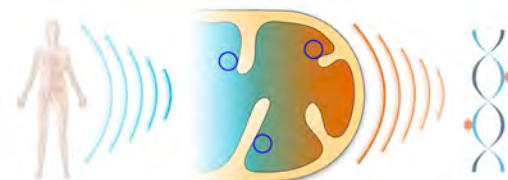


Profiling multifunctional mitochondria in the brain and beyond



Martin Picard, Ph.D.
Department of Psychiatry, Division of Behavioral Medicine
Department of Neurology, H. Houston Merritt Center
Robert N Butler Columbia Aging Center
Columbia Translational Neuroscience Initiative
New York State Psychiatric Institute (NYSPI)

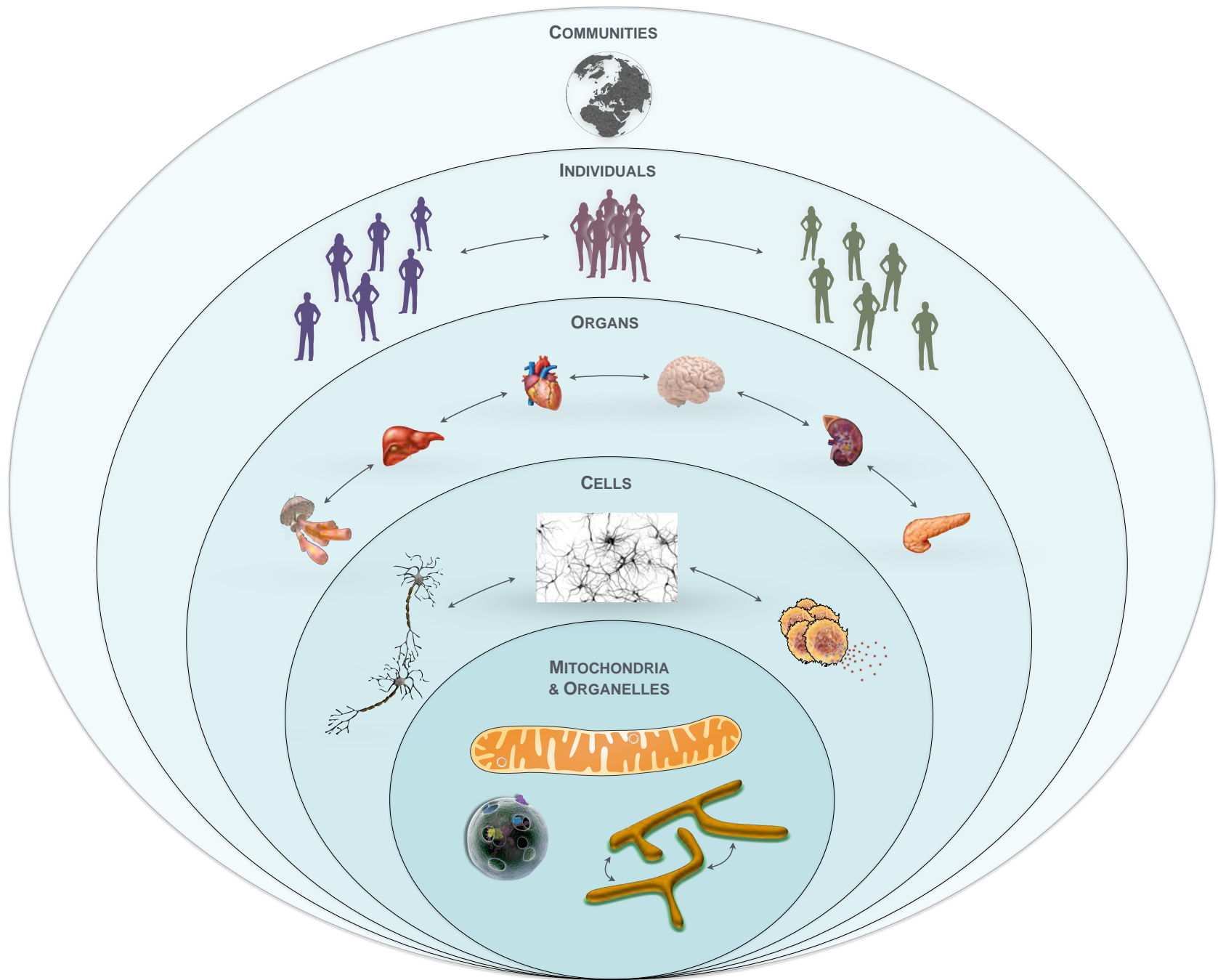
 **COLUMBIA**
COLUMBIA UNIVERSITY
IRVING MEDICAL CENTER

 **New York State
Psychiatric Institute**

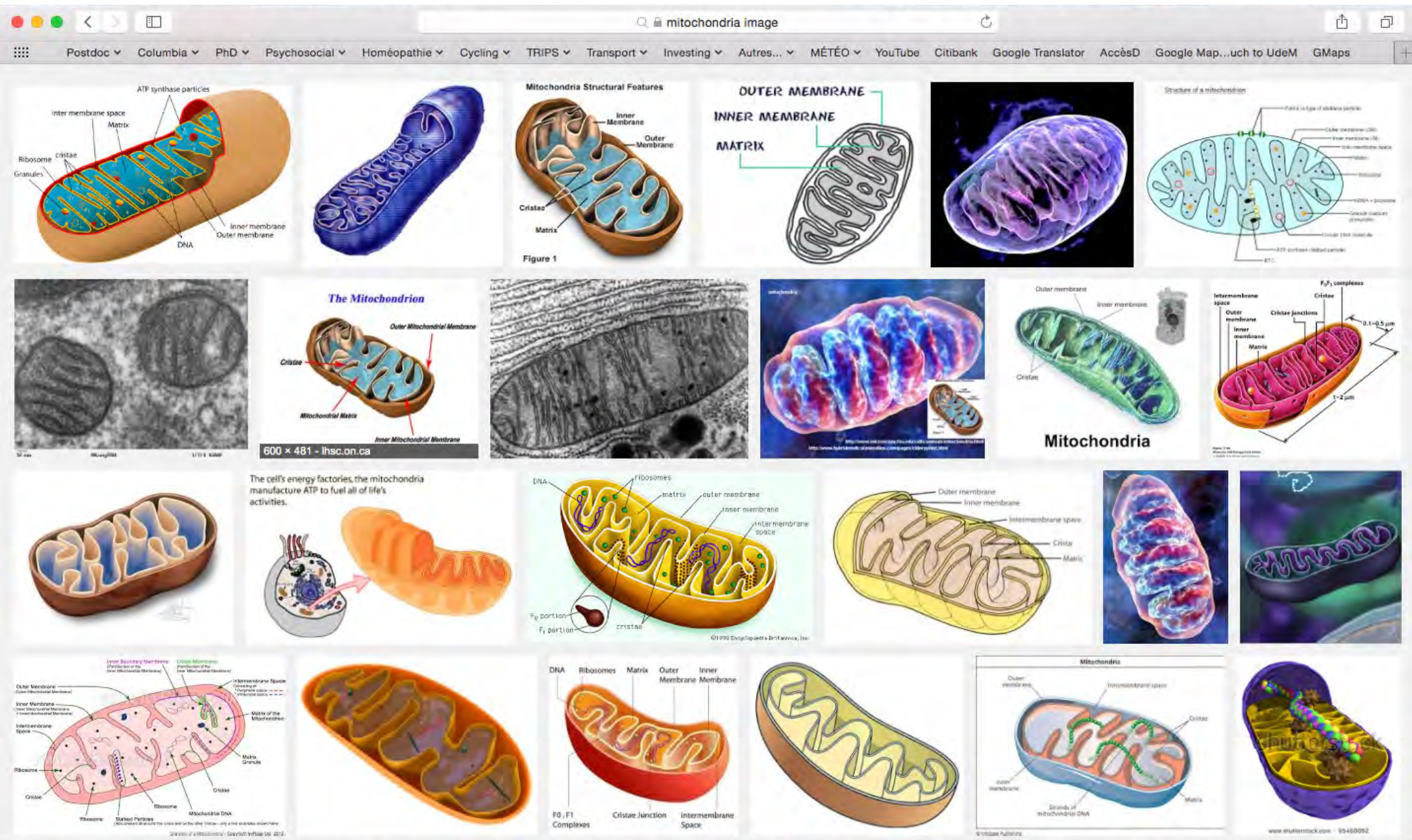
1.Role of mitochondria in organisms

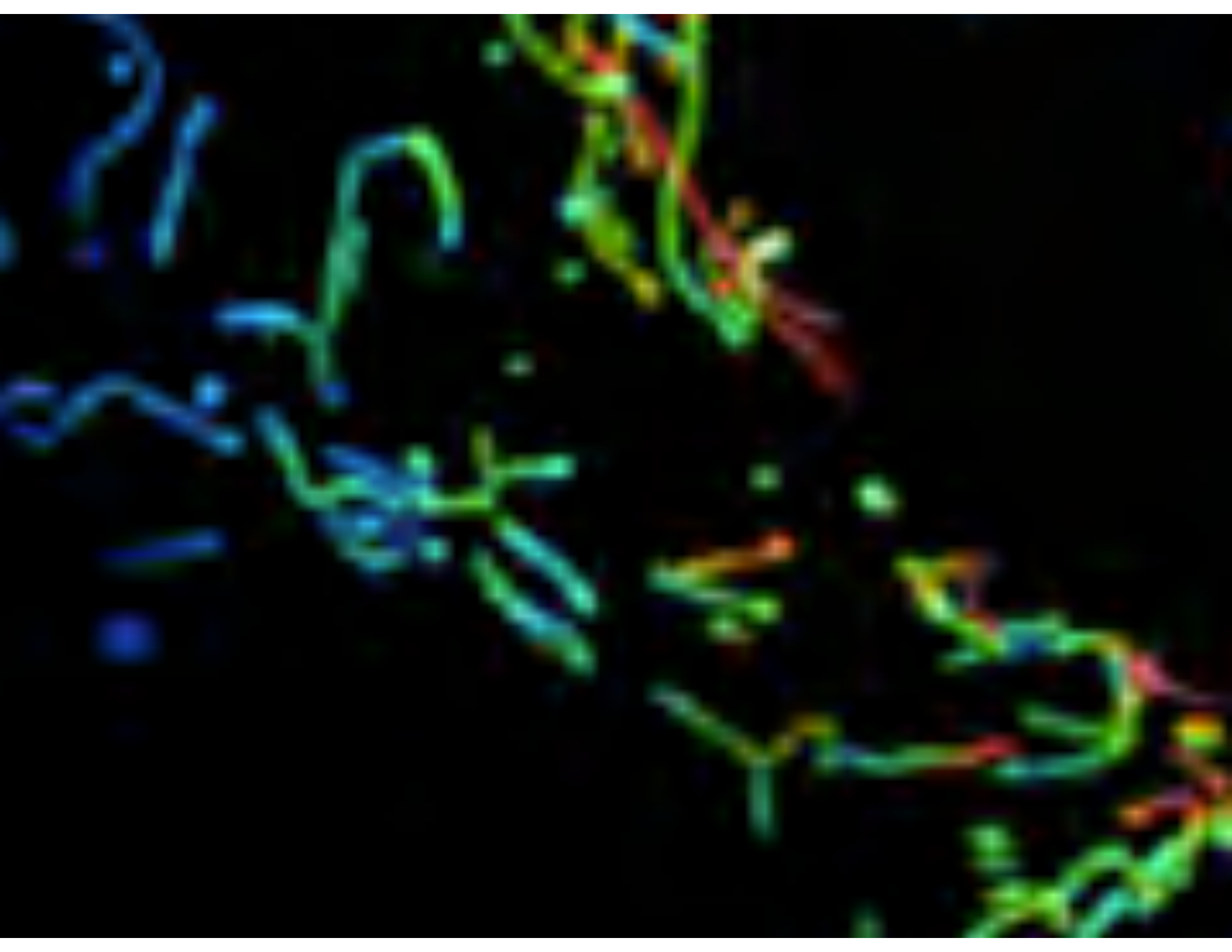
2.Celebrating diversity of mitochondria

3.Human brain mitochondria



What do mitochondria look like?





