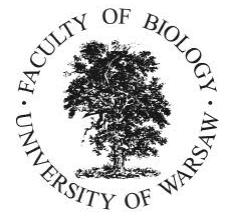




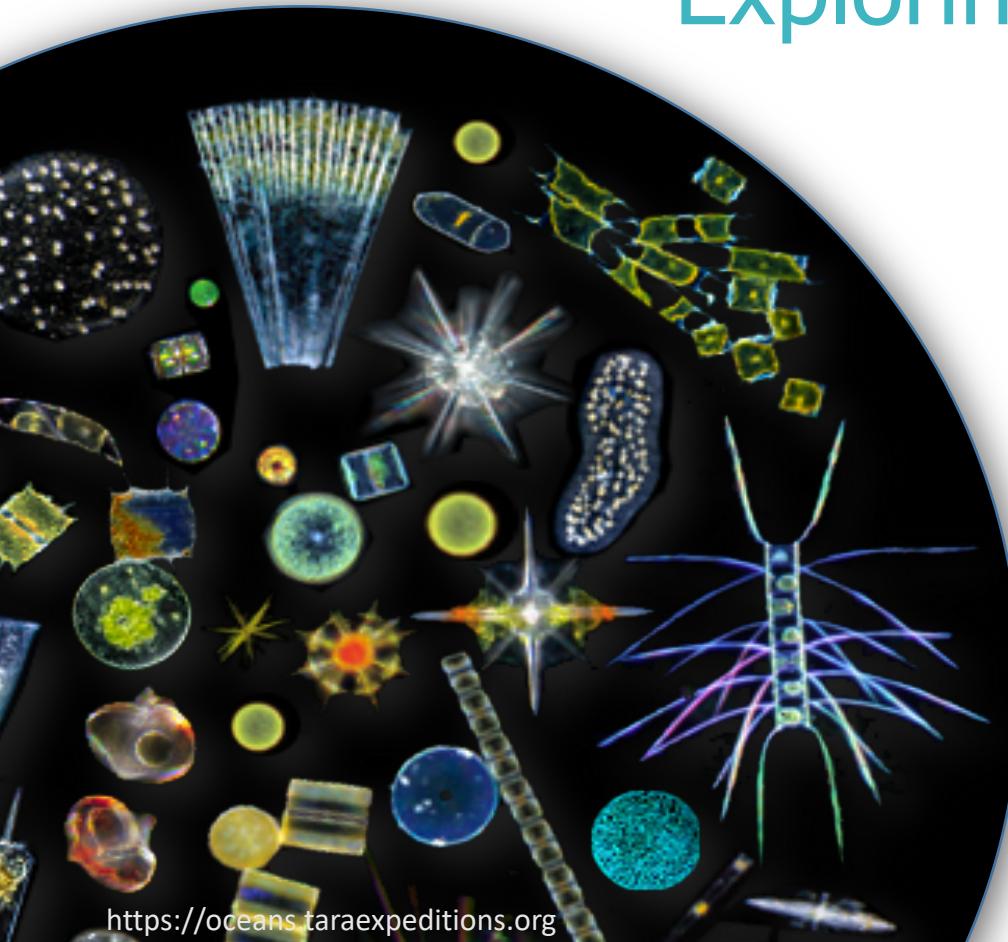
UNIVERSITY
OF WARSAW



Institute
of
Evolutionary
Biology

Exploring microbial diversity to understand the evolution of eukaryotic cell

Anna Karnkowska



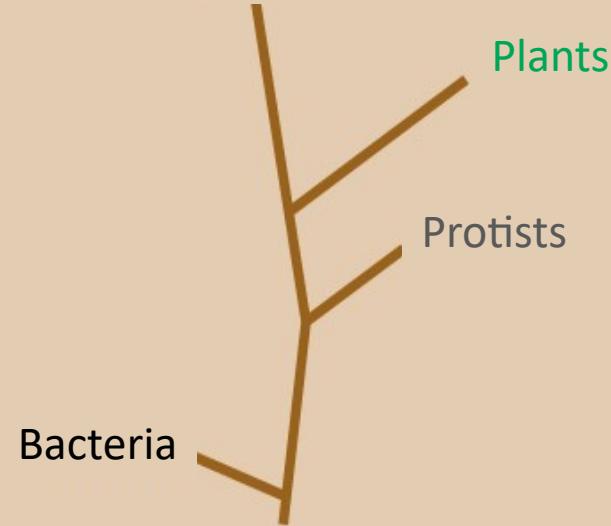




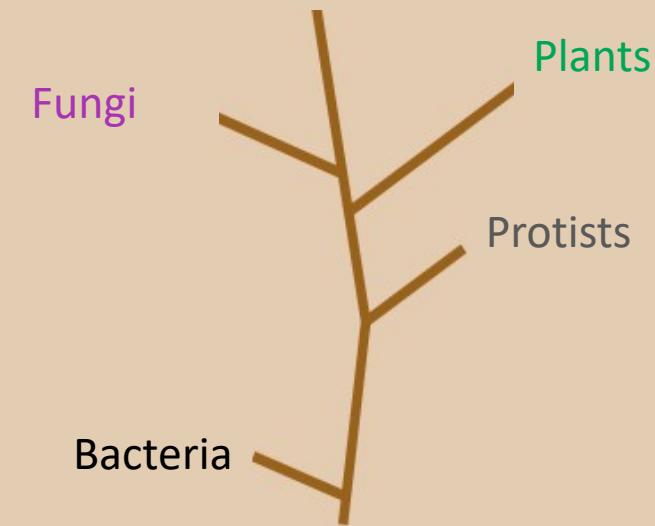
Animals



Animals



Animals



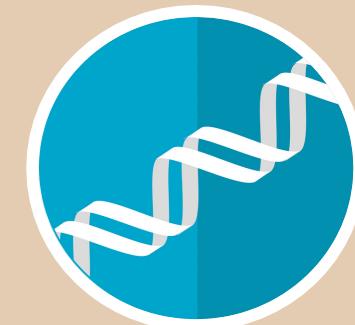
1758

Carolus Linnaeus



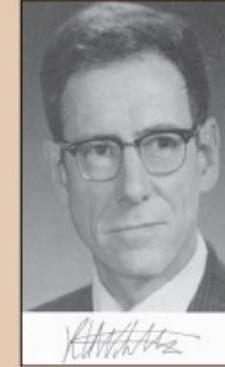
1866

Ernst Haeckel

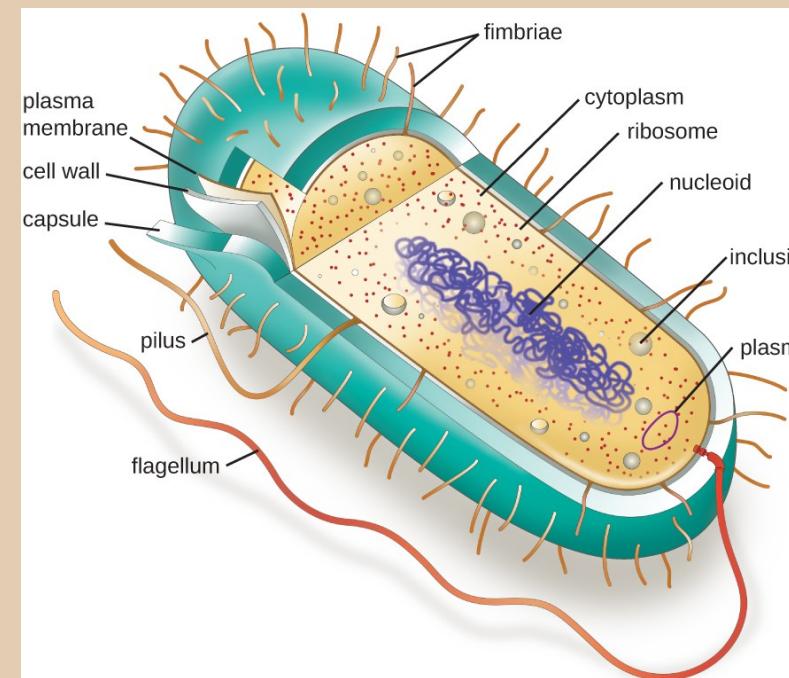
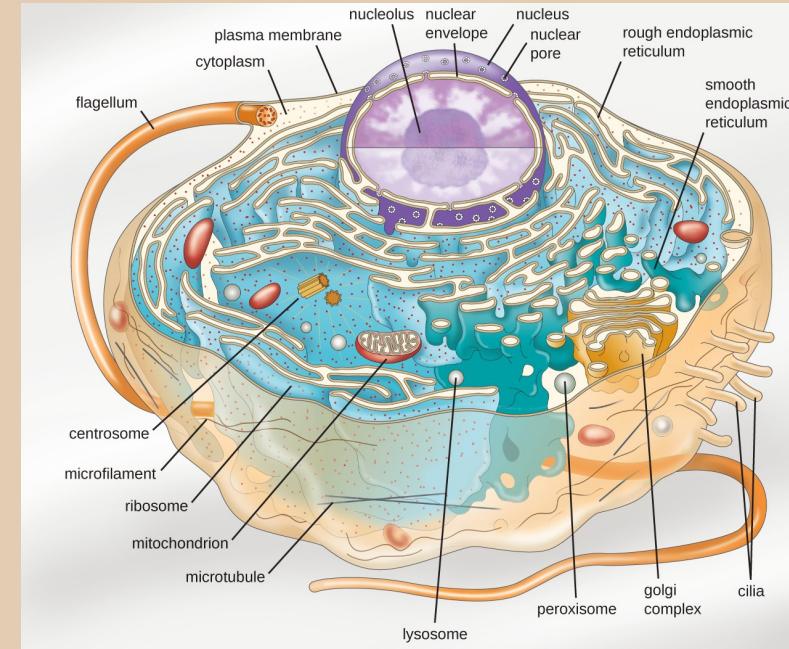
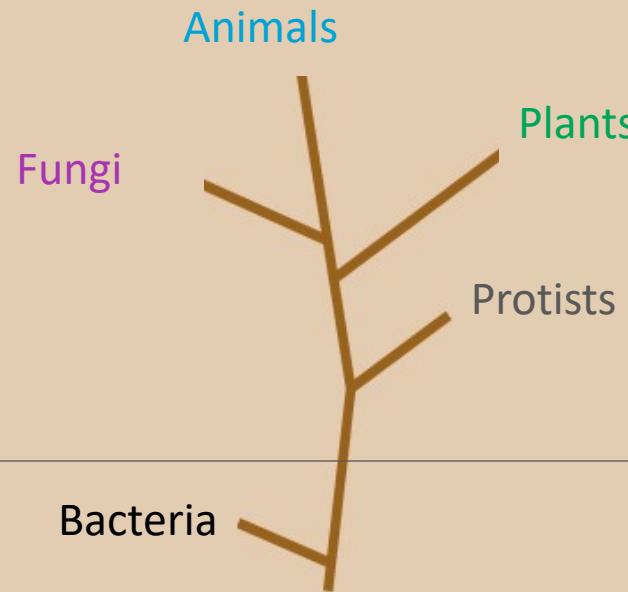


