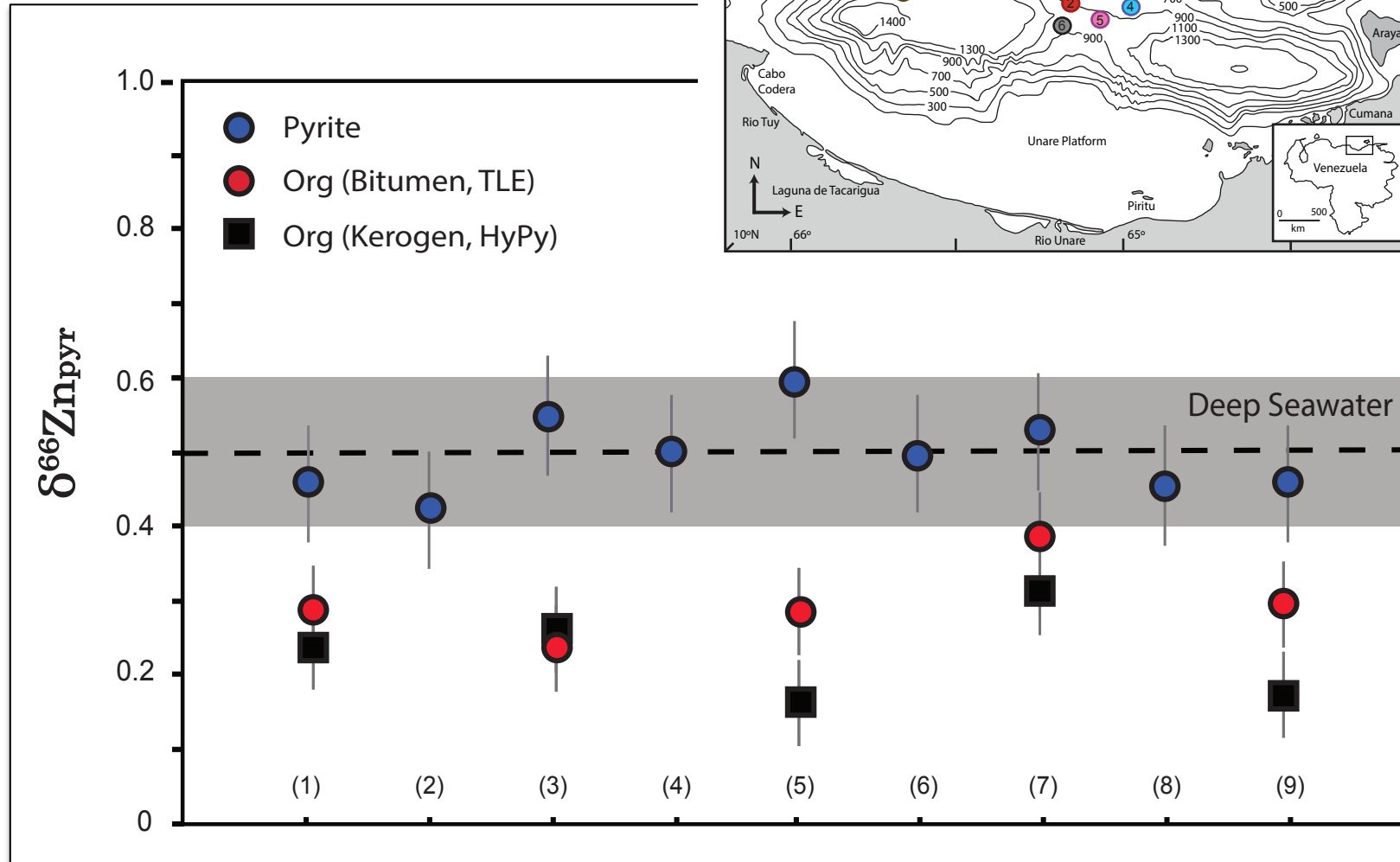
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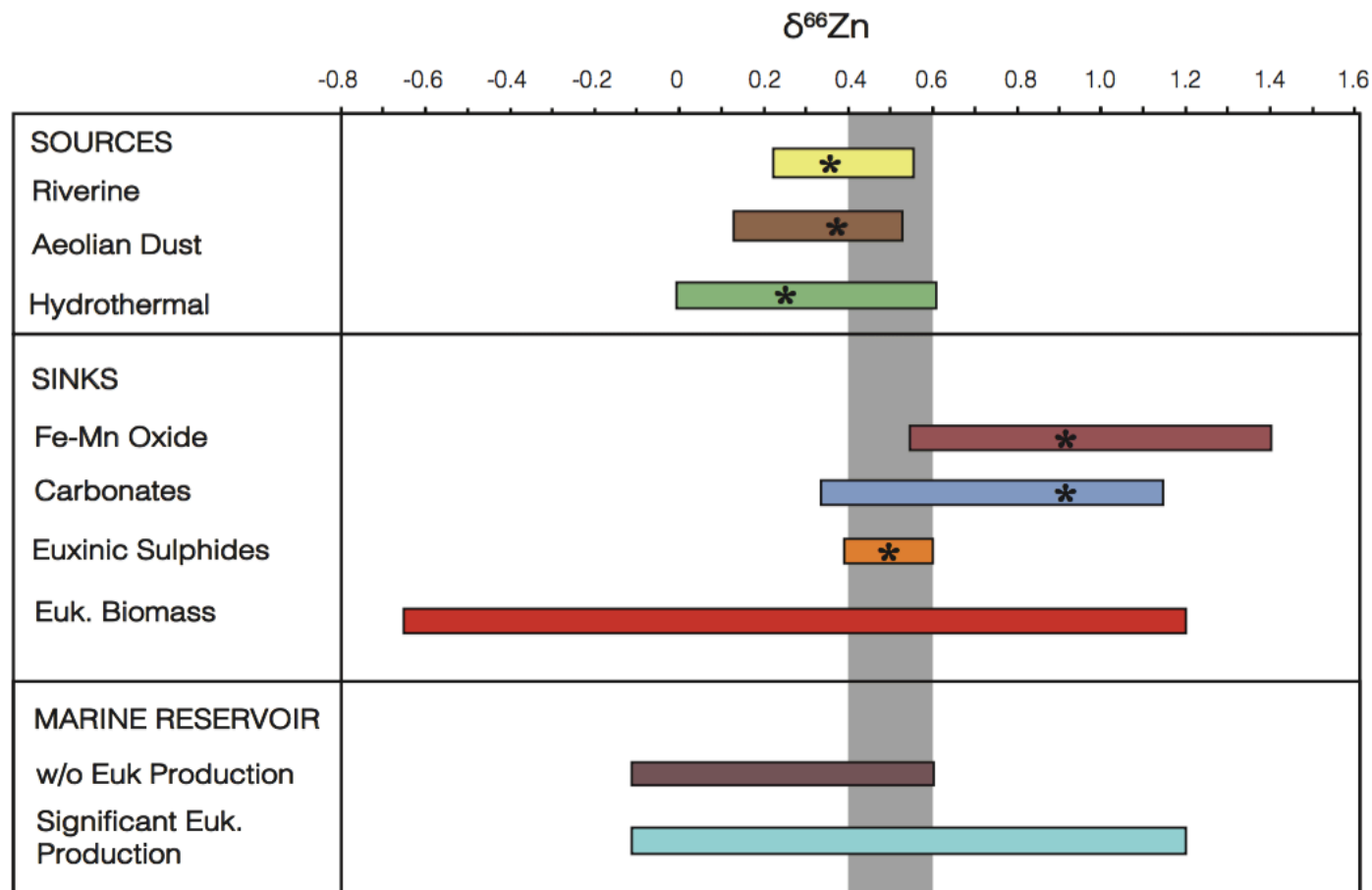
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## Core Top Calibration Cariaco Basin (Euxinic)