What do mitochondria do?

SCIENTIFIC
AMERICAN®

HEALTH

Powerhouse of the Cell

It is the mitochondrion, a small body which appears to play a central role in the oxidation of foodstuff. Its structure, as revealed by the electron microscope, mirrors its function

By Philip Siekevitz on July 1, 1957



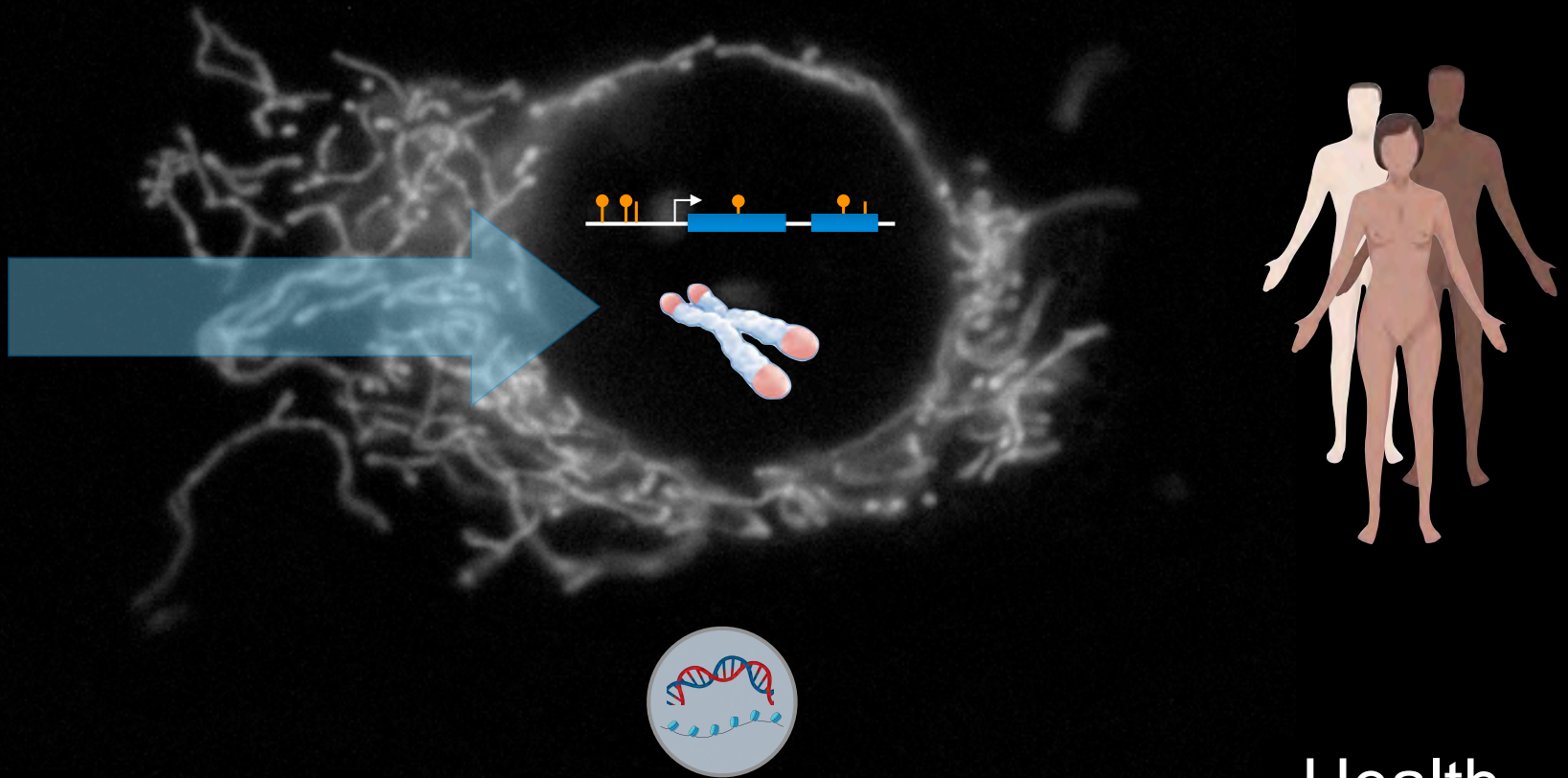
Environment

X

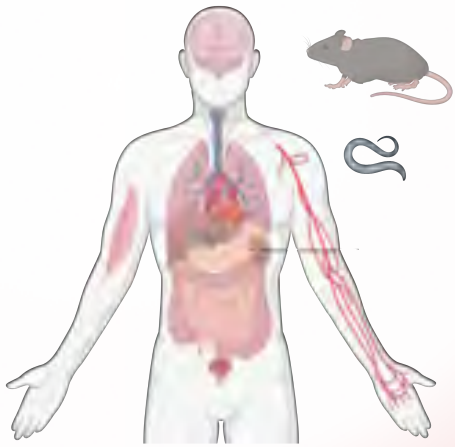
Gene

Health
Lifespan

“Are mitochondria the X factor?”



Multicellular organisms



OUTPUTS (mitochondrial, DIRECT)

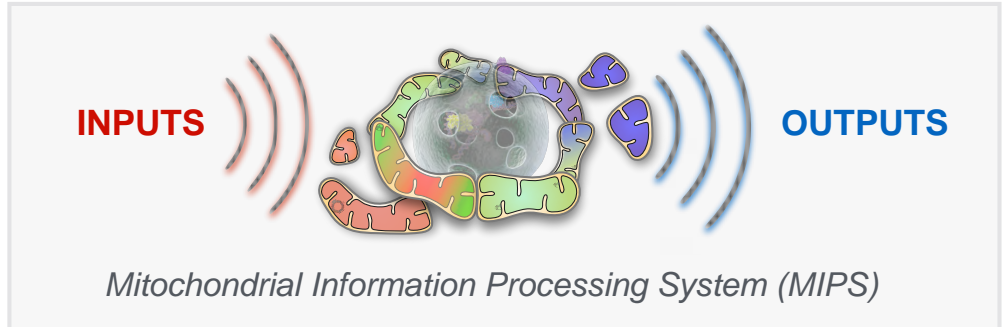
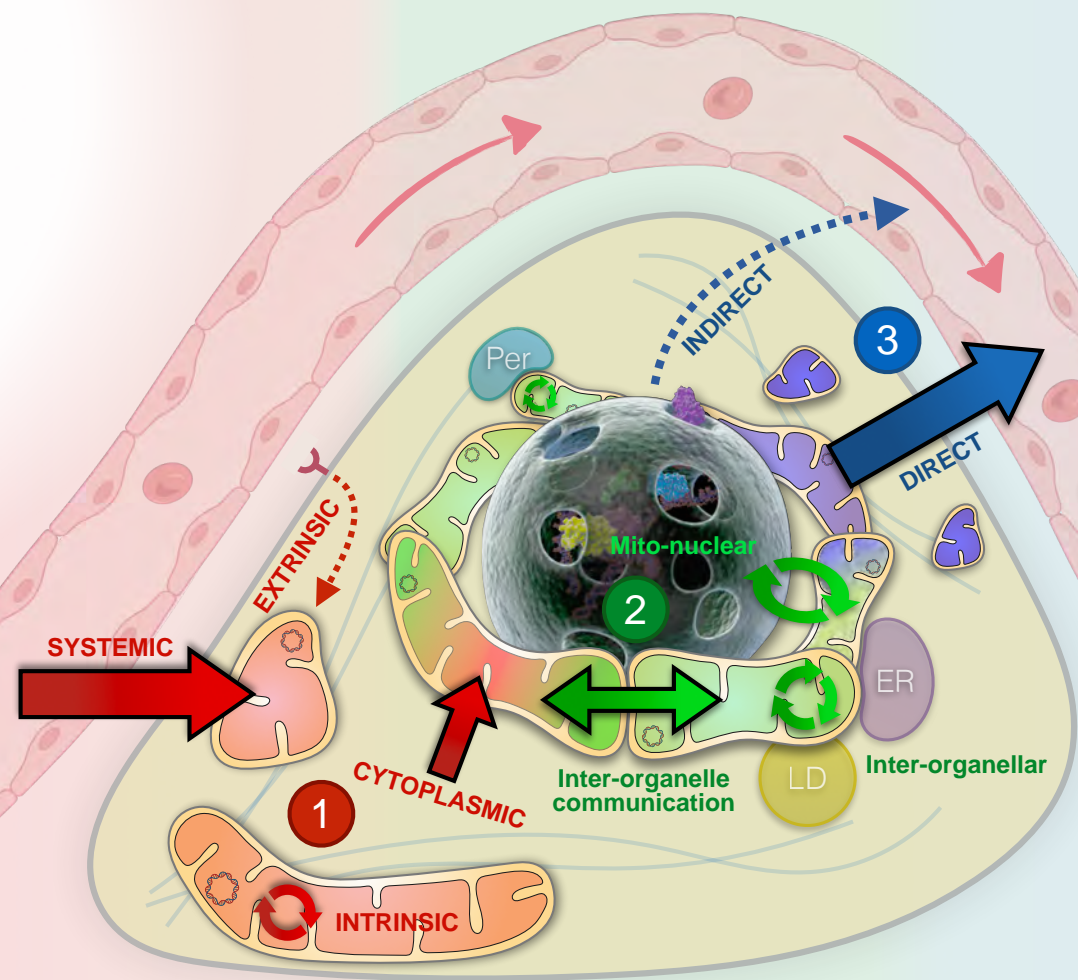
- Metabolites
- Lipids
- DNA and RNA
- cf-mtDNA (whole, fragments)
- ATP (ΔG_p)
- Ions
- ROS
- Gases
- Heat
- Steroid hormones
- Small peptides
- Others

OUTPUTS (via nucleus, INDIRECT)