1969

Robert Whittaker



eukaryotes



prokaryotes





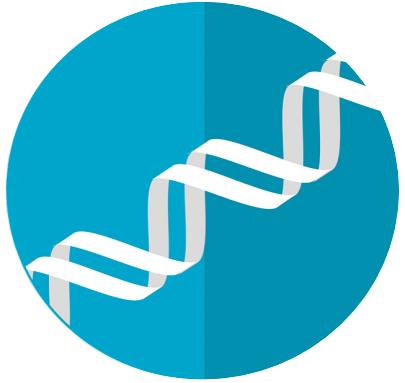
multicellular
eukaryotes

microbial eukaryotes

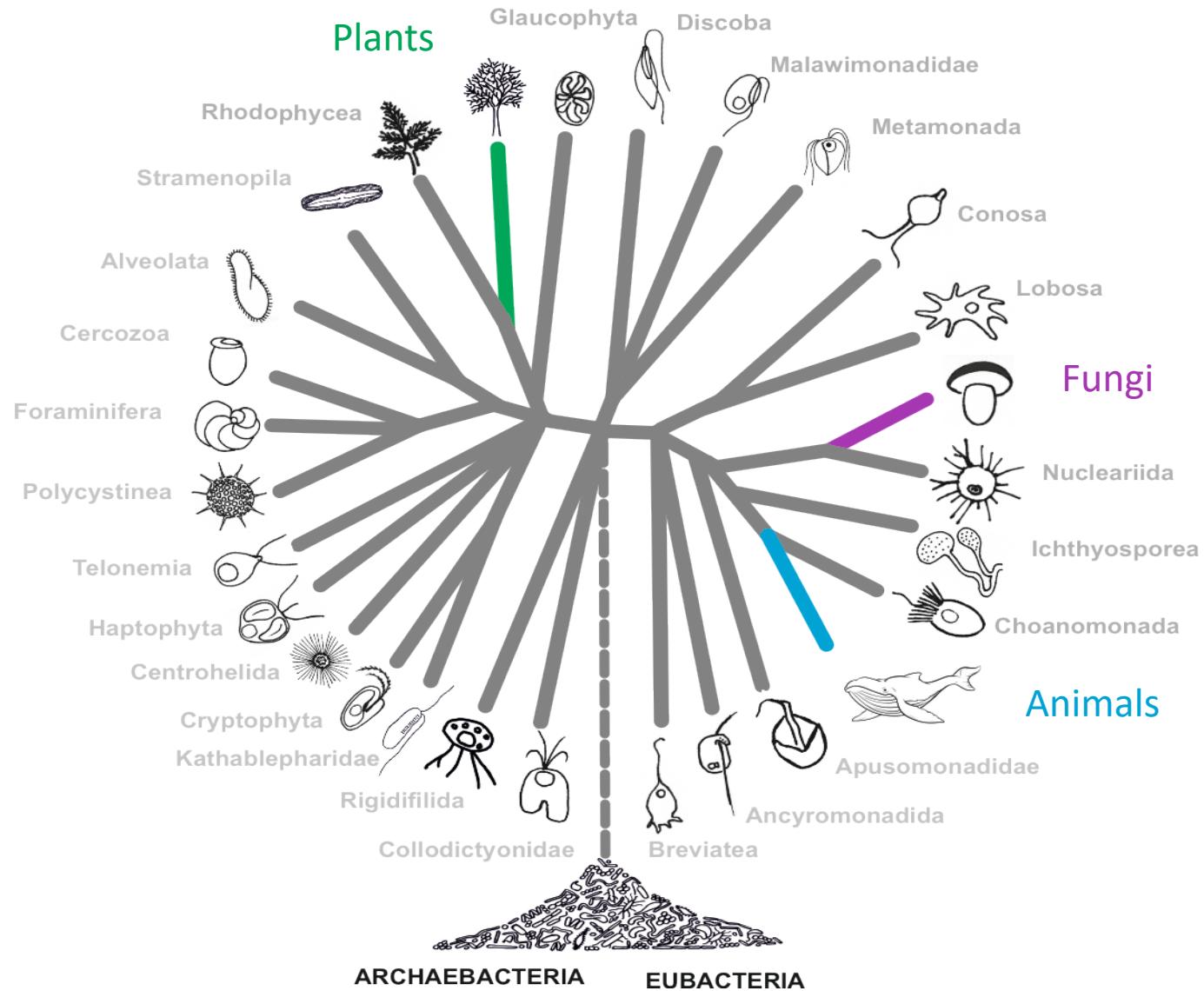
microbial eukaryotes = protists



<https://oceans.taraexpeditions.org>



diversity of eukaryotes

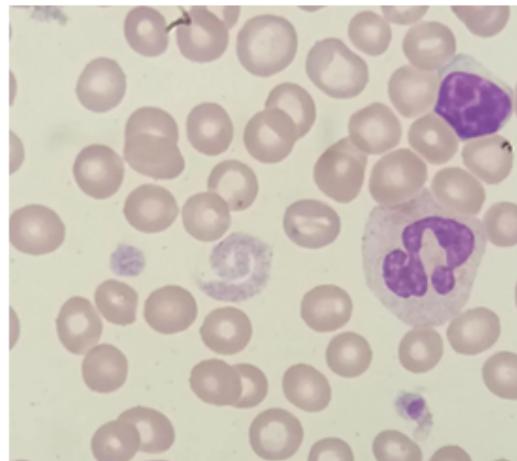


The view of a cell by:

a lab manager



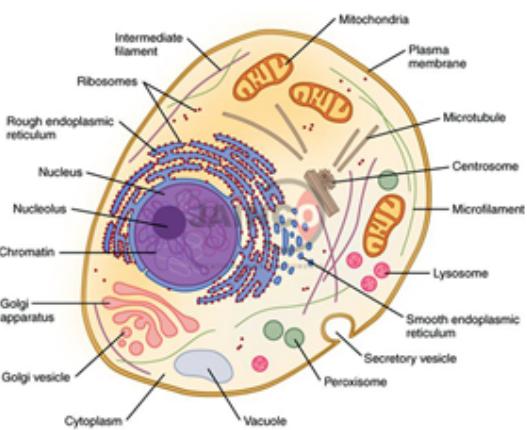
a med lab technician



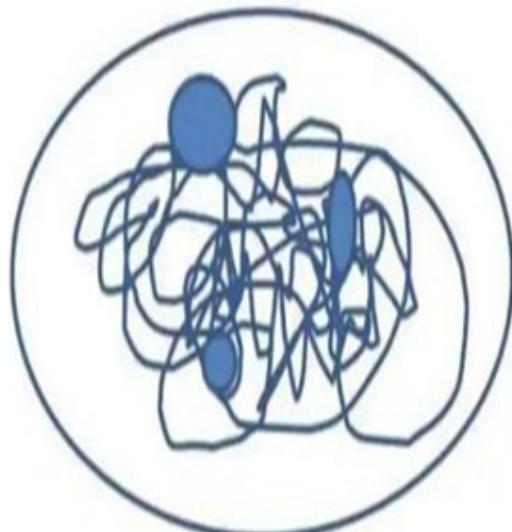
a microbiologist



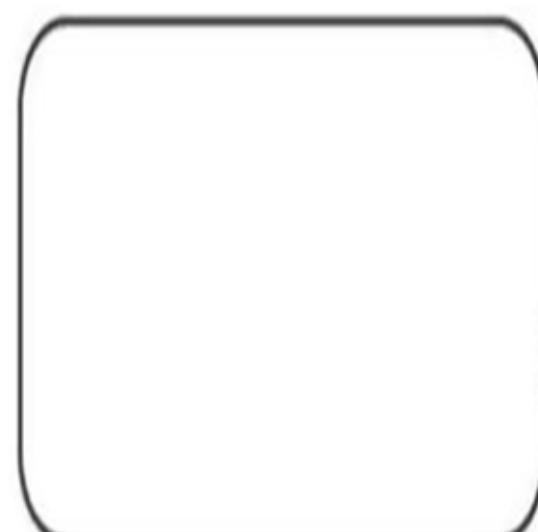
a cell biologist

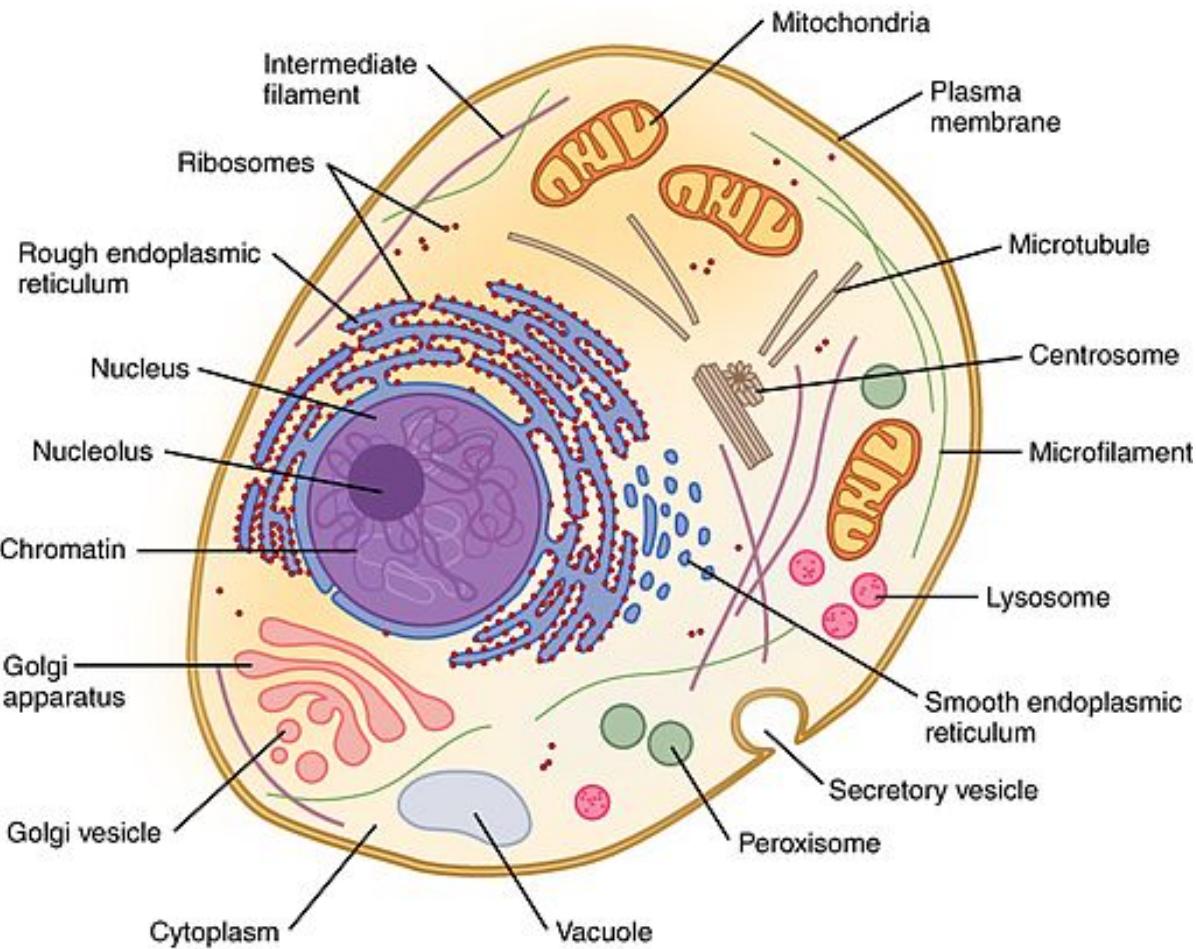


a chemist



a physicist

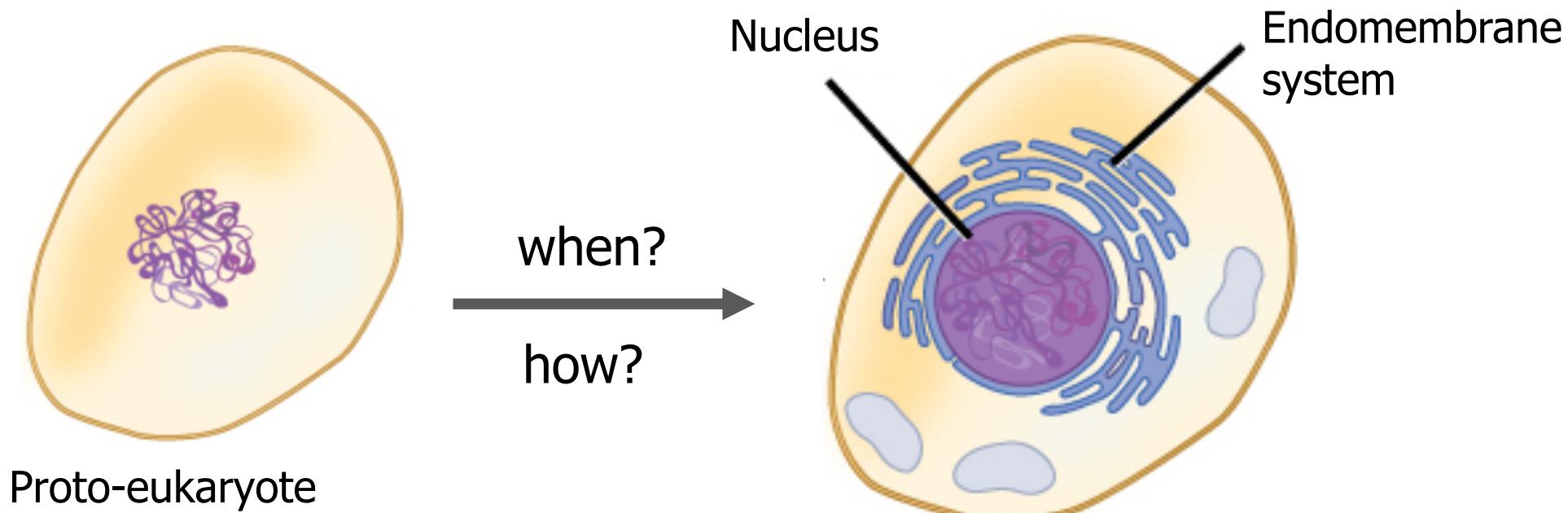




How the eukaryotic cell complexity evolved?