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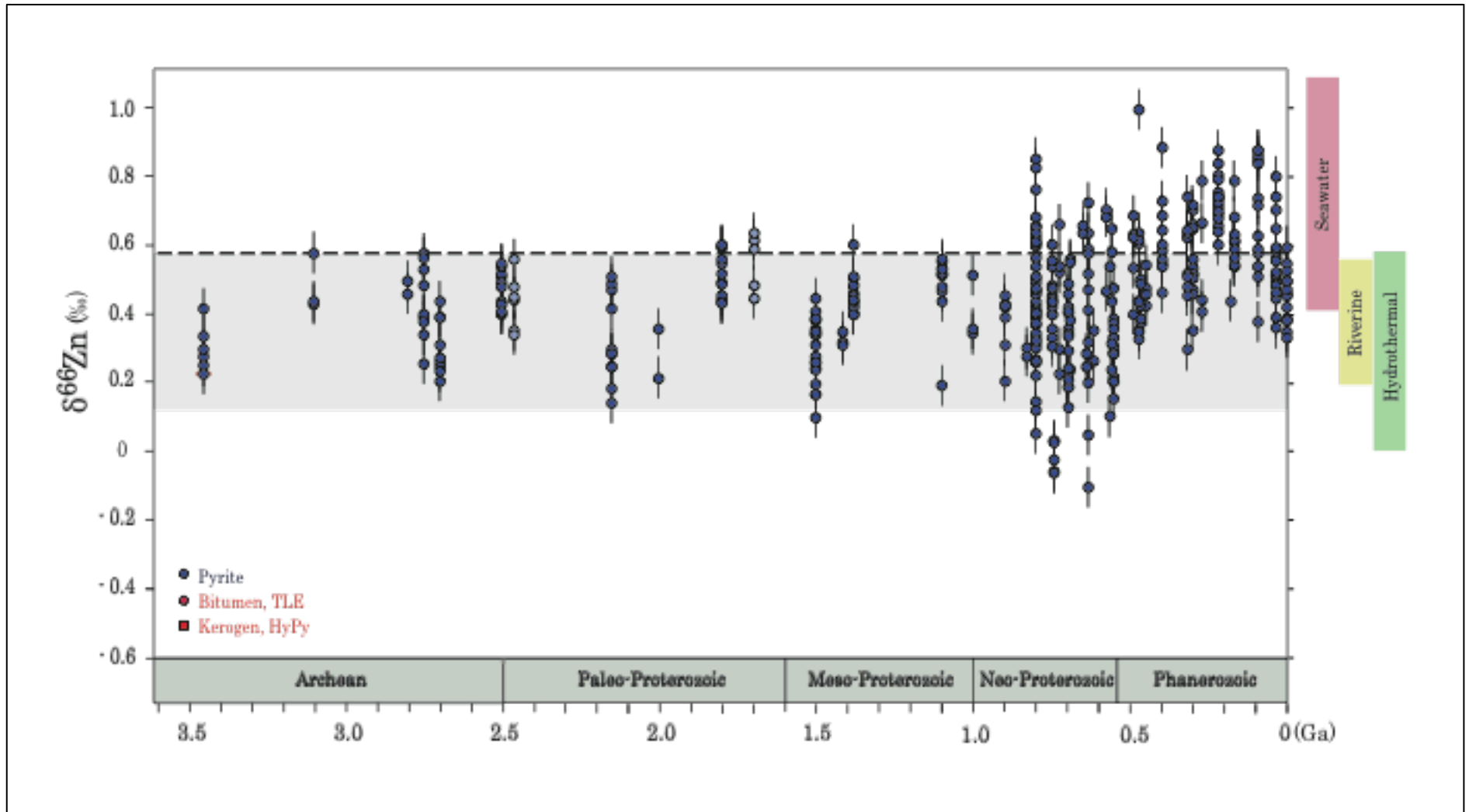
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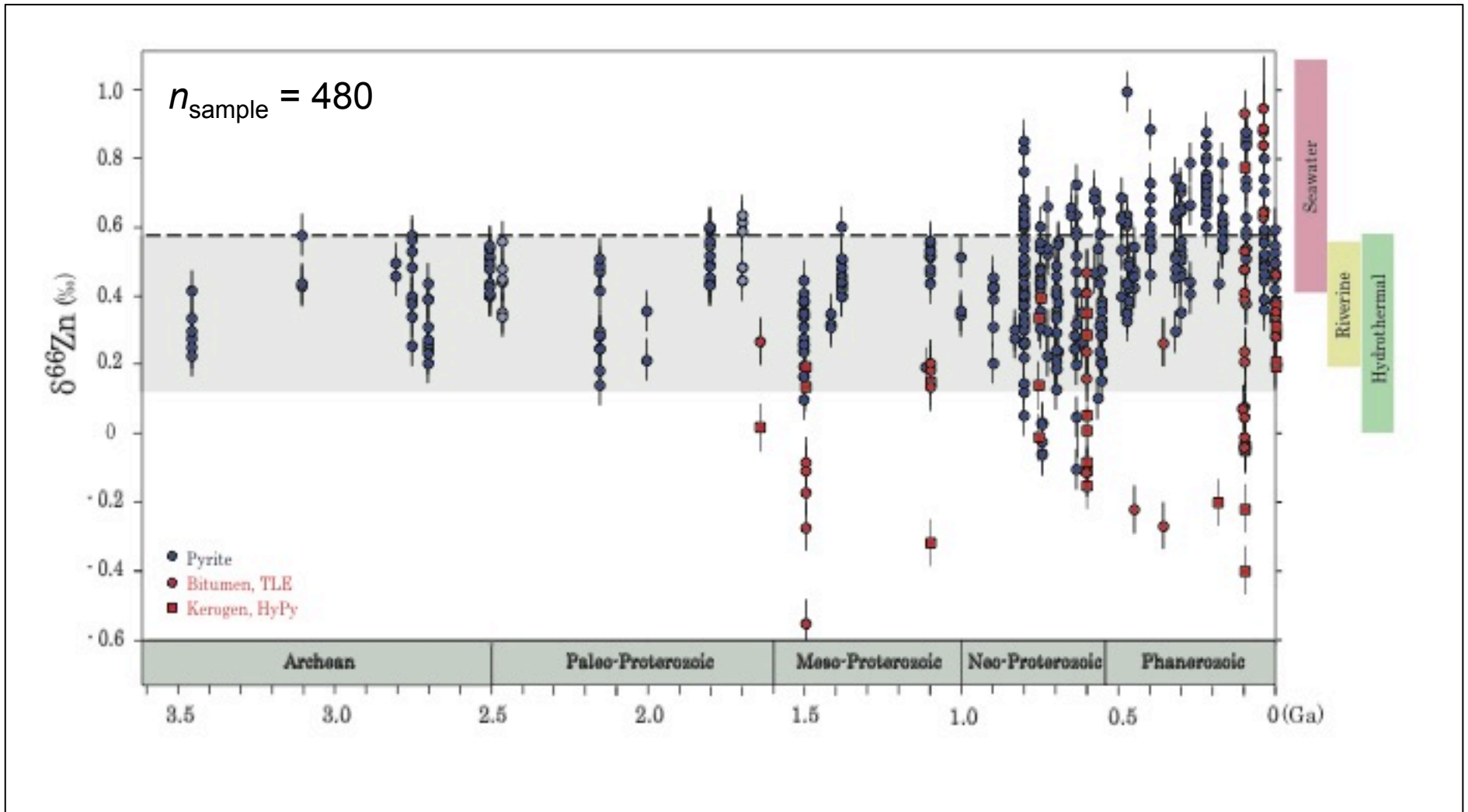