- Peptide hormones

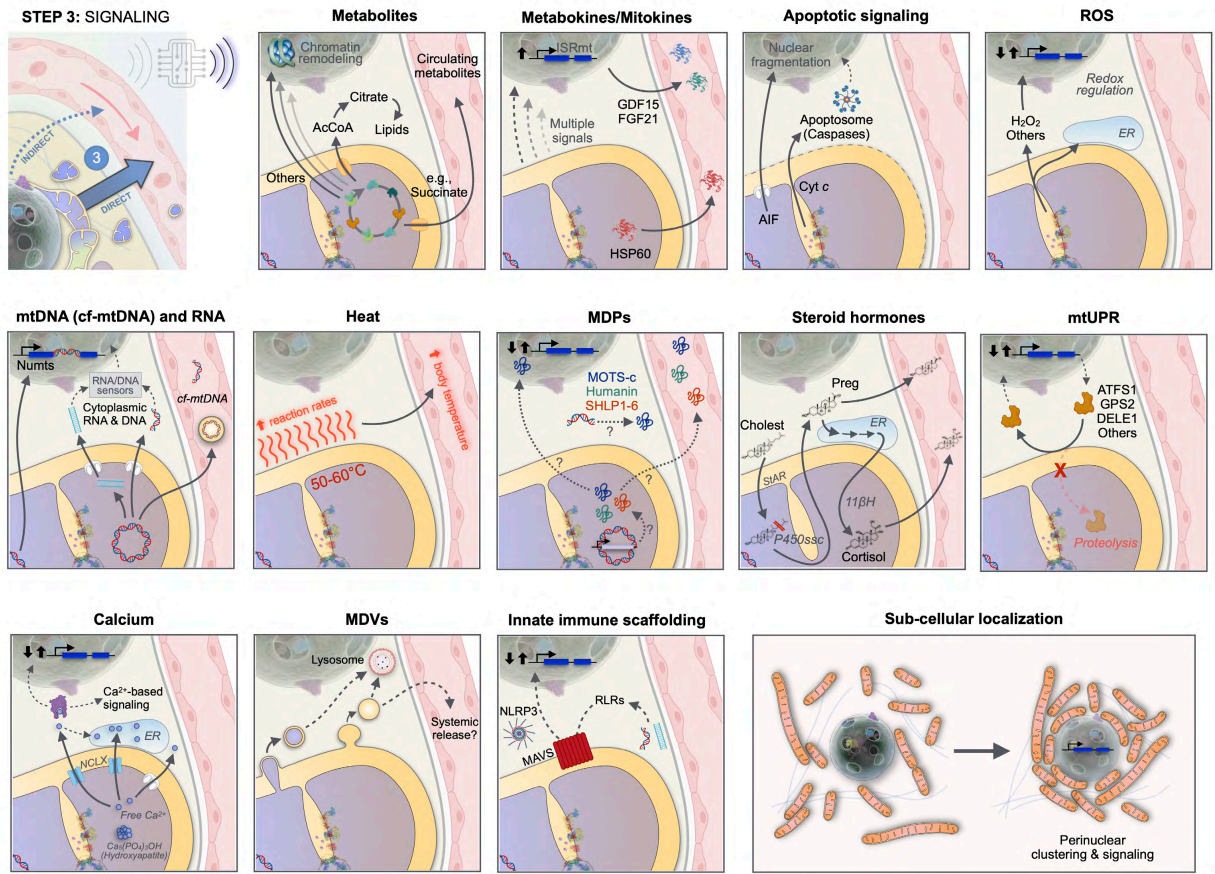
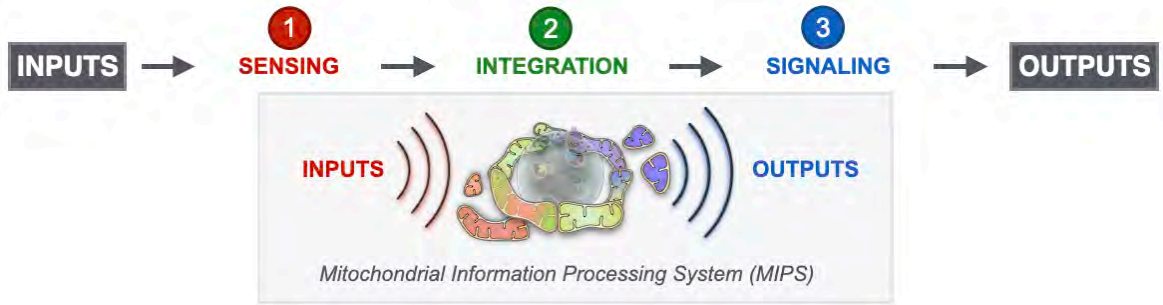
INPUTS

- Peptide hormones
- Steroid & other hormones
- Nutrients levels
- Metabolites
- Ions
- Gases (e.g., O_2 , NO)
- ATP/ADP (ΔG_p)
- NAD(P)⁺/NAD(P)H ratio
- mtDNA variations
- Others

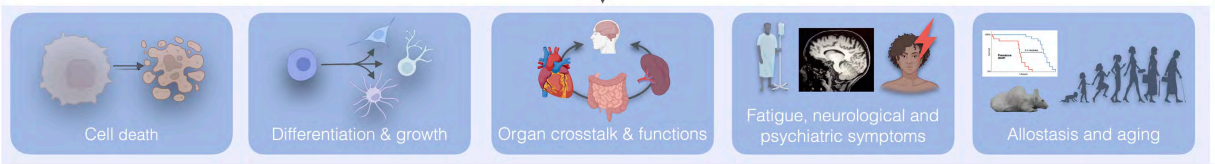


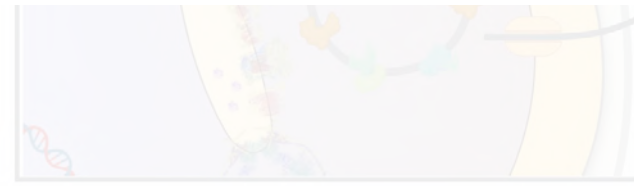
The hallmarks of mitochondrial signal transduction