early evolution of eukaryotes

emergence of nucleus



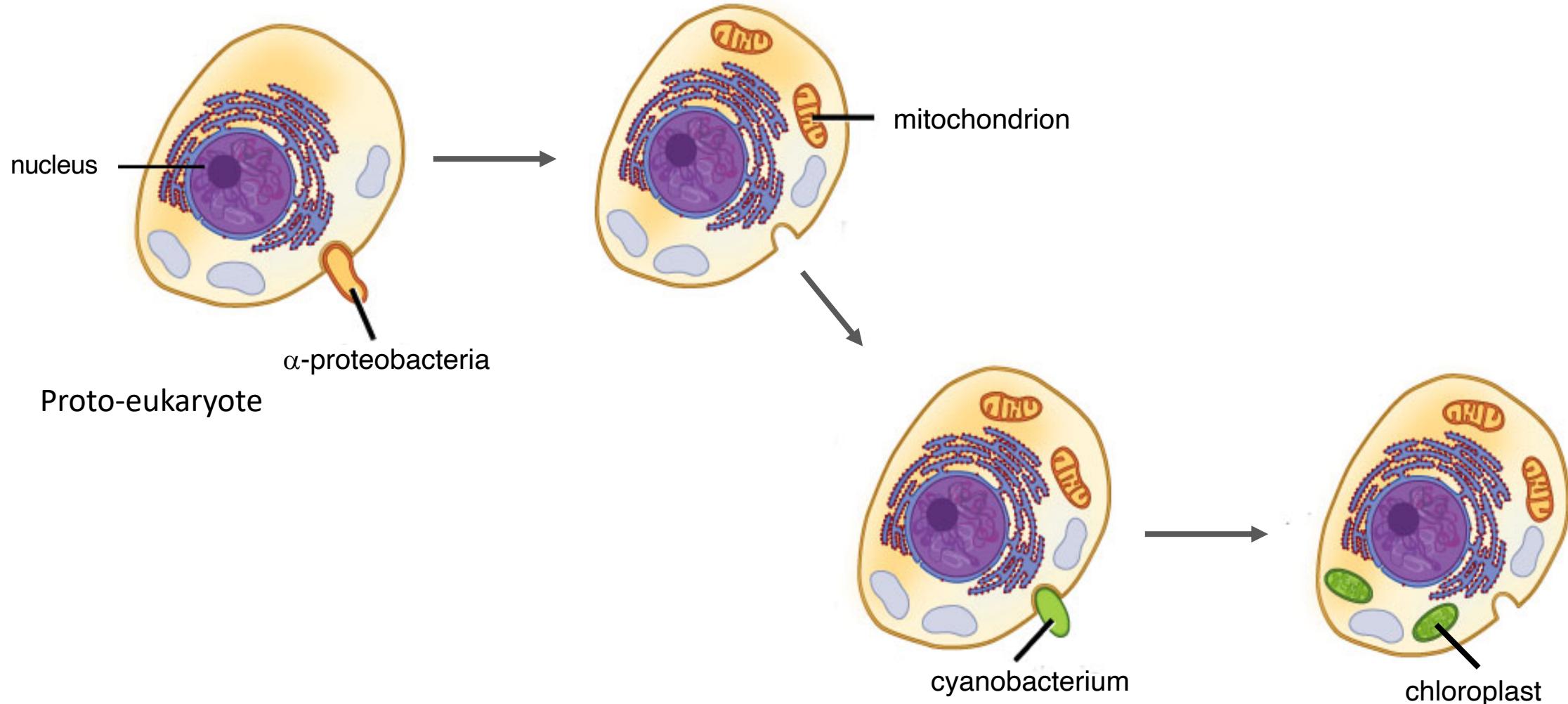
Which **came first**, the **nucleus** or the **mitochondrion**?

symbiosis



By Jan Derk - Own work, Public Domain, <https://commons.wikimedia.org/w/index.php?curid=1728598>