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## A Mid Neoproterozoic Ecological Revolution



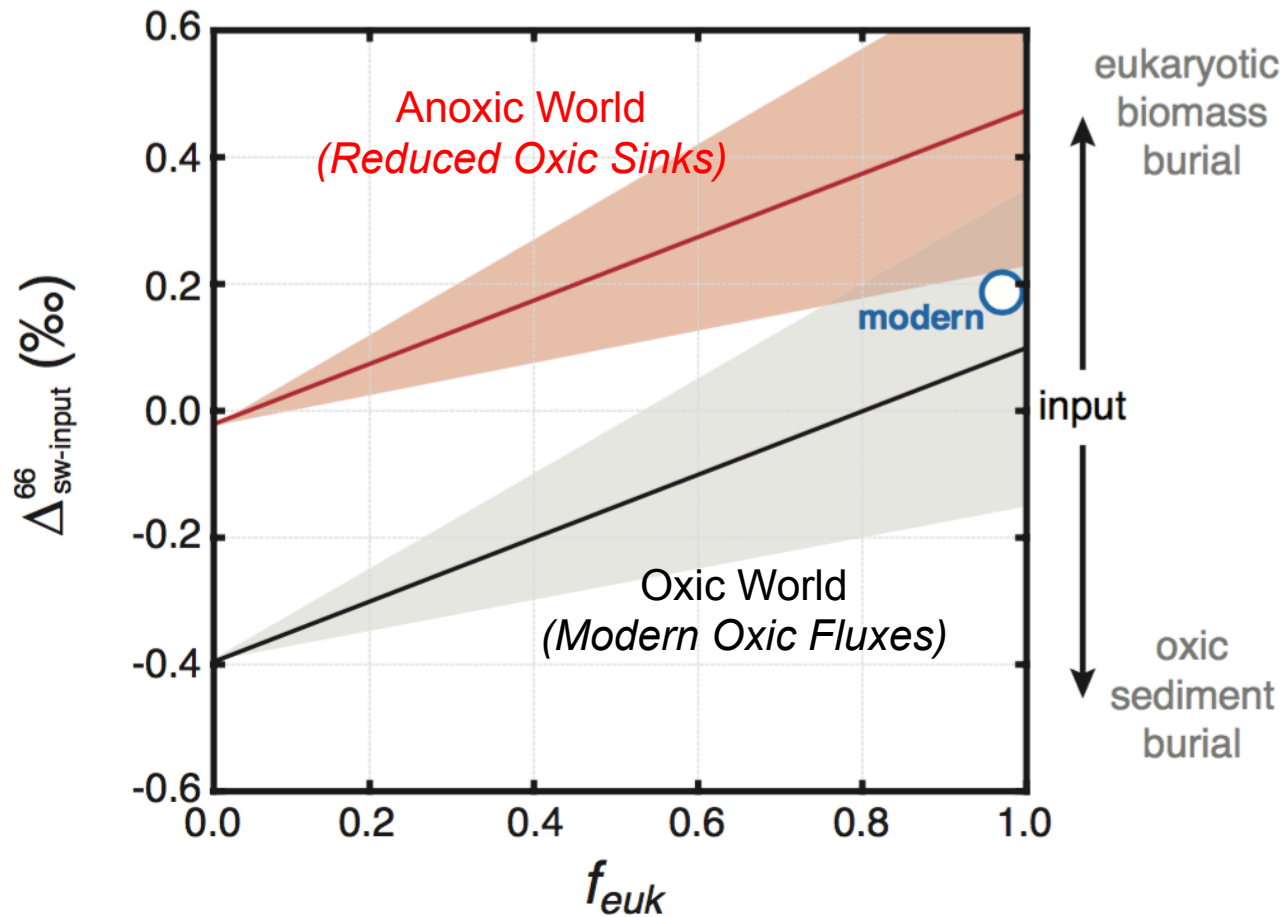
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## A Mid Neoproterozoic Ecological Revolution



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$$f_{euk} = \frac{Euk}{Euk + Sulph}$$

# tracking the rise of eukaryotes – geochemical proxies

- 1) Zn isotopes provide a powerful method of tracking ecological shifts between cyanobacterial and eukaryotic productivity
- 2) 800 Ma Ecological Revolution
  - 80 Myrs before the onset of Cryogenian Snowball events
  - Coincident with the onset of significant carbon cycle perturbations