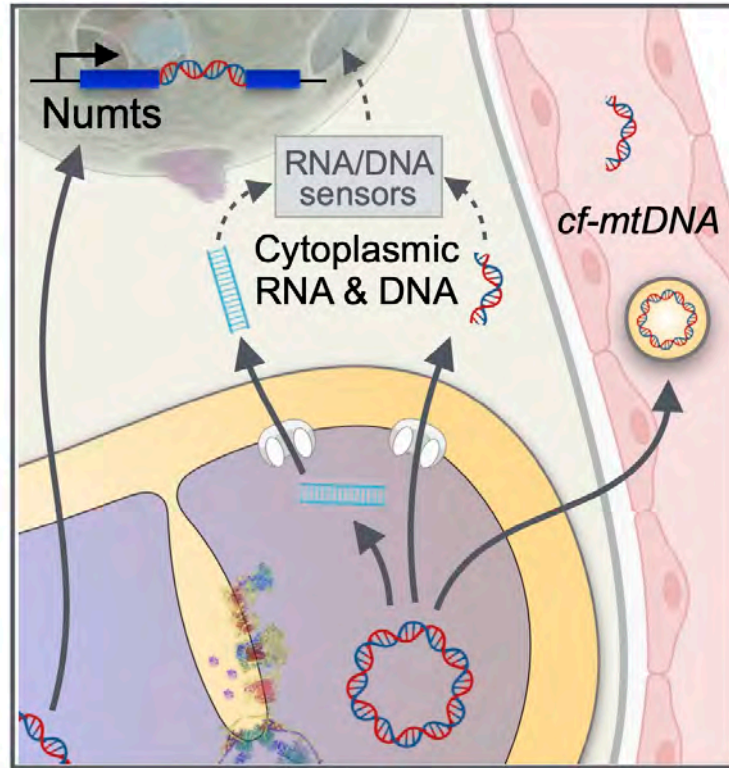


MIPS-derived intracellular and systemic signals





mtDNA (cf-mtDNA) and RNA



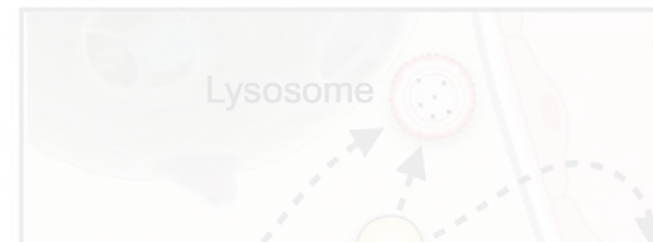
Heat



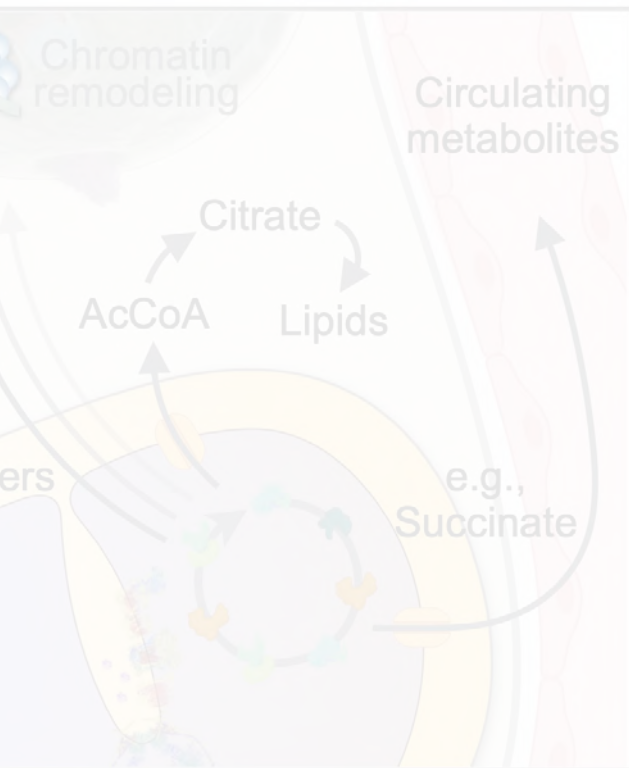
Calcium



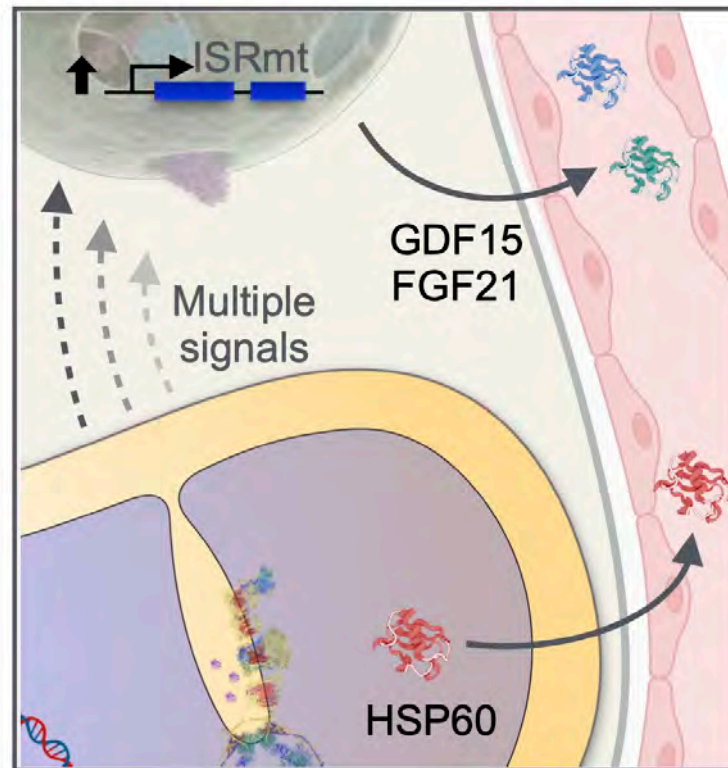
MDVs



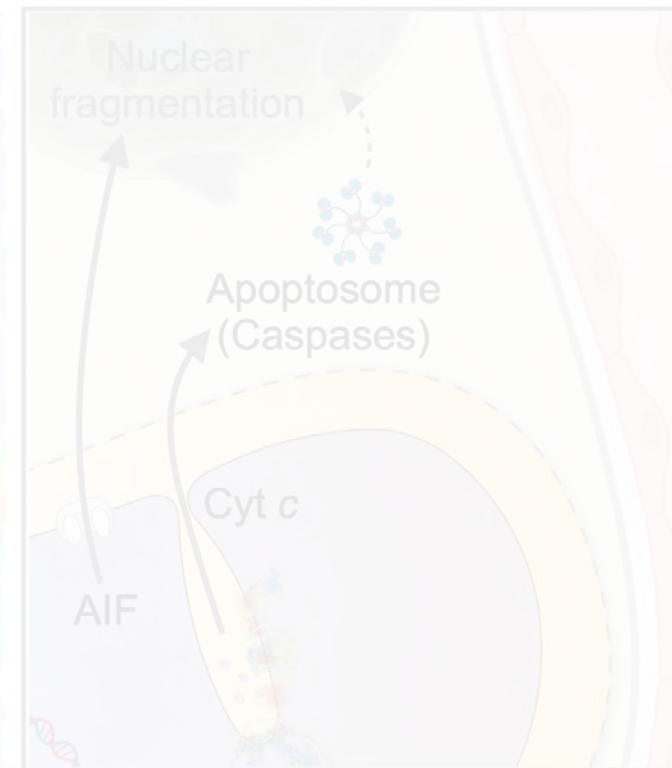
Metabolites



Metabokines/Mitokines



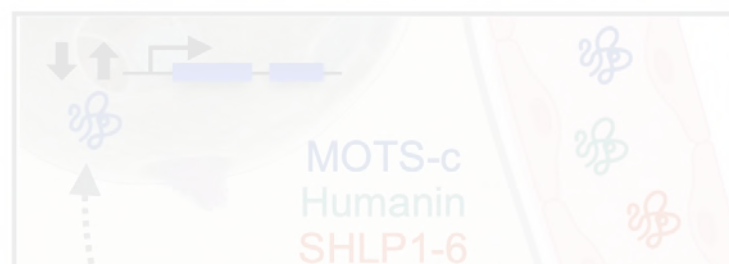
Apoptotic signaling



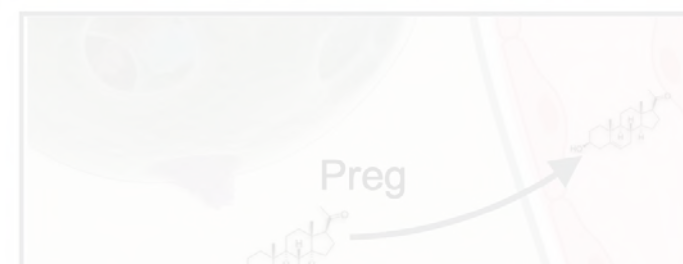
Heat

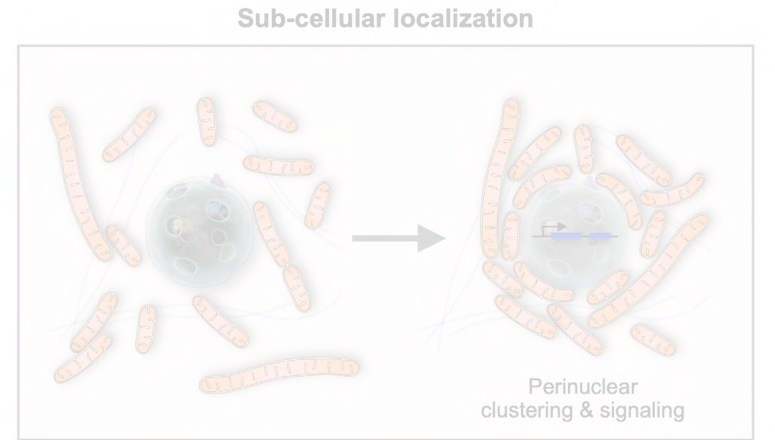
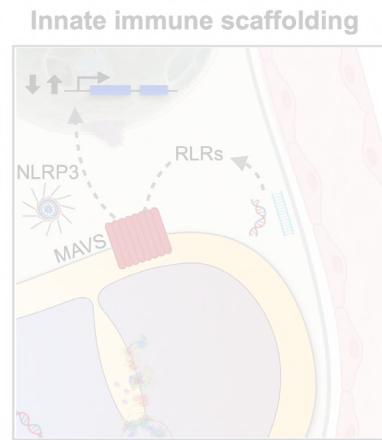
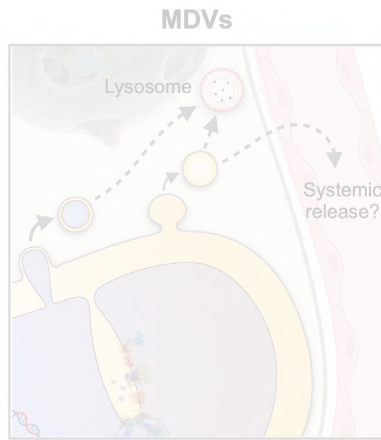
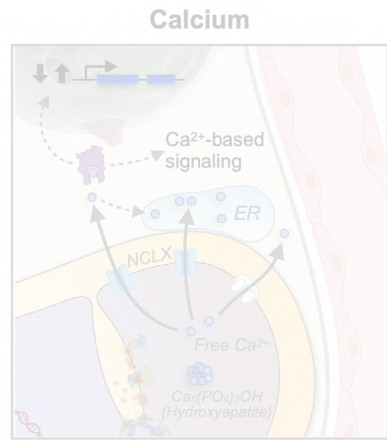


MDPs

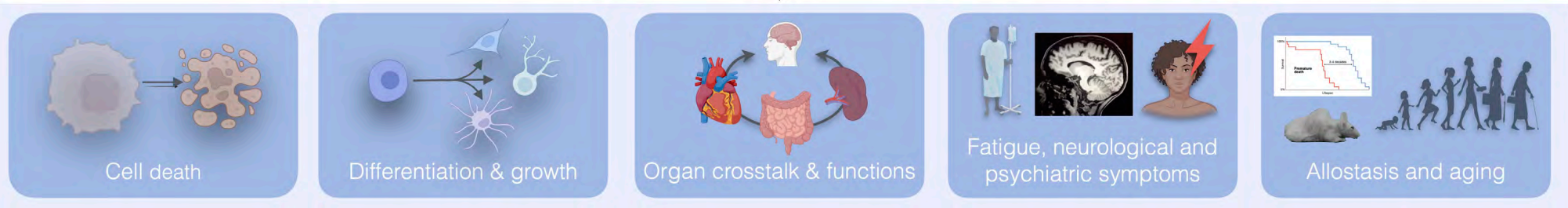


Steroid hormones



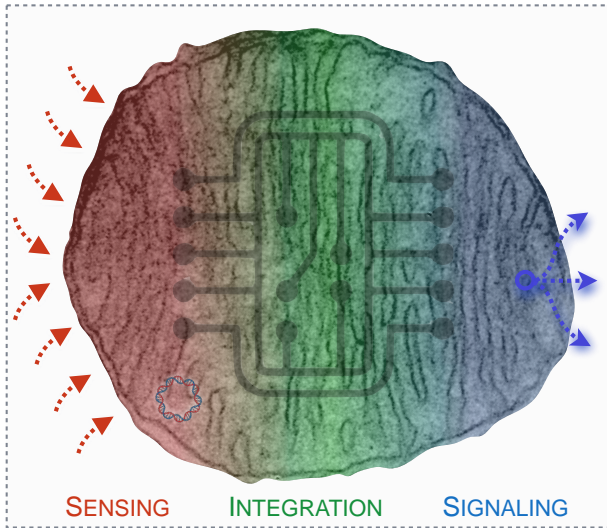


MIPs-derived intracellular and systemic signals

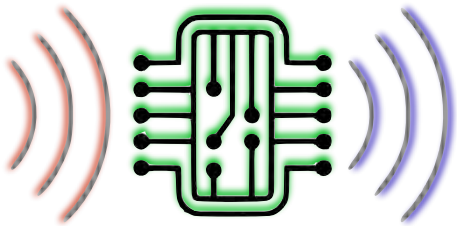


Mitochondrial Information Processing System — MIPS

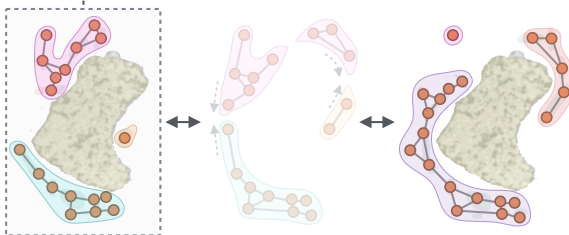
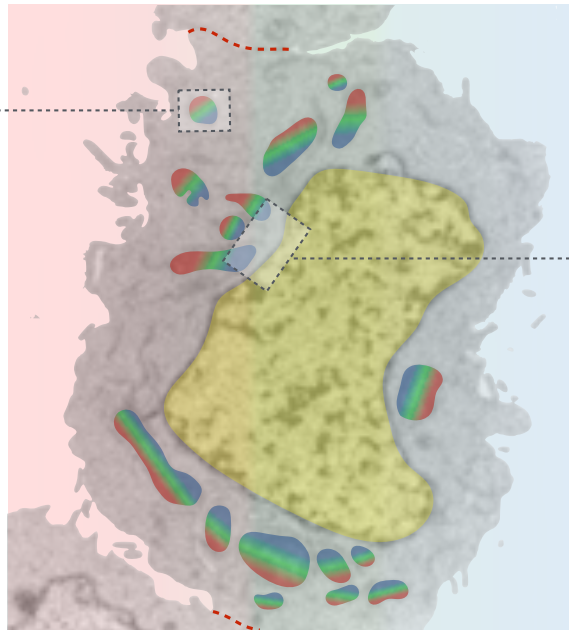
Signal transducing mitochondrion