endosymbiotic origin of organelles



endosymbiotic organelles

mitochondrion



carbohydrates
metabolism

amino acid
metabolism

lipids
metabolism

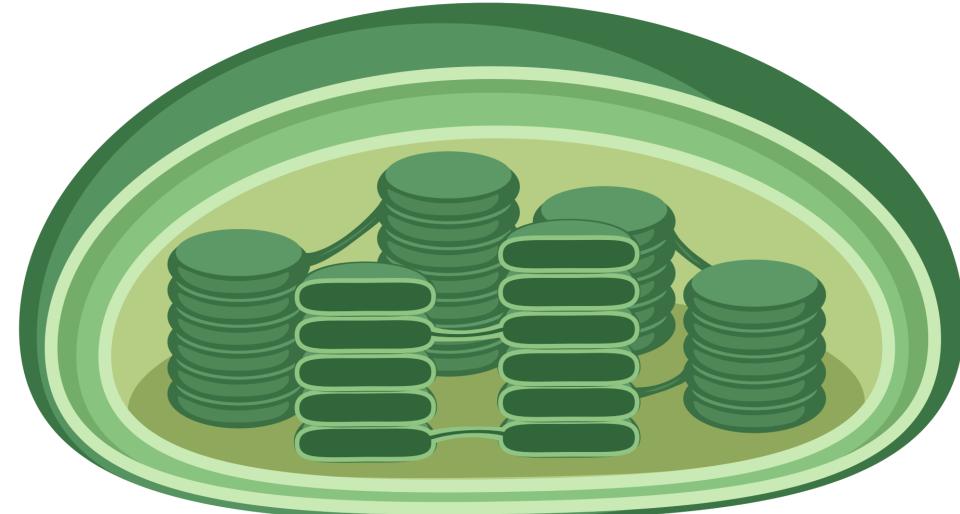
nucleotides
metabolism

Fe-S
synthesis

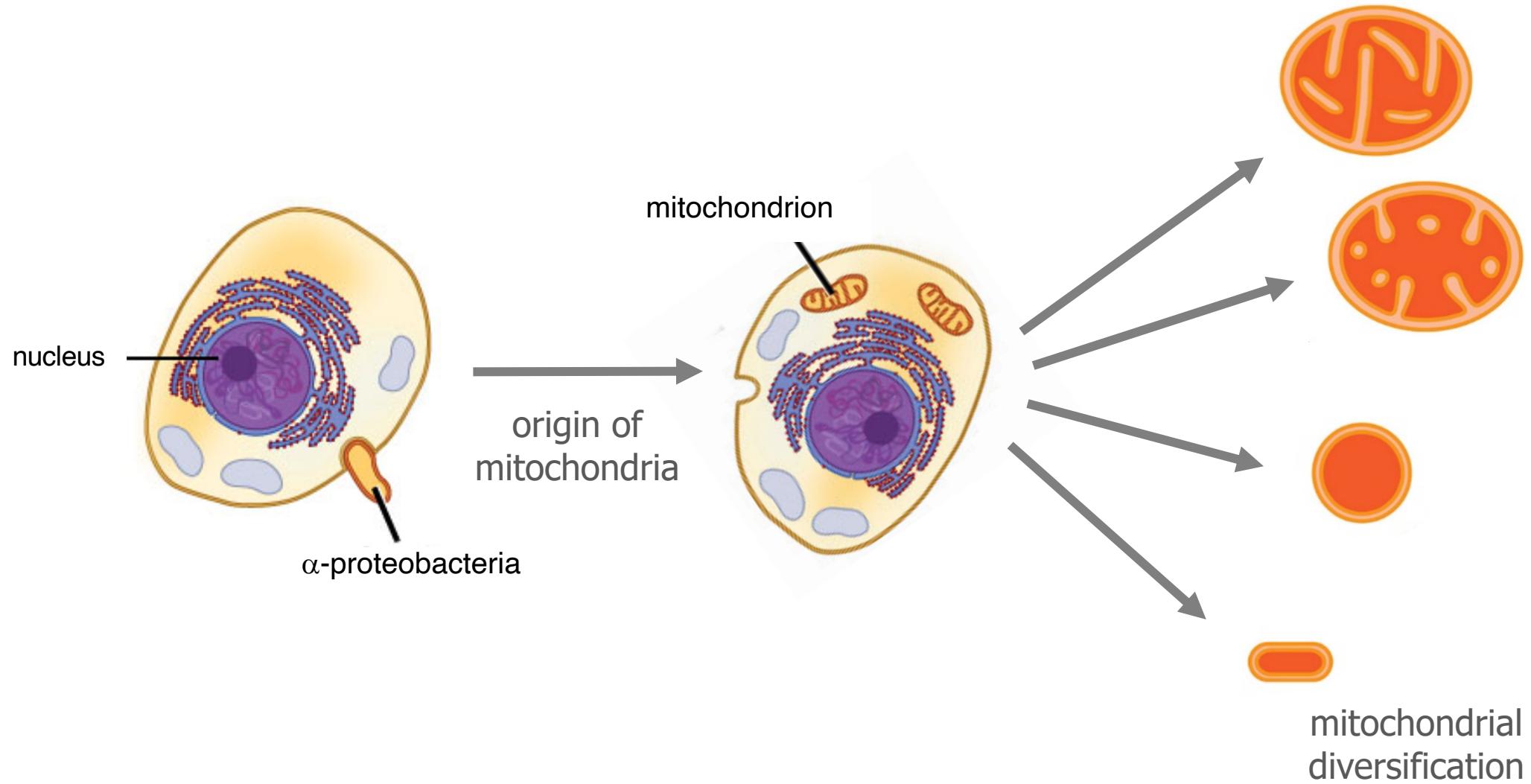
Heme
synthesis



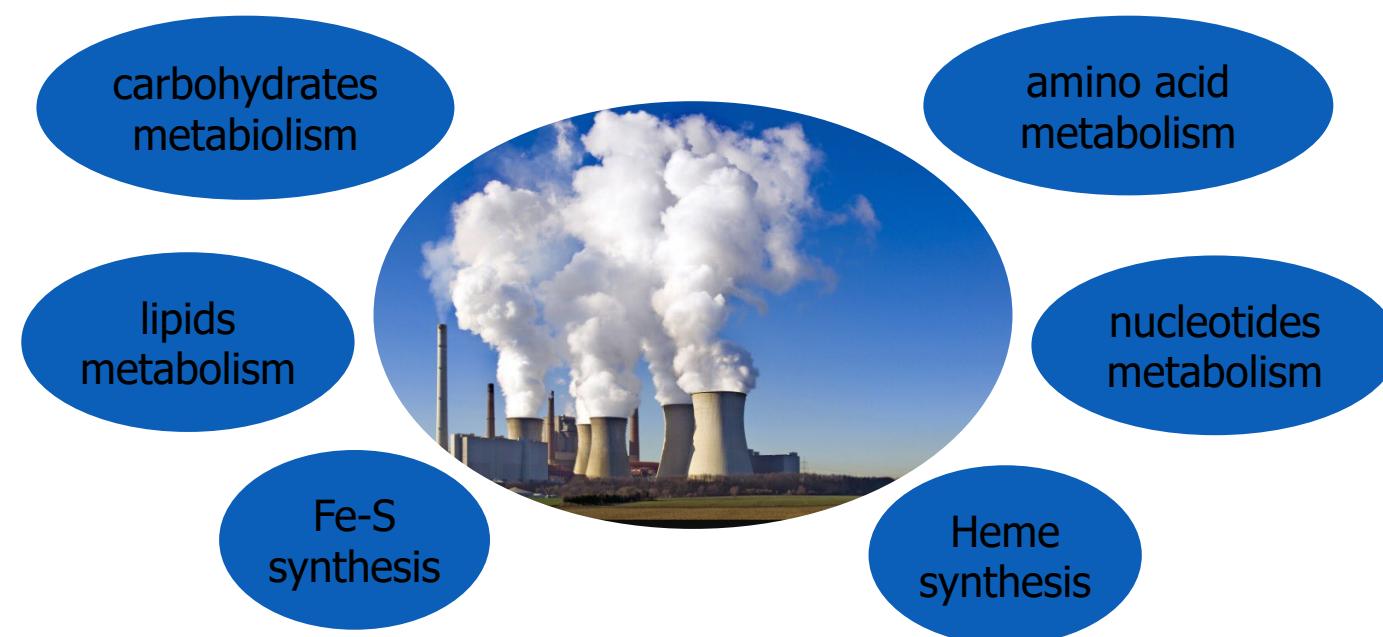
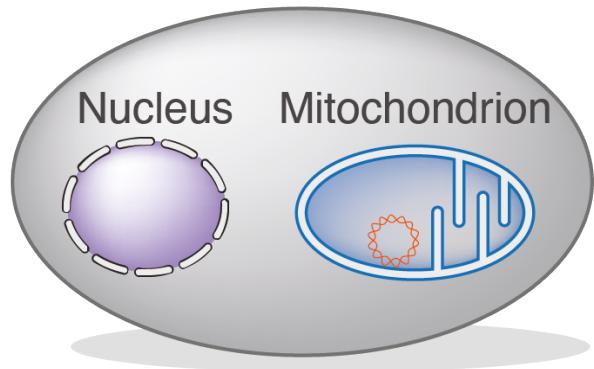
chloroplast



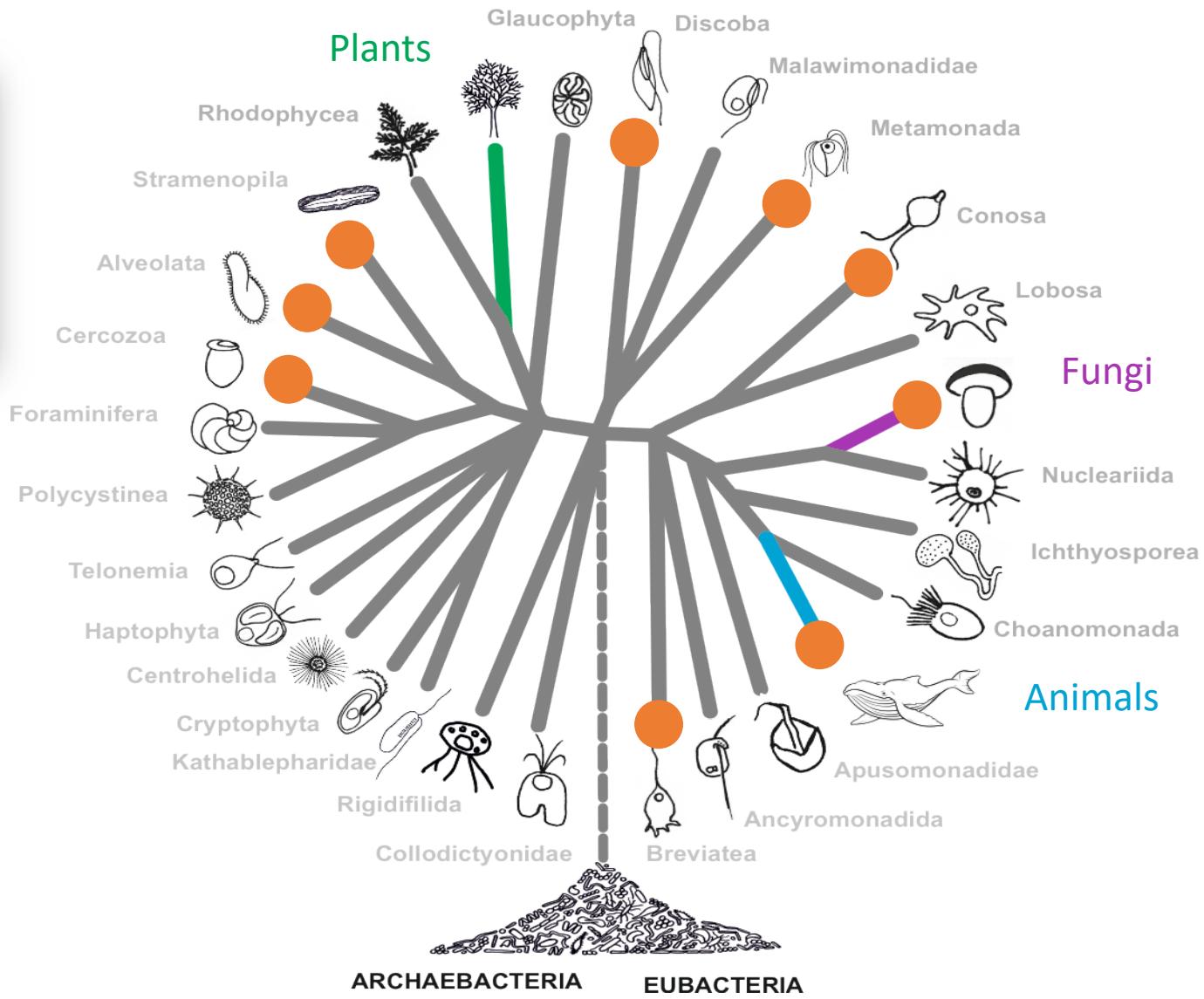
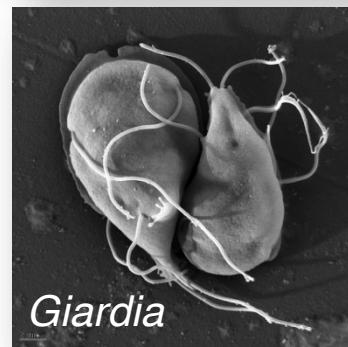
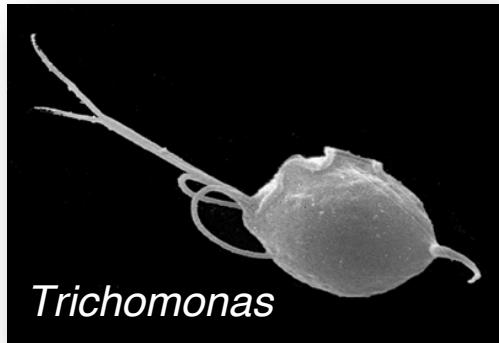
evolution of endosymbiotic organelles



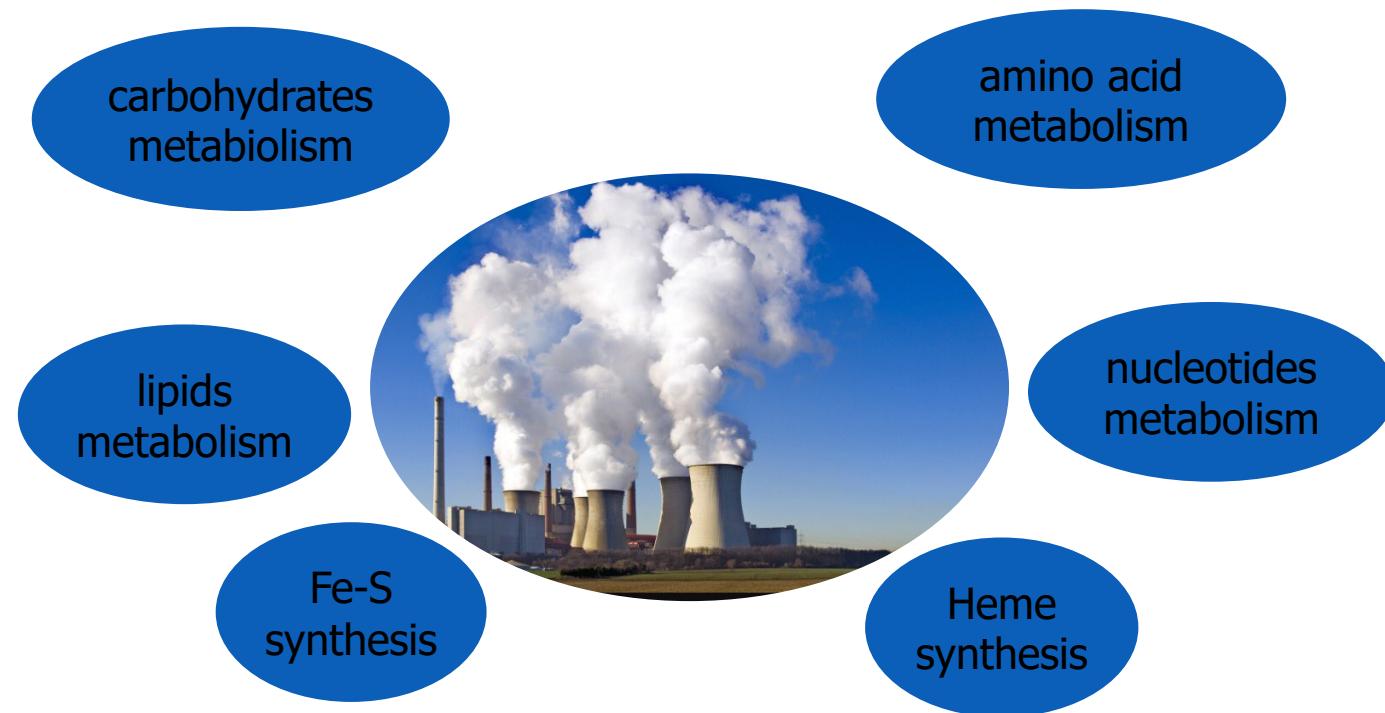
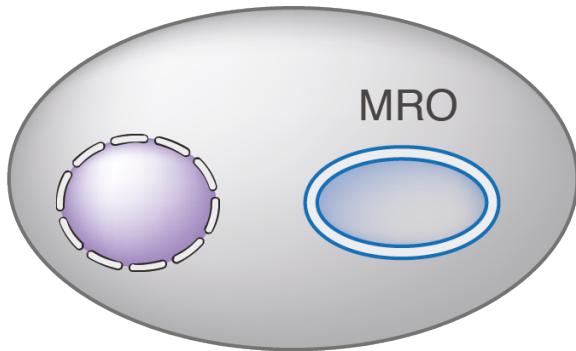
A eukaryotic cell with typical mitochondria



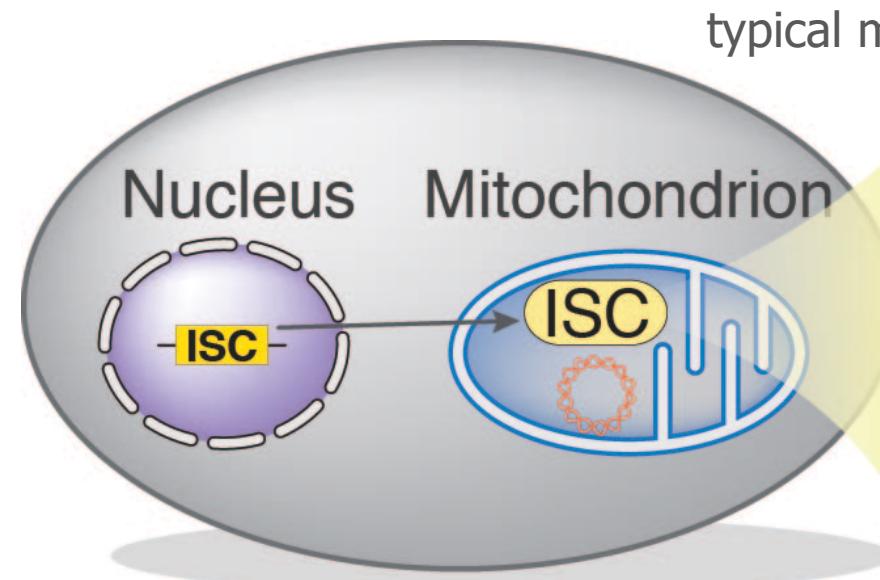
anaerobic eukaryotes



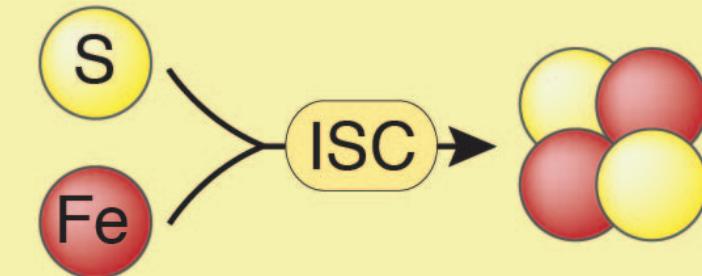
An anaerobic eukaryotes with mitochondrion related organelles (MROs)