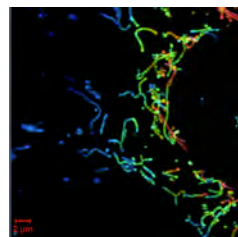
Incoming data Outgoing data



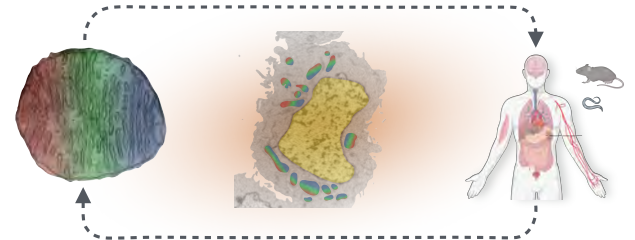
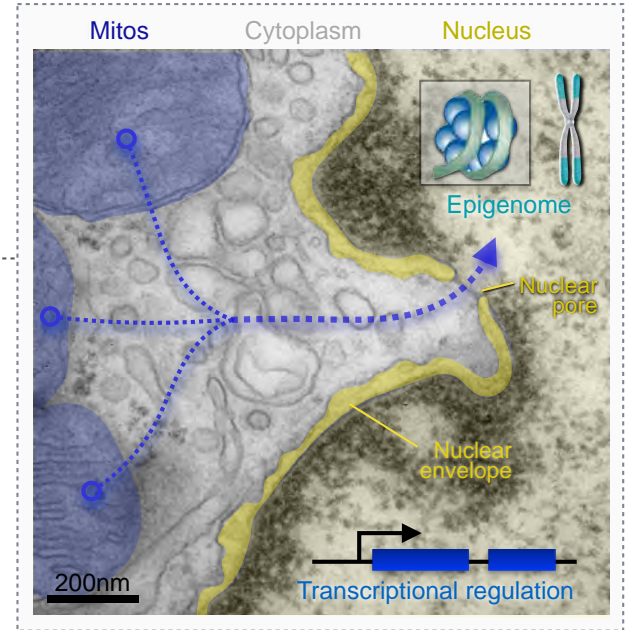
"Mitochondria are the processor of the cell"



Dynamic remodeling of mito networks

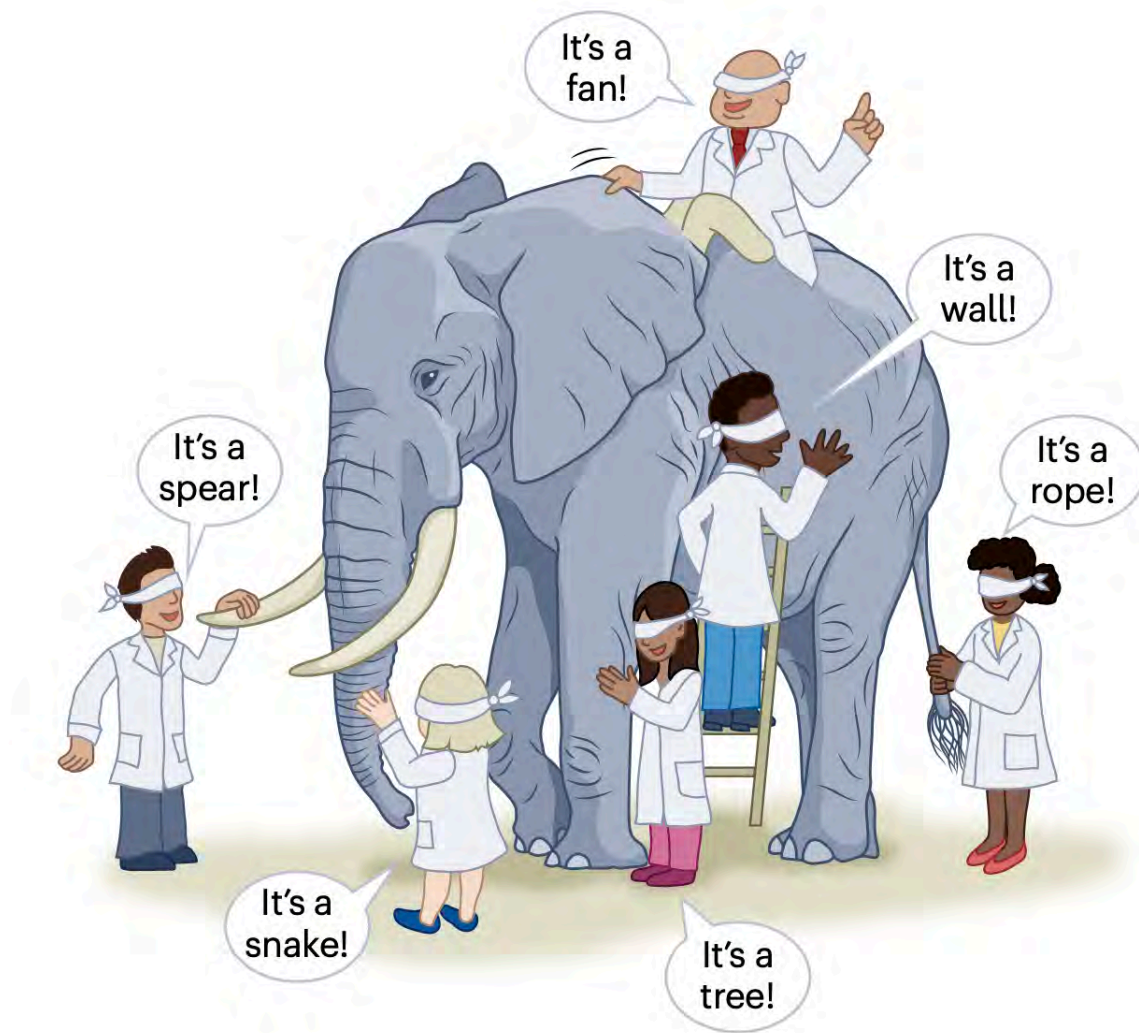


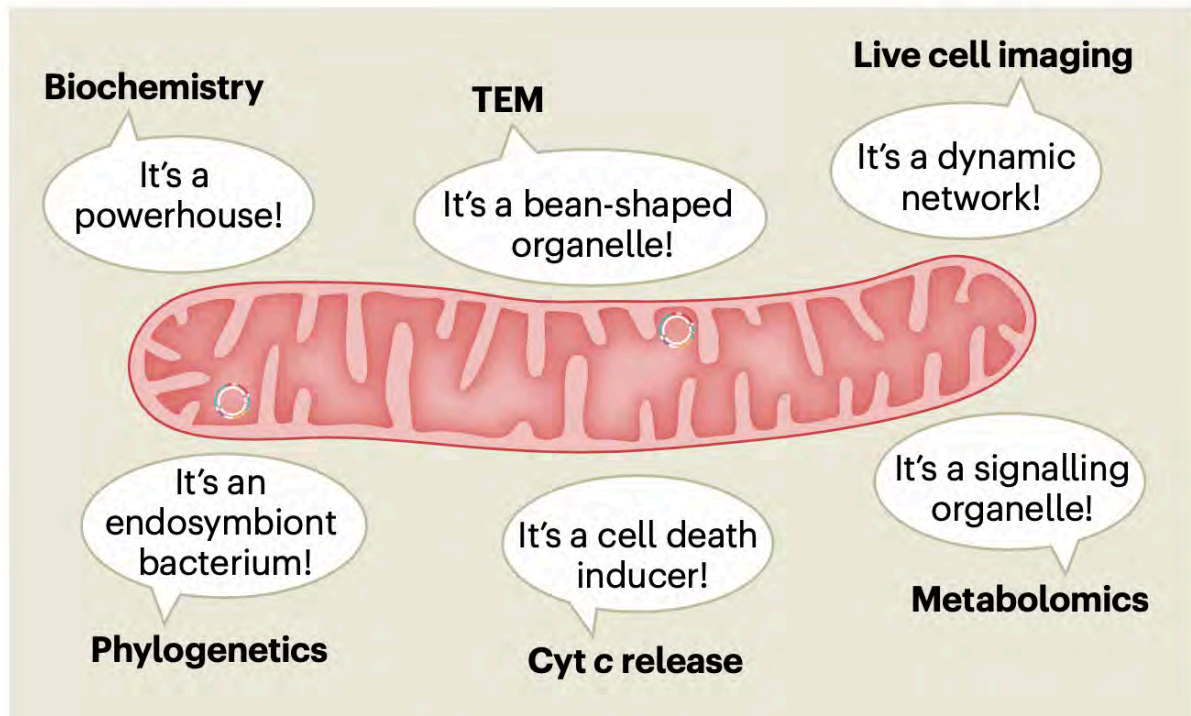
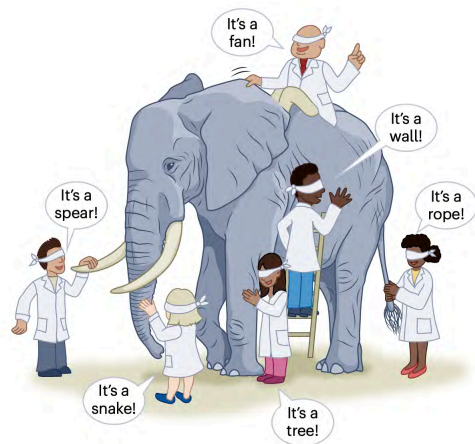
Mito-nuclear unit



Organelle Cell Organism

How do we research mitochondria?





Integrated perspective

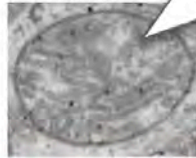
It's a family of organelles that exist as distinct mitochondrial phenotypes, defined by their molecular and morphological features, activities, functions and behaviours

Biochemistry



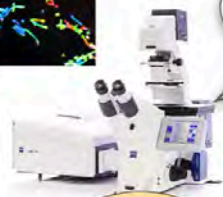
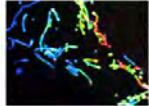
It's a powerhouse!

TEM



It's a bean-shaped organelle!

Live cell imaging

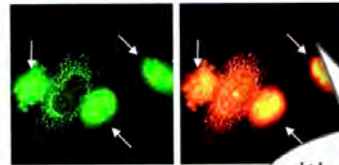
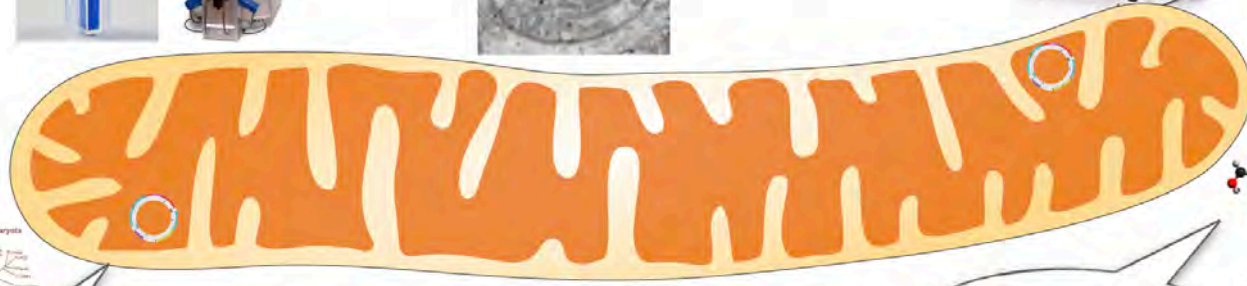


It's a dynamic network!

Phylogenetics



It's an endosymbiont bacterium!



Cyt c release

It's a cell death inducer!

It's a signaling organelle!



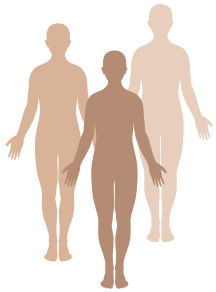
Metabolomics

It's a genetically-related family of organelles that perform dozens of functions, and specialize as cell type- and tissue-specific **mitotypes**.