O_2



typical mitochondria

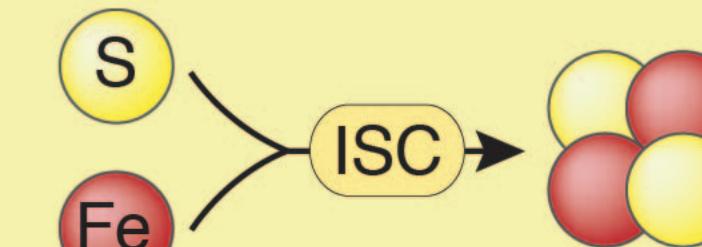
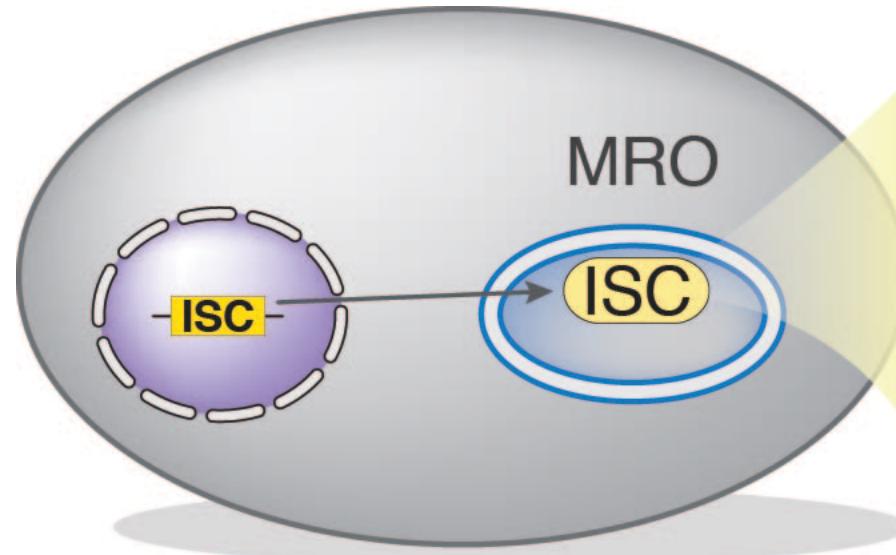


Mitochondrial Fe-S cluster assembly (ISC) system

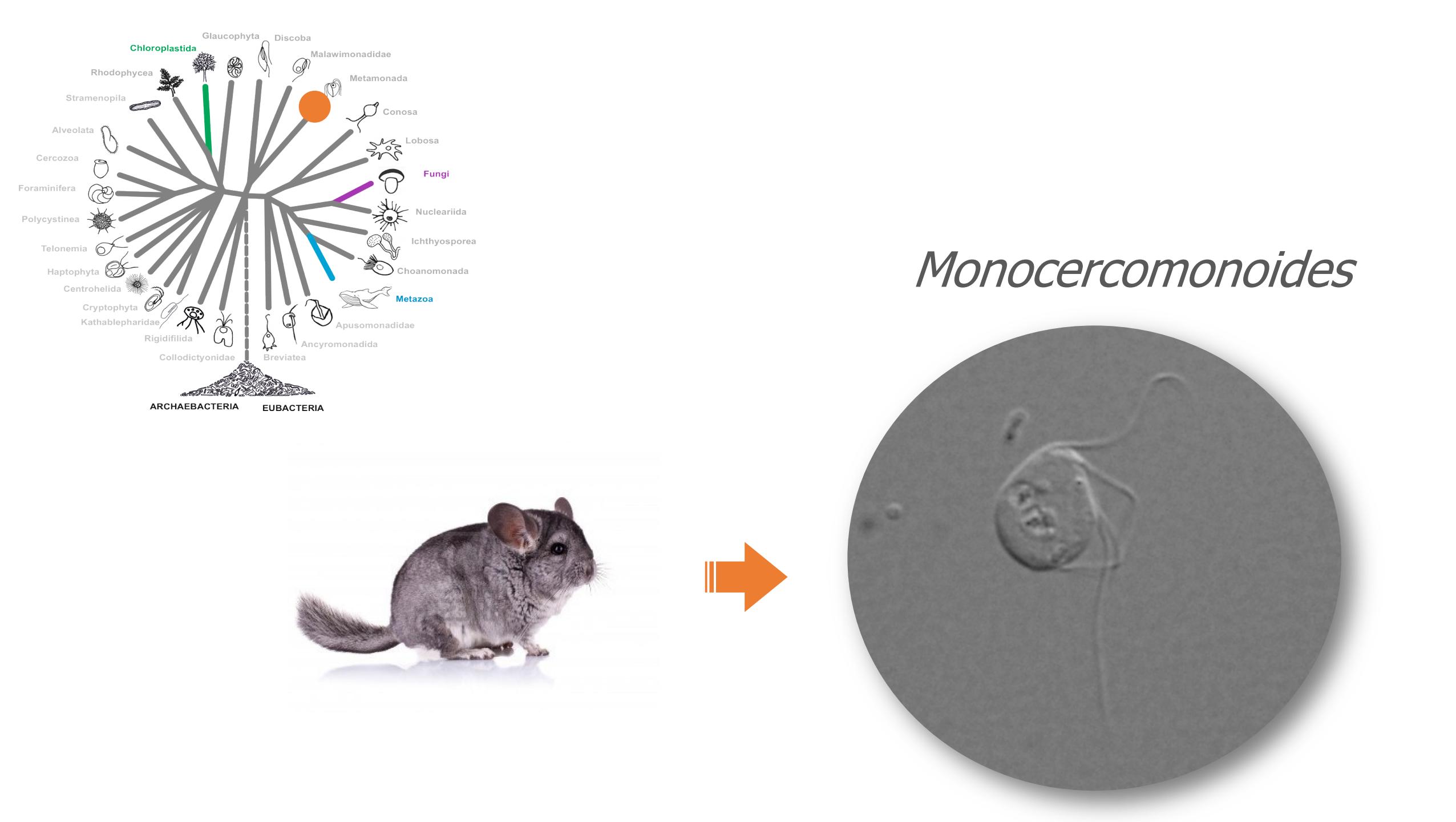


mitochondrion related organelle

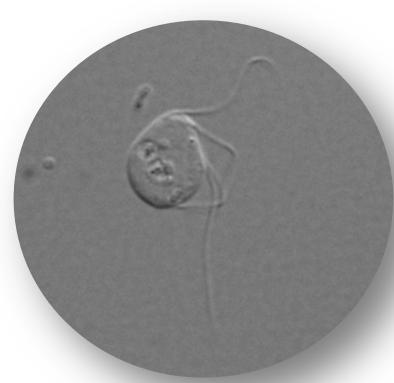
$\otimes O_2$



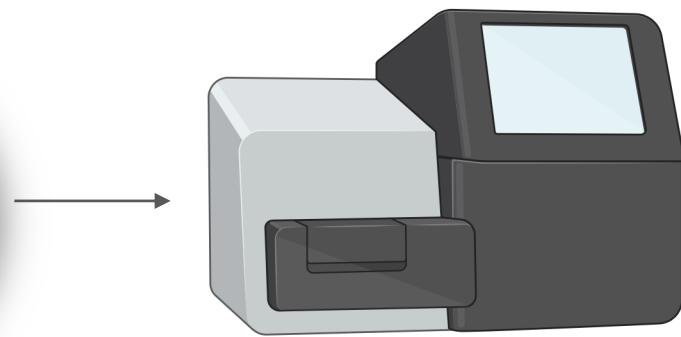
Mitochondrial Fe-S cluster assembly (ISC) system



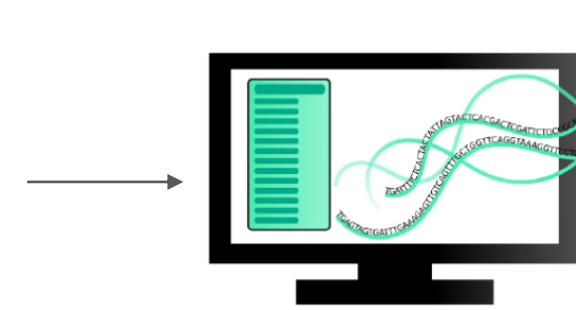
high-throughput genome sequencing & bioinformatics



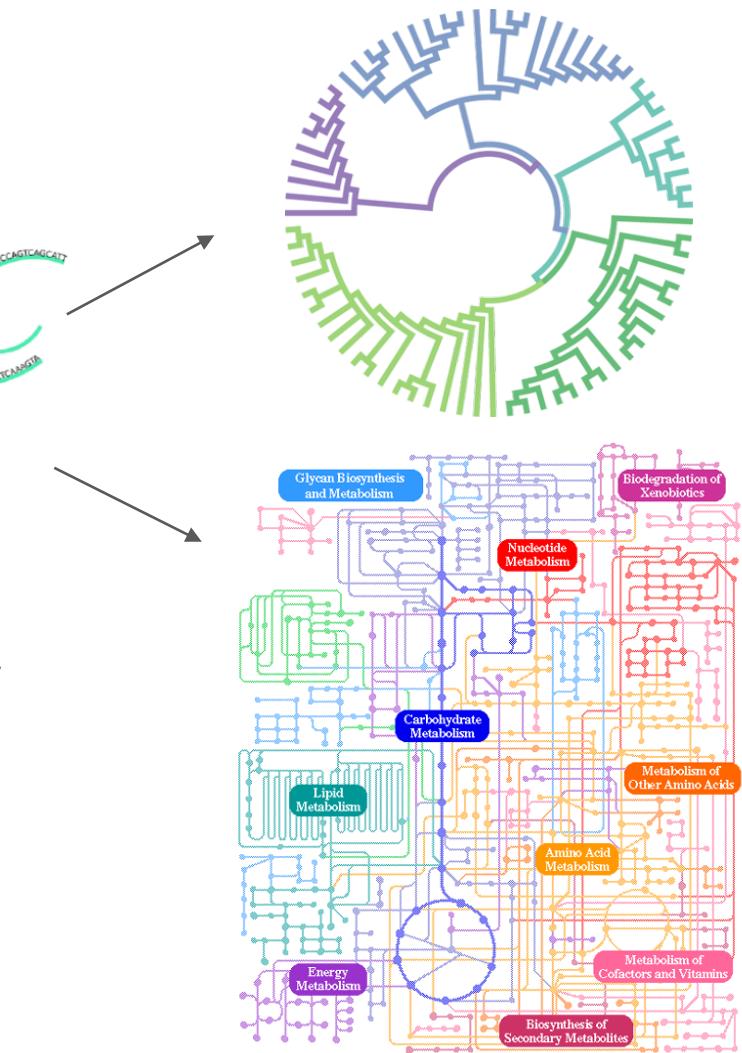
genomic DNA



sequencing

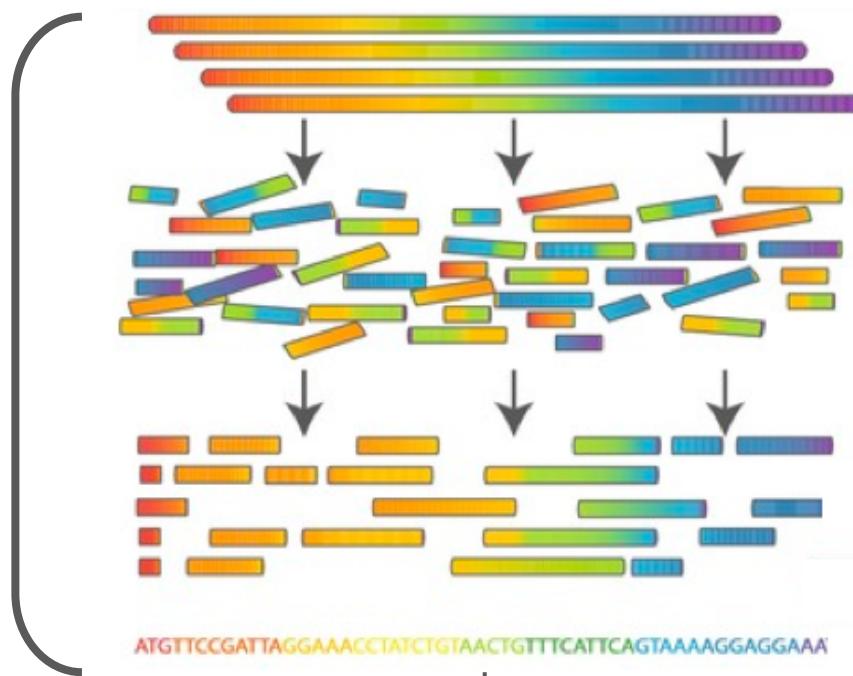


genome assembly
& annotation



reconstruction of the cell systems
& cell metabolism

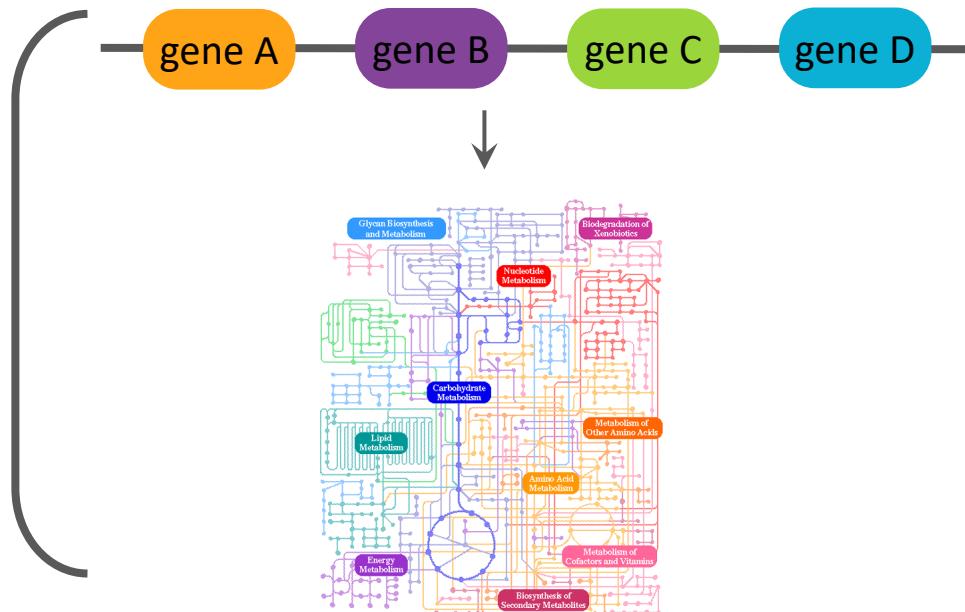
genome assembly



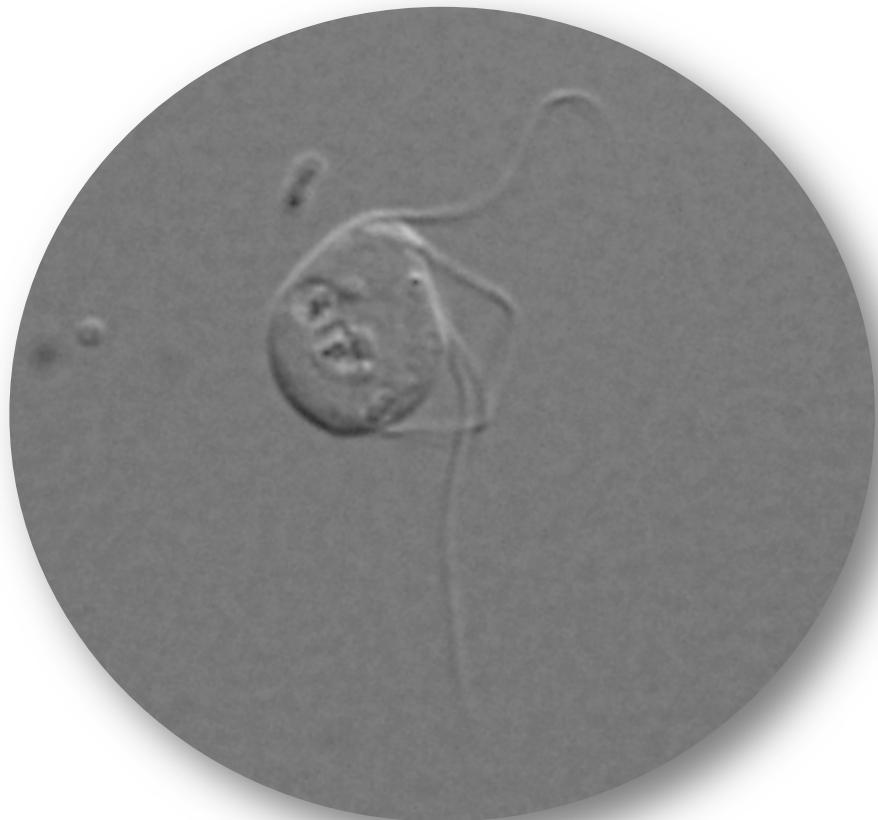
2095 scaffolds

75 000 000 base pairs

genome annotation

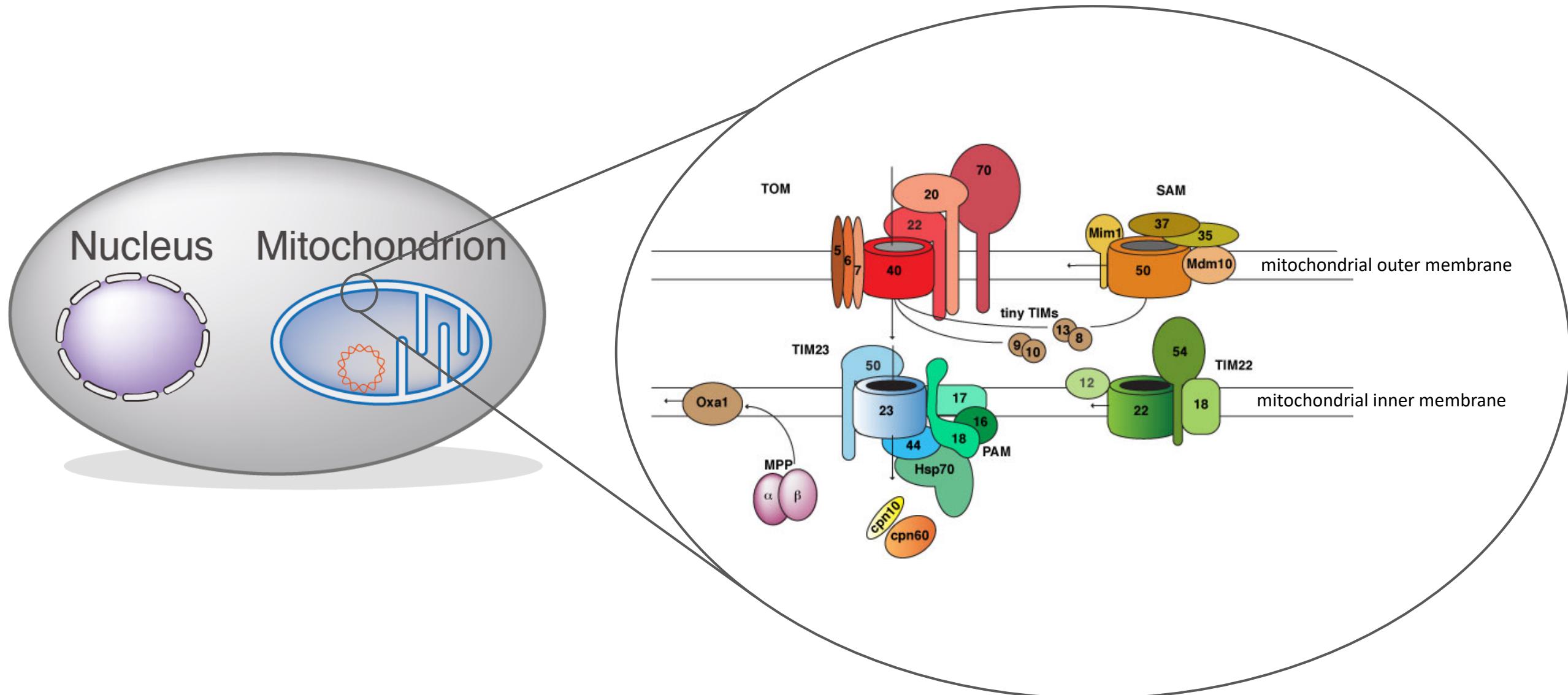


16 629 genes

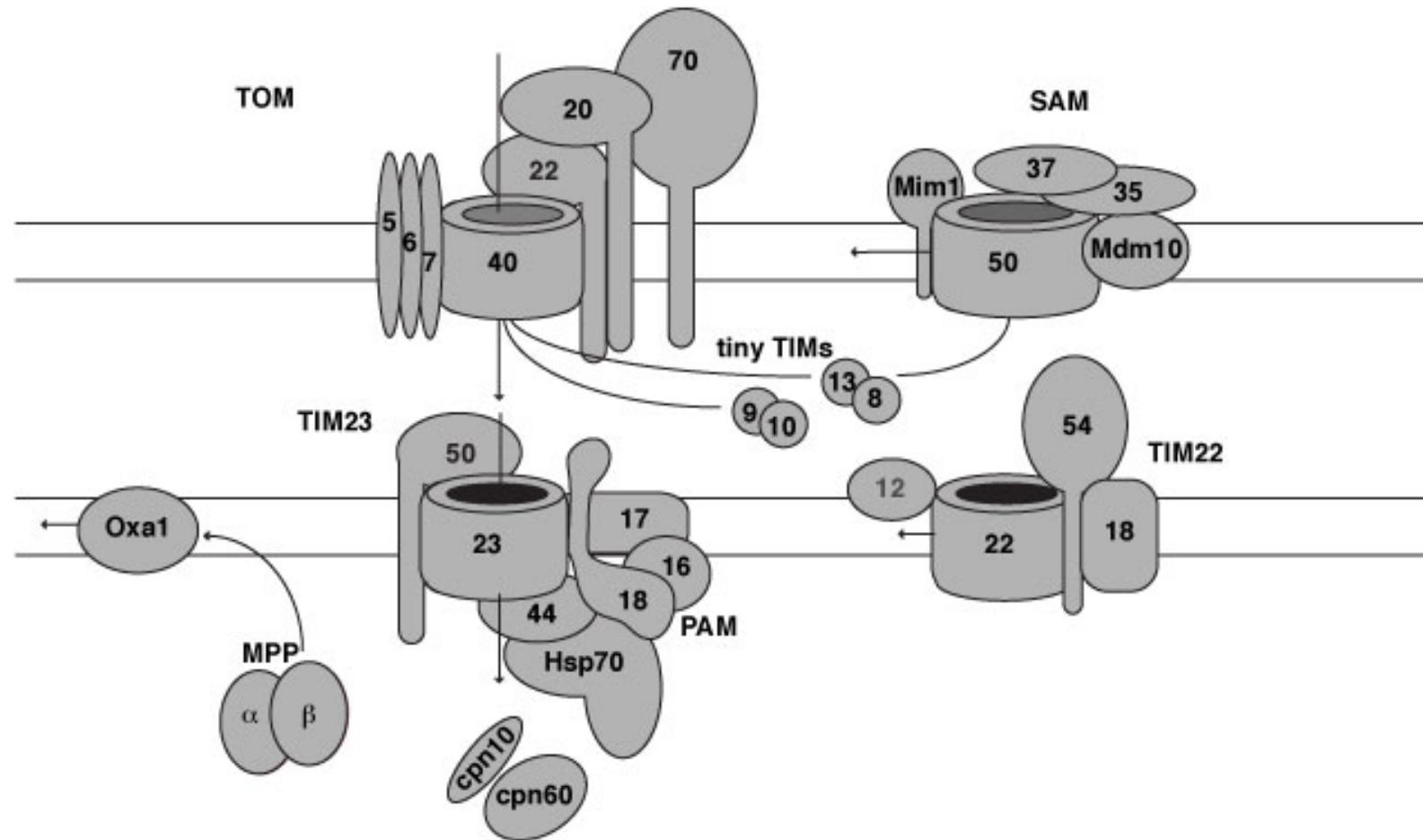
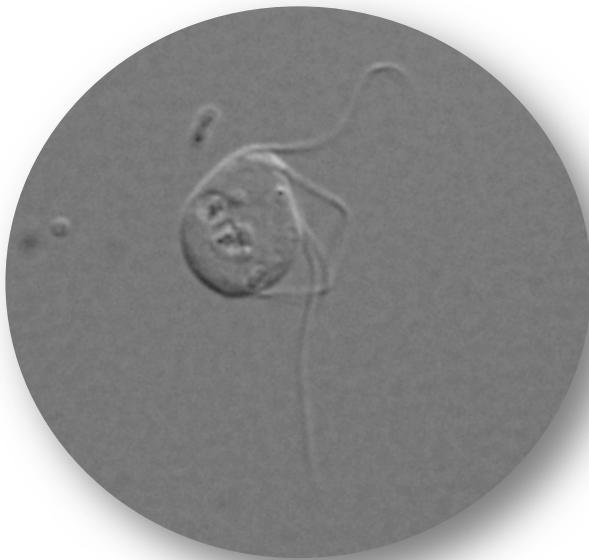