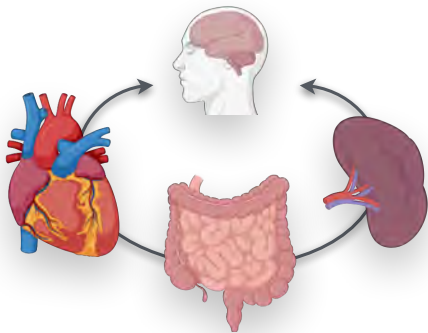


INTERGRADED PERSPECTIVE

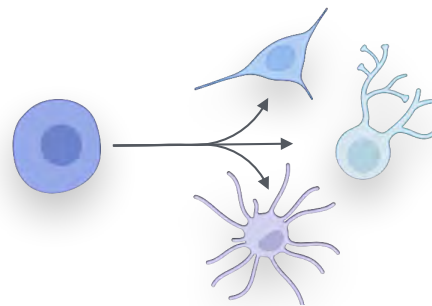
a Domains of human health



- Development and growth
- Physical activity
- Wound healing
- Immunity
- Cardiovascular fitness
- Locomotion
- Digestion
- Sleep
- Cognition
- Learning and memory
- Social interactions
- Others...



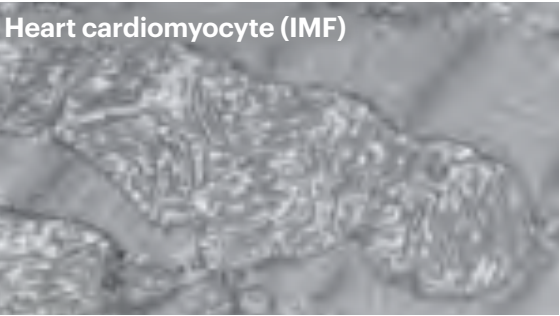
Organ systems



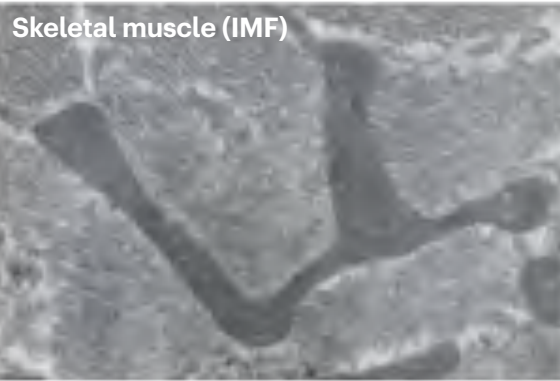
Cell types

Mitotypes?

Heart cardiomyocyte (IMF)



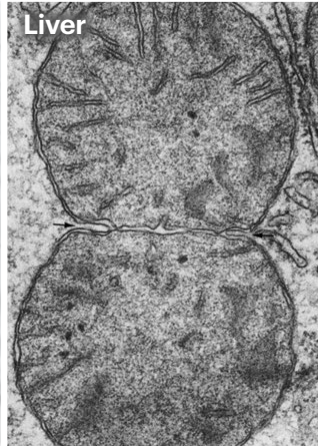
Skeletal muscle (IMF)



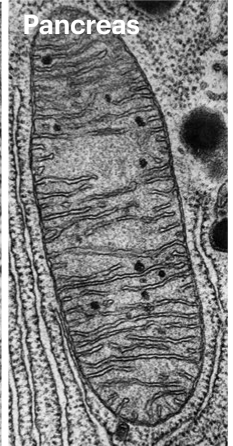
143B-ρ⁰



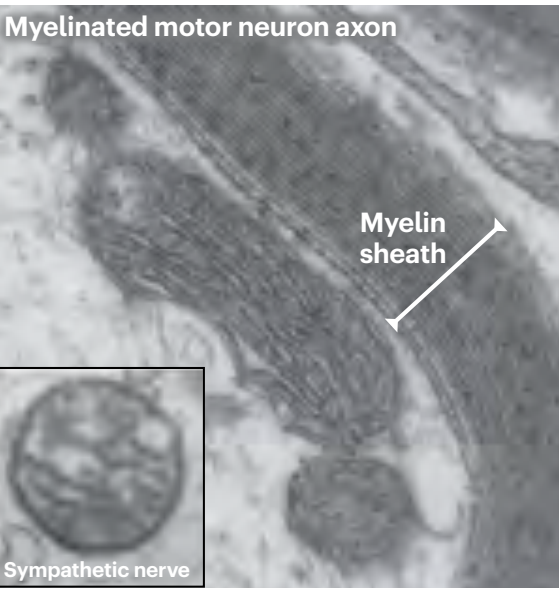
Liver



Pancreas

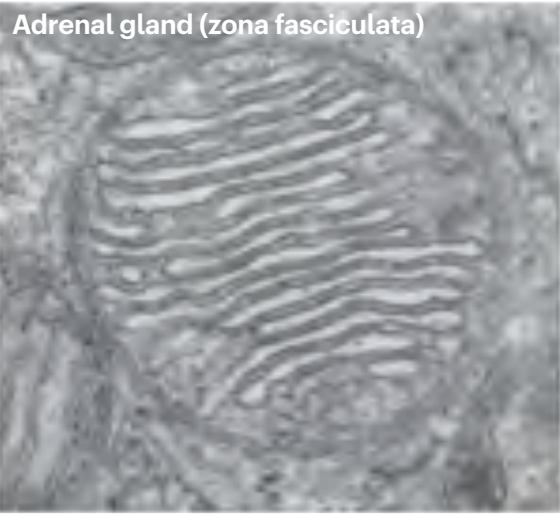


Myelinated motor neuron axon

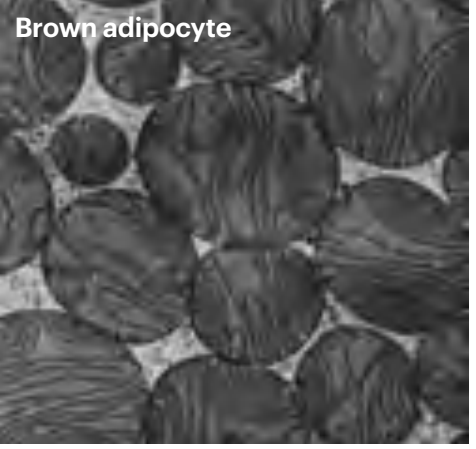


Myelin sheath

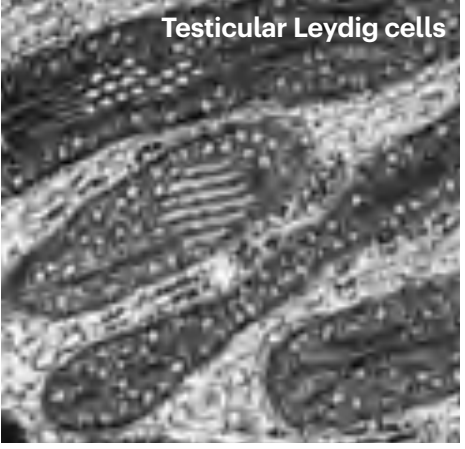
Adrenal gland (zona fasciculata)



Brown adipocyte



Testicular Leydig cells



Sympathetic nerve

Heart cardiomyocyte (IMF)

This electron micrograph shows a heart cardiomyocyte (IMF) and a myelinated motor neuron axon. The cardiomyocyte is characterized by its striated appearance, with numerous mitochondria and a complex network of sarcoplasmic reticulum. The myelinated motor neuron axon is visible as a dark, cylindrical structure with a distinct myelin sheath. The image is presented in grayscale, highlighting the intricate cellular structures.

Myelinated motor neuron axon

Myelinated motor neuron axon



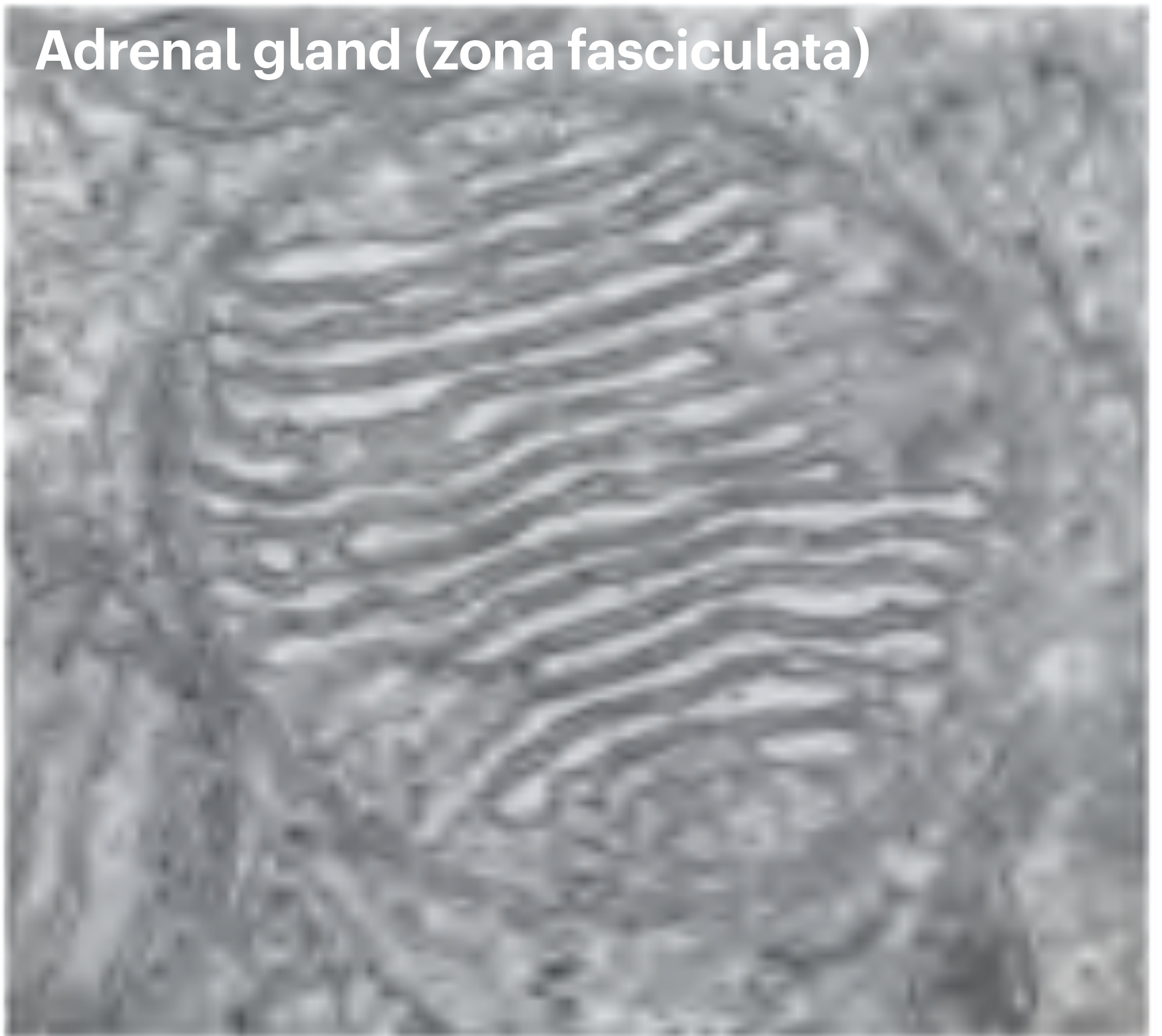
Myelin sheath

Sympathetic nerve

Adrena



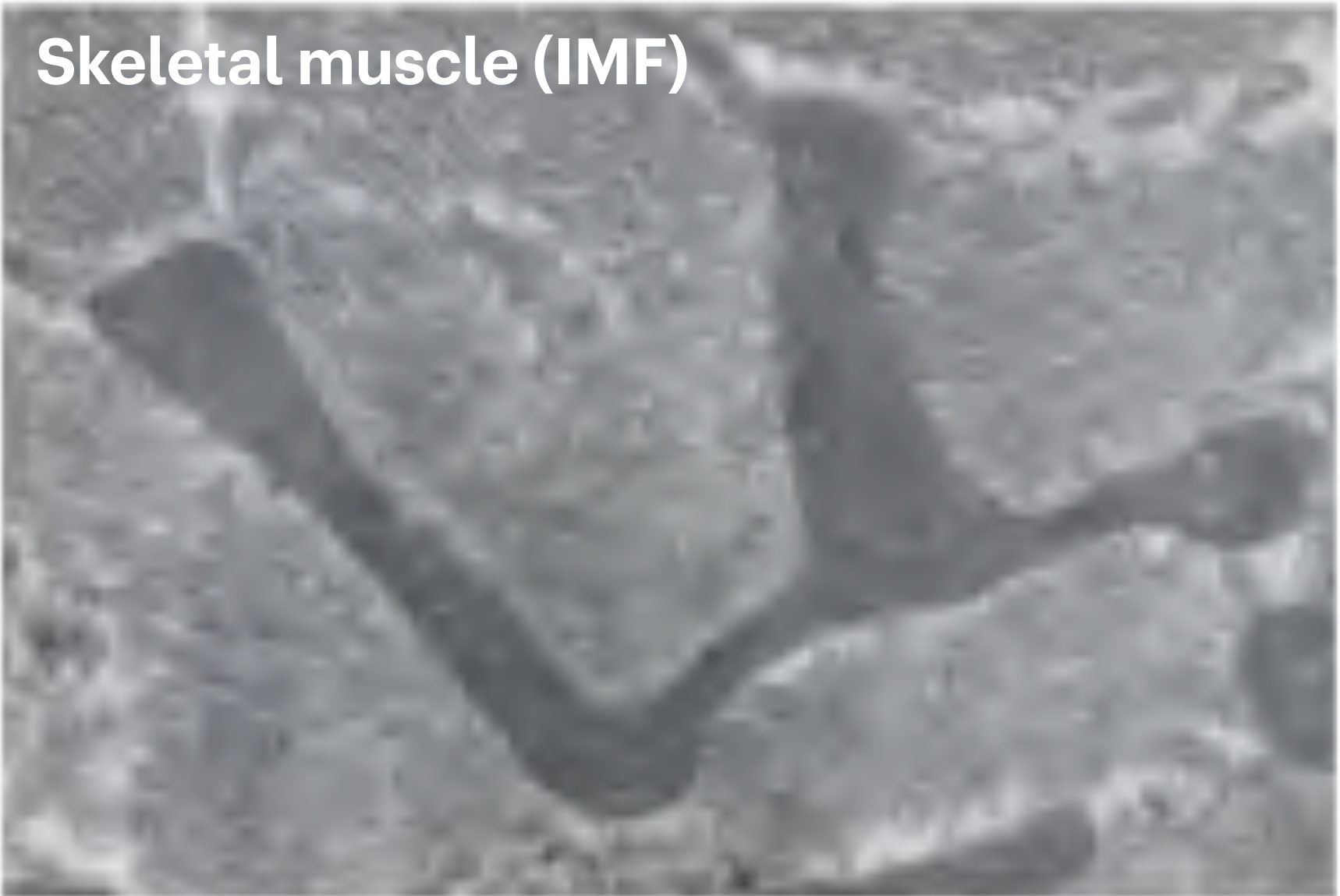
Adrenal gland (zona fasciculata)



Br



Skeletal muscle (IMF)



143B

Adrenal gland (zona fasciculata)