Monocercomonoides
searching for mitochondria

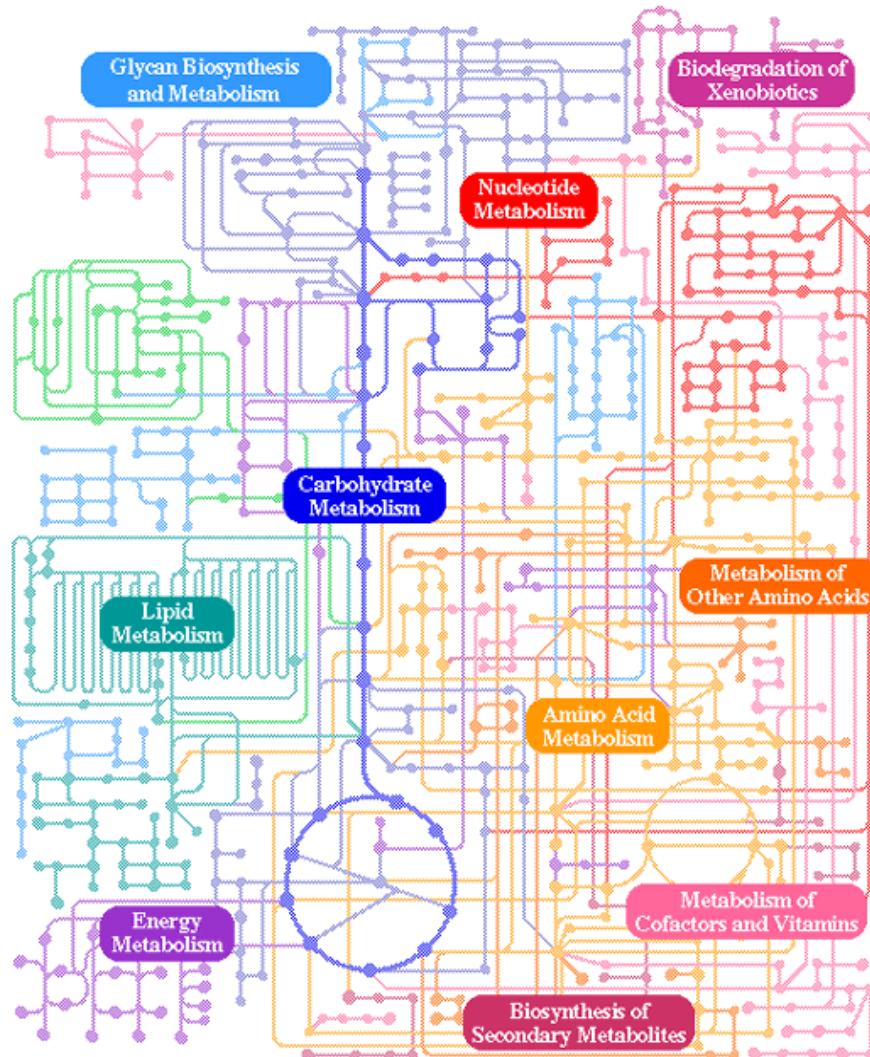
Searching for hallmark mitochondrial proteins



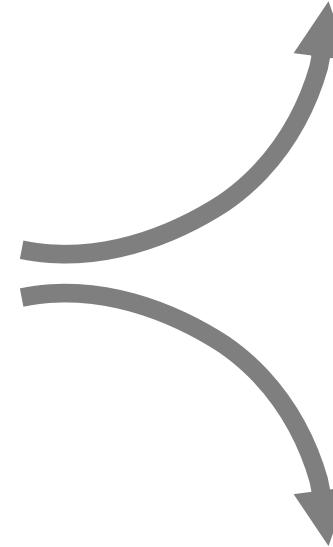
Searching for hallmark mitochondrial proteins



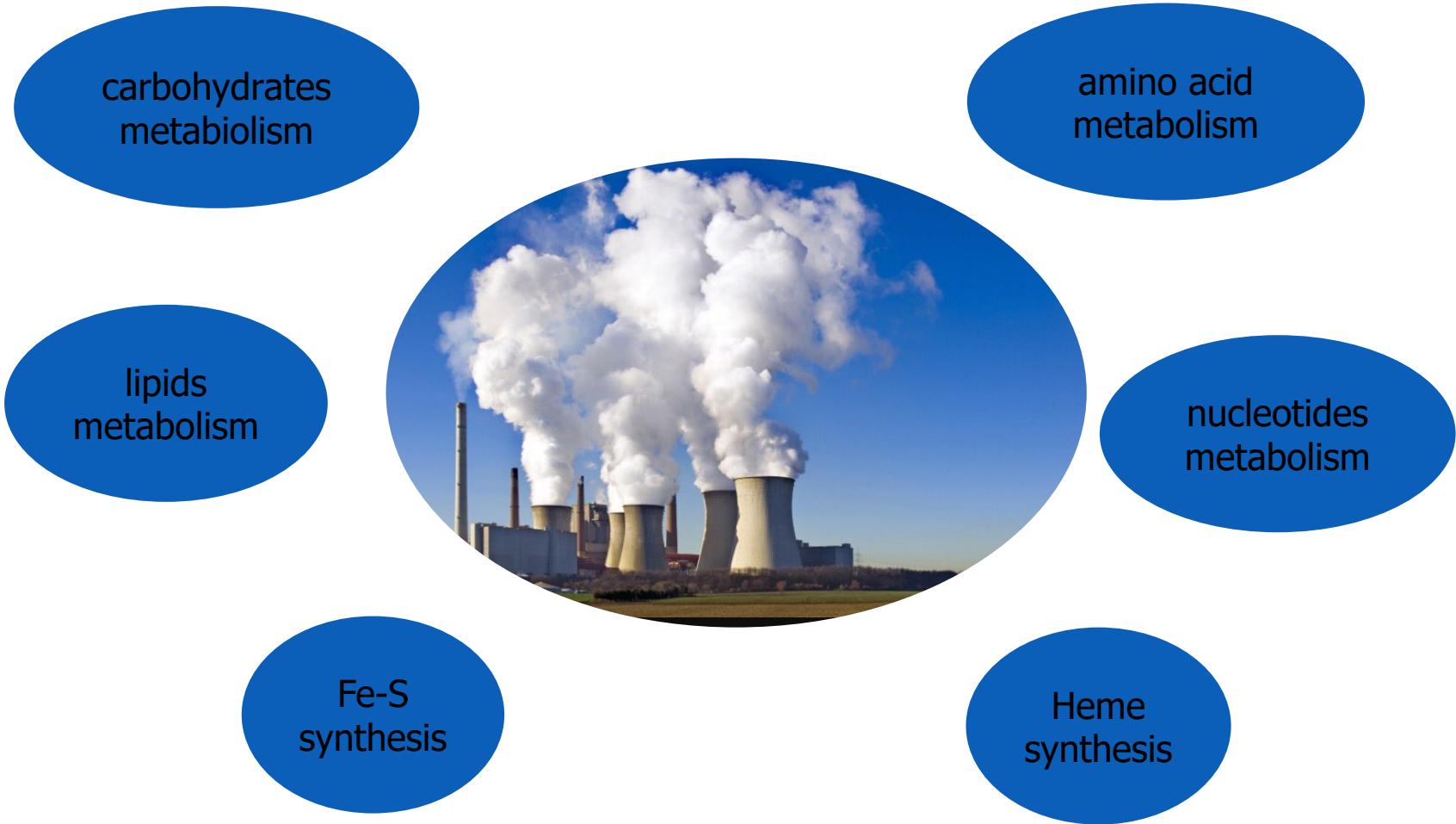
Searching for proteins involved mitochondrial metabolism



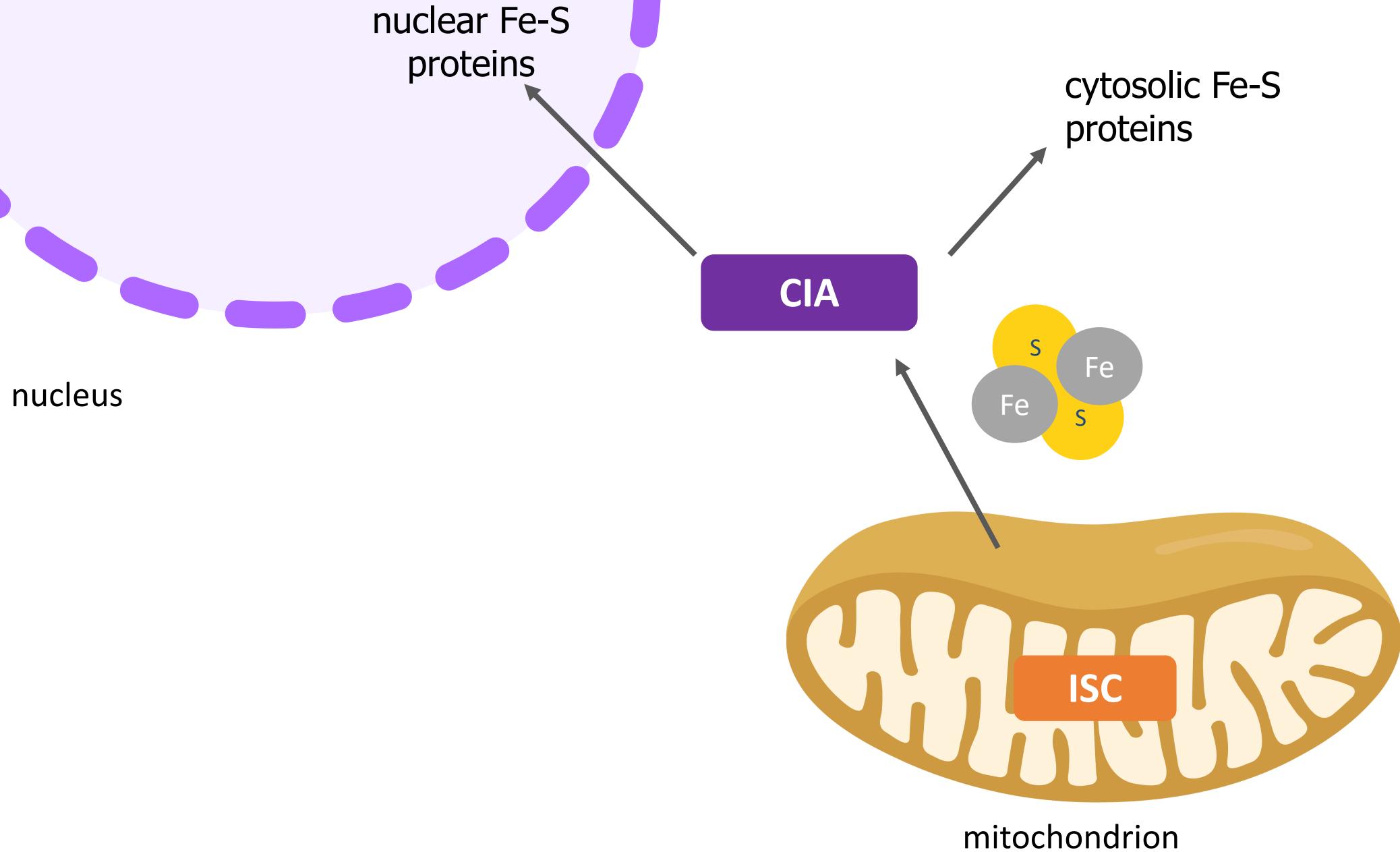
metabolic pathways in cytosol



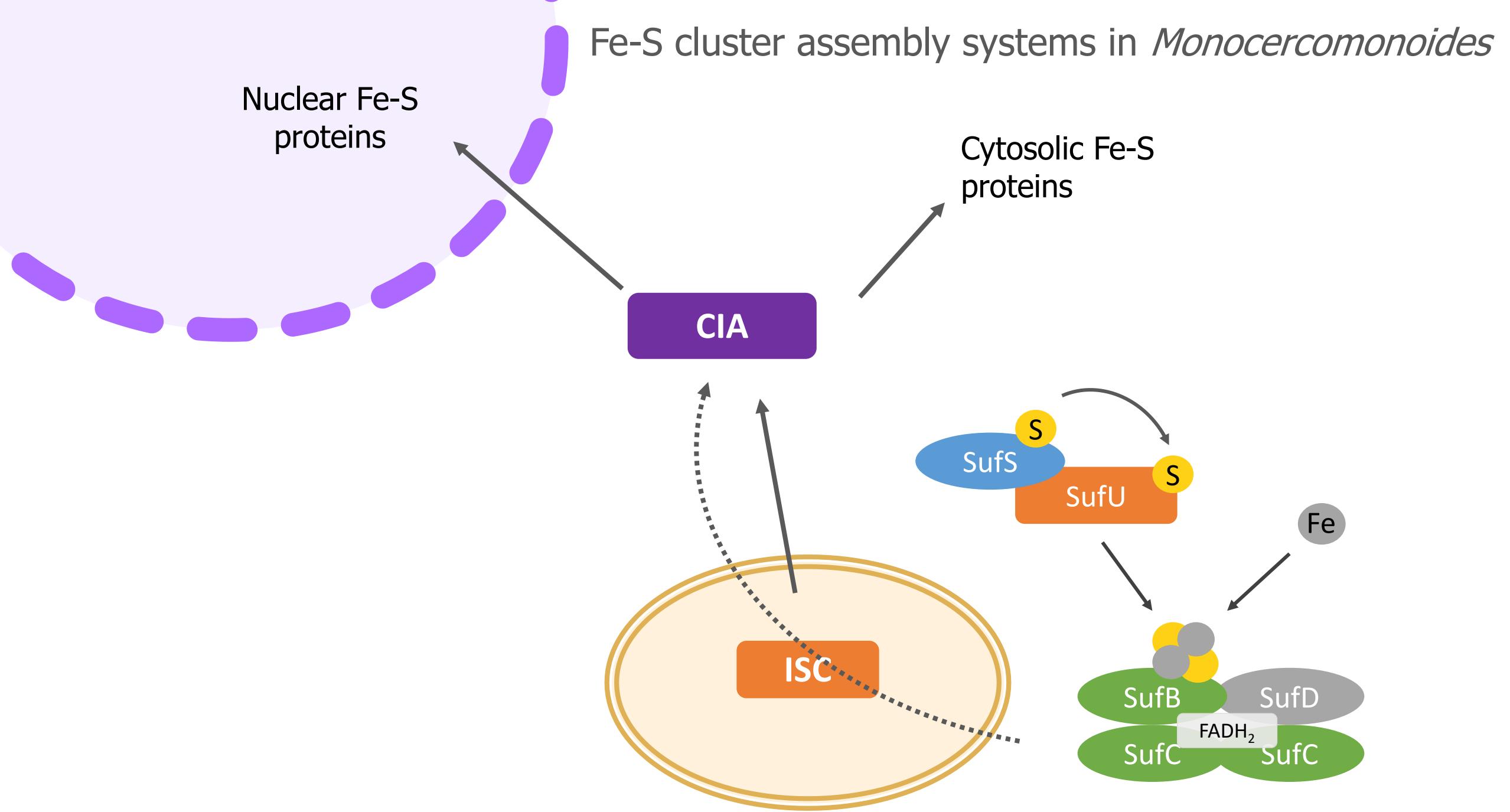
transport from the host



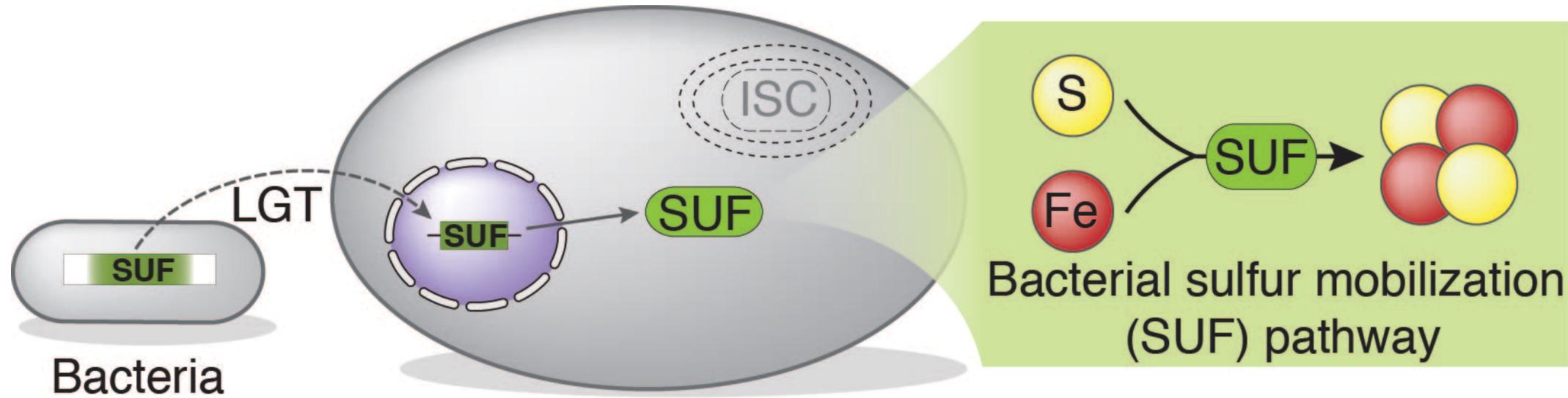
Fe-S cluster assembly systems in eukaryotes



Fe-S cluster assembly systems in *Monocercomonoides*



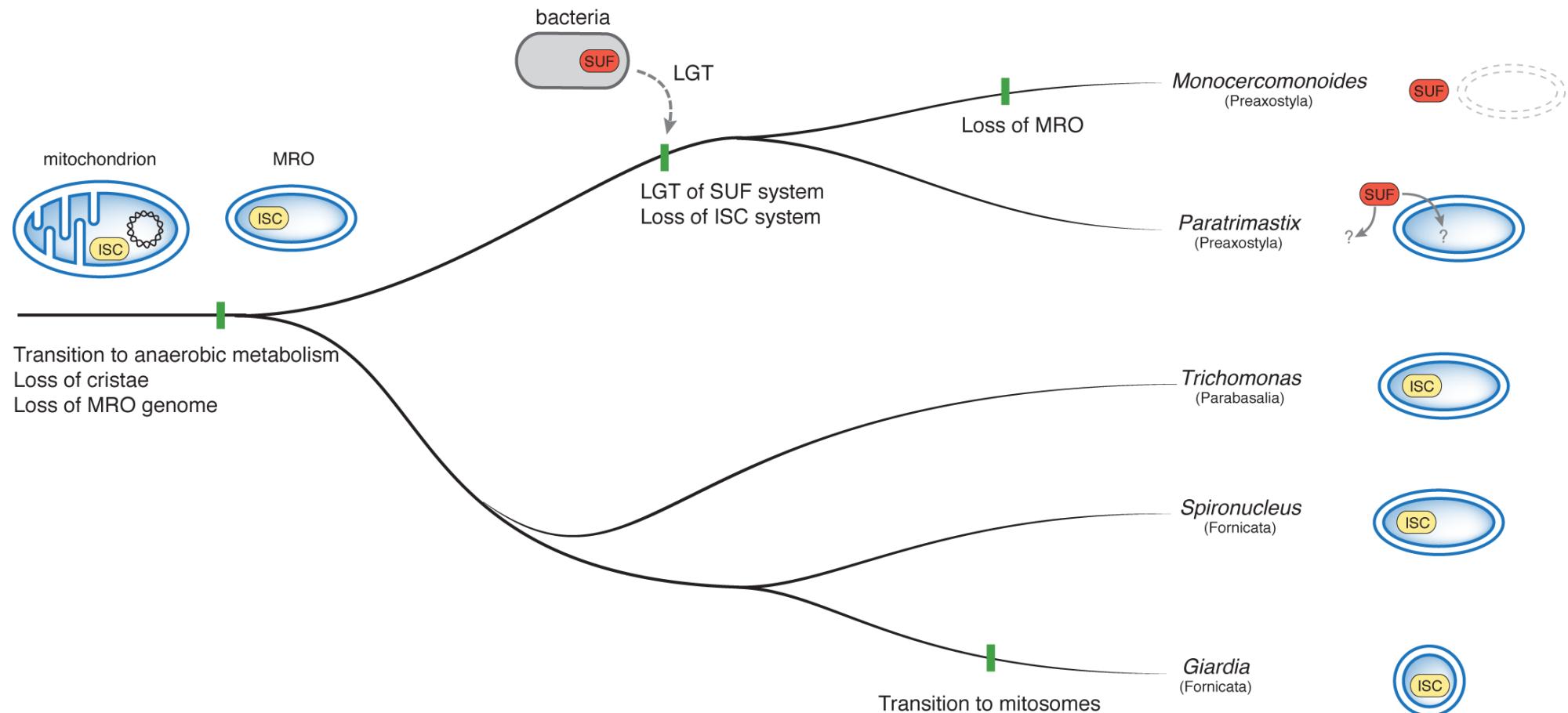
Monocercomonoides an amitochondriate eukaryote



Lateral gene transfer (LGT) of bacterial genes encoding sulfur mobilization pathway (SUF) and loss of reduced mitochondrion

Loss of a mitochondrial organelle

LGT of SUF system resulted in relocation of the pathway to the cytosol

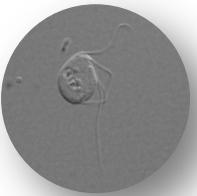


Endosymbiosis can be undone!

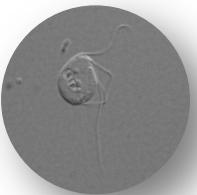
Take home message



(almost all) eukaryotic cells have mitochondria



complex eukaryotic cell can live without a mitochondrion



Monocercomonoides can serve as a model for cell biology studies



studies on eukaryotic microbial diversity help to understand evolution of eukaryotes

Acknowledgements



Hampl Lab



Keeling Lab





University of Warsaw
Biological and Chemical
Research Centre



Exploring microbial diversity to understand the evolution of eukaryotes