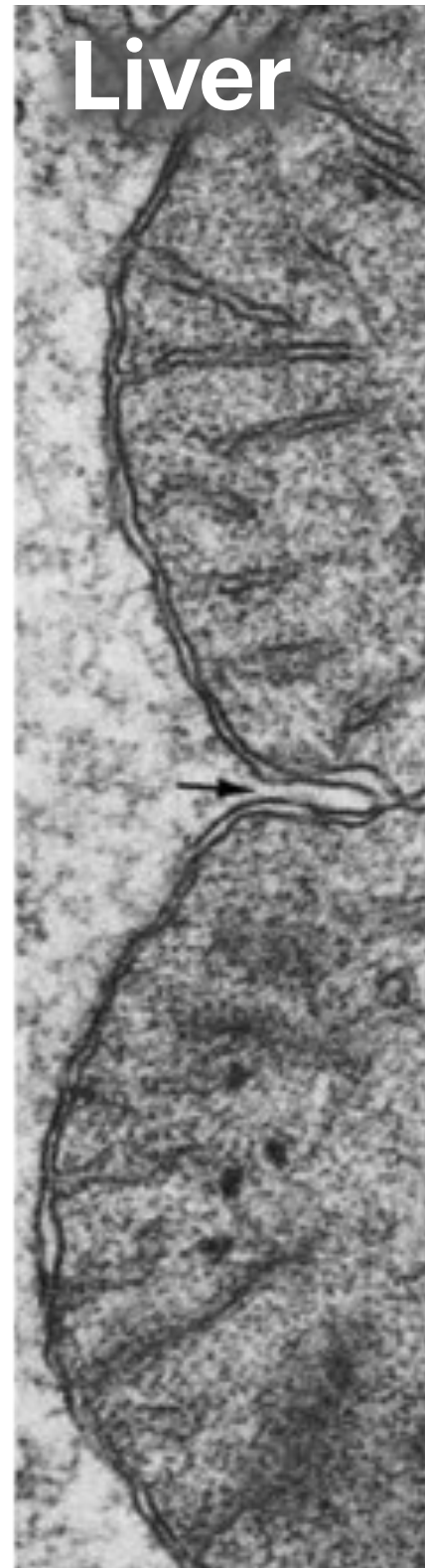


Brow

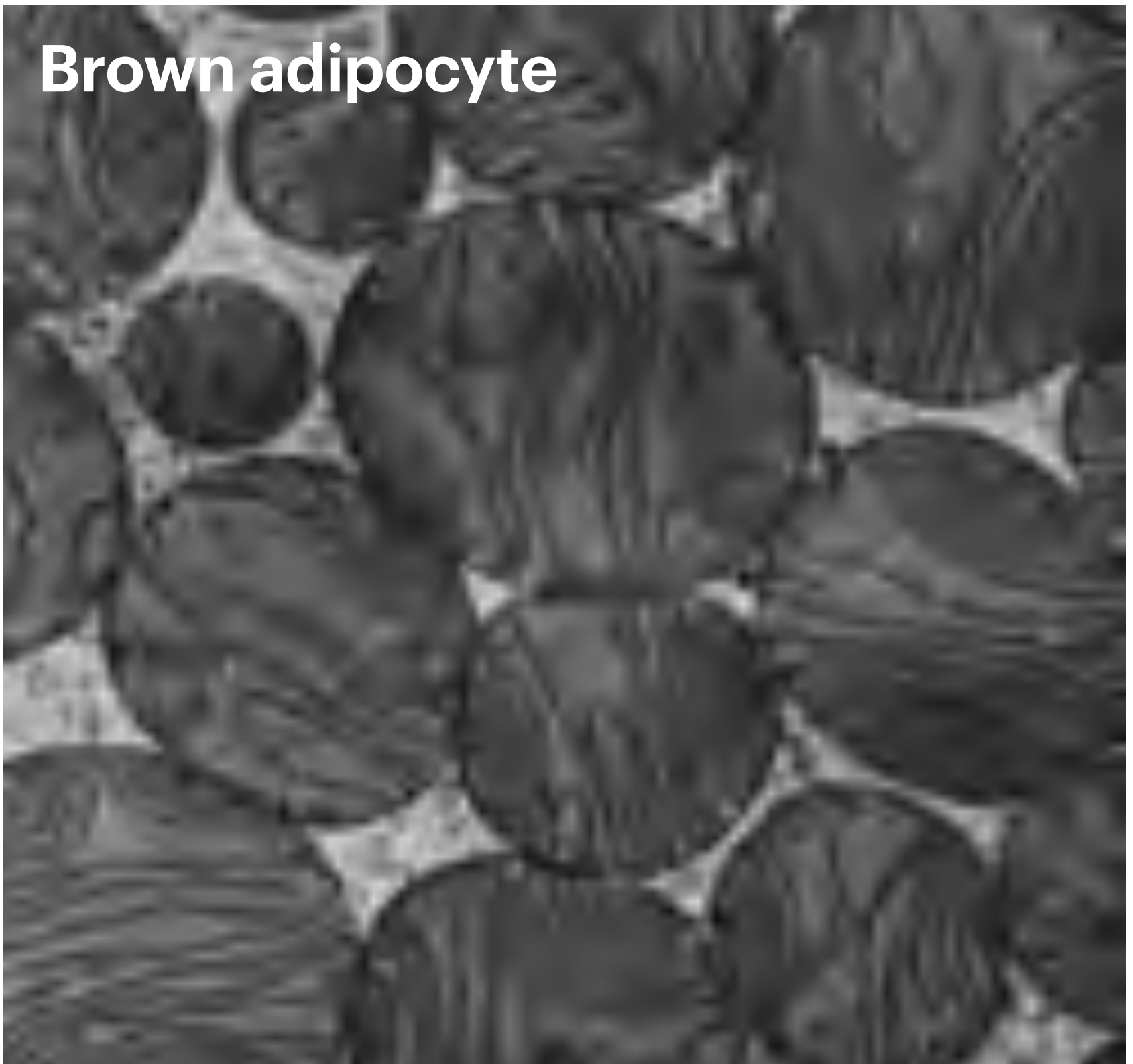
143B- ρ^0



Liver



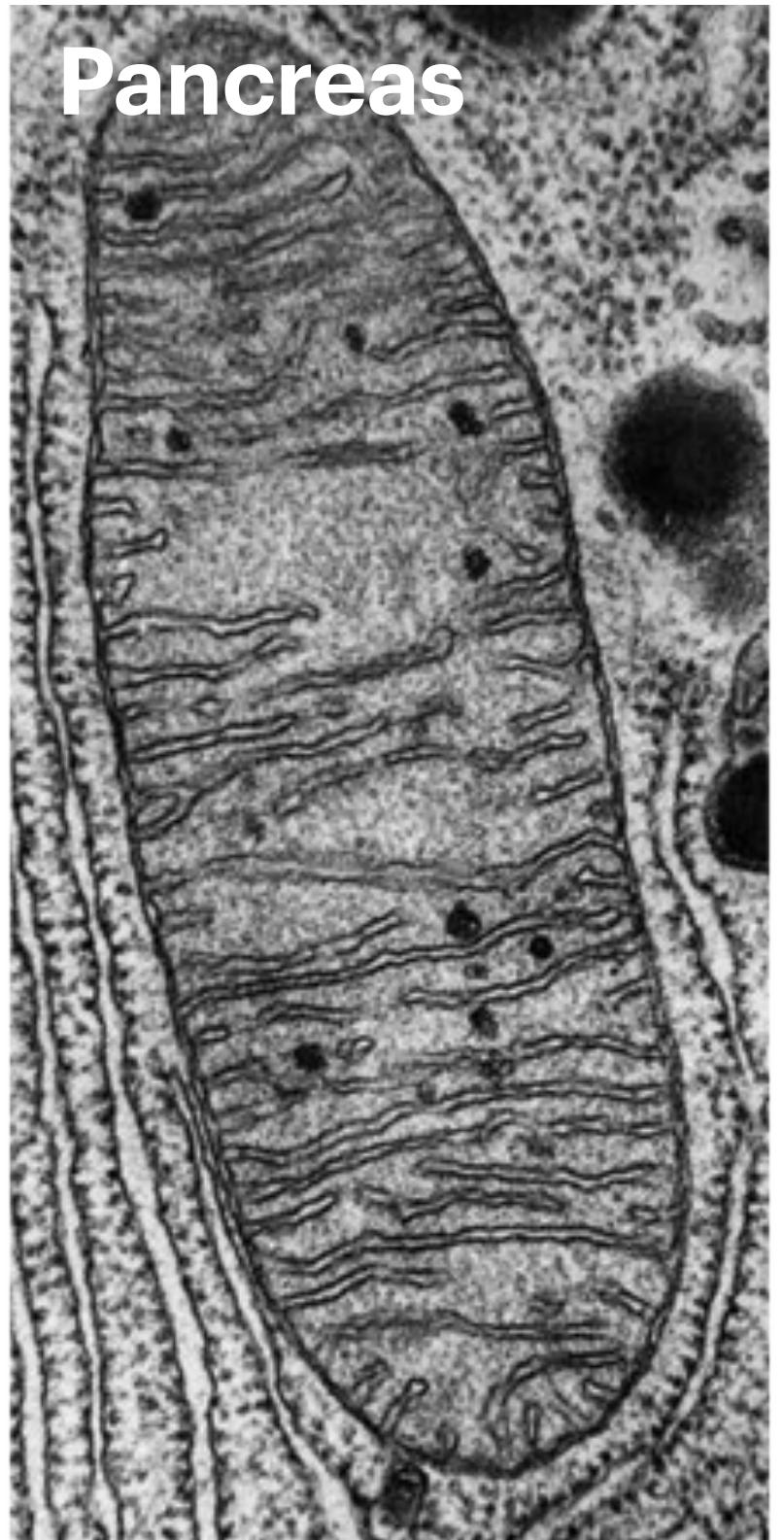
Brown adipocyte



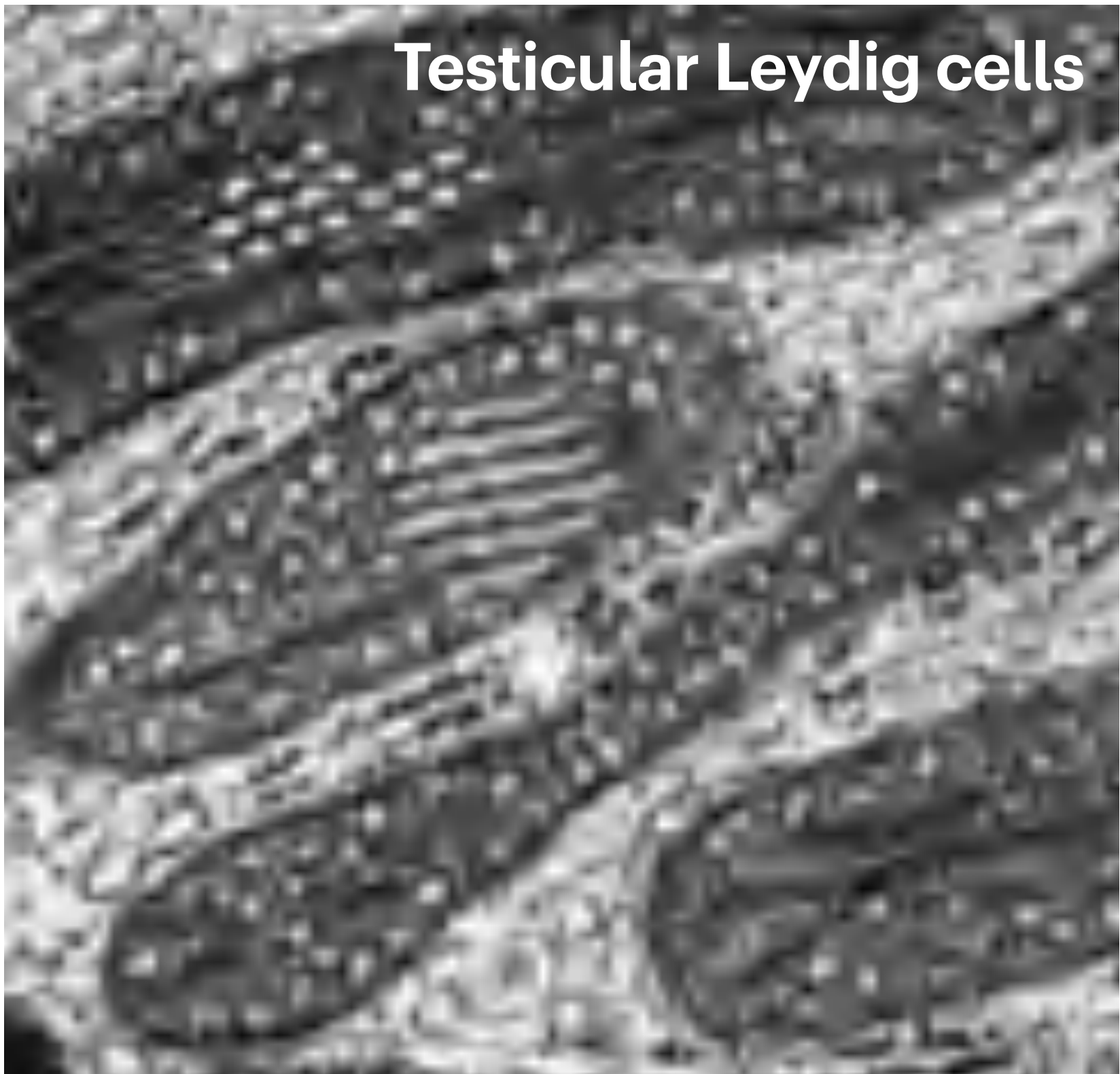
Liver



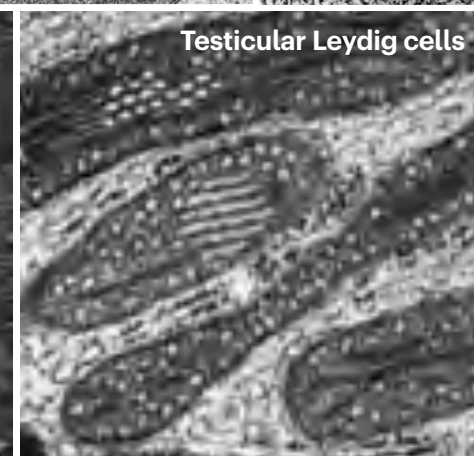
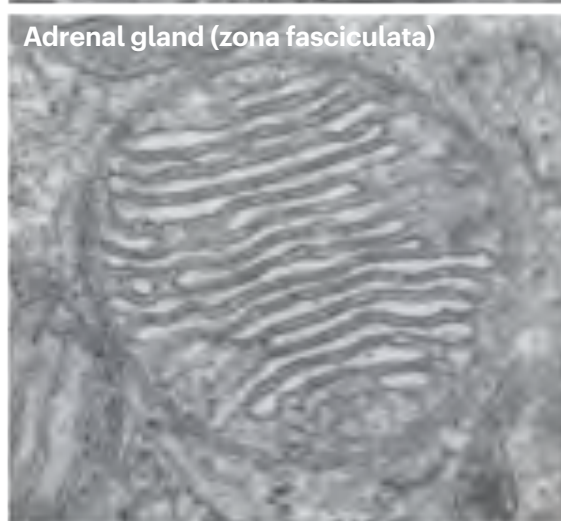
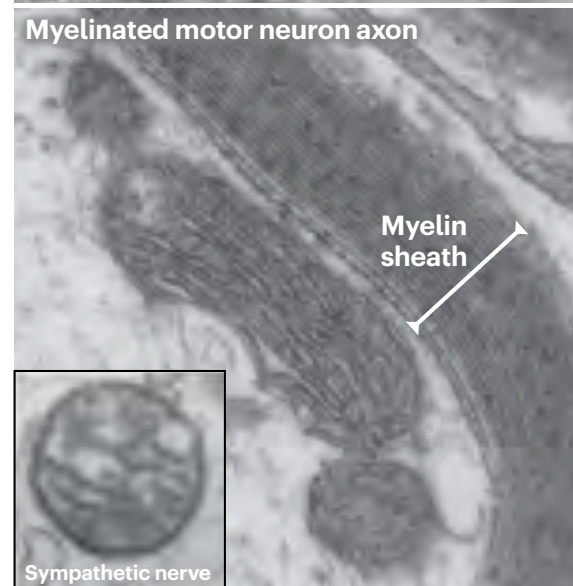
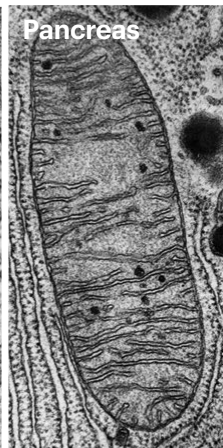
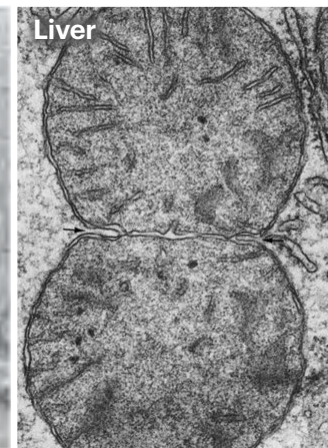
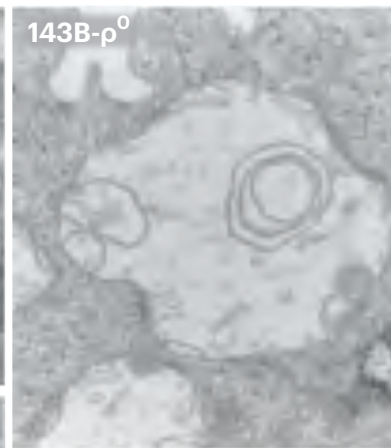
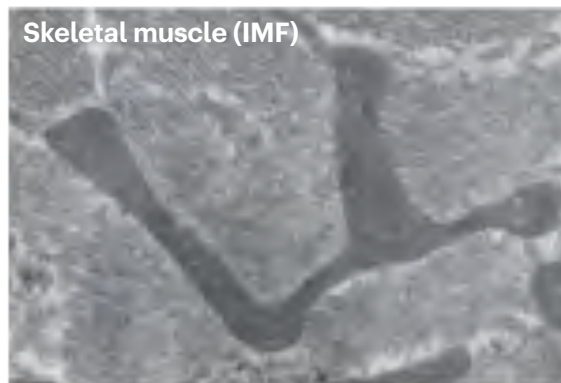
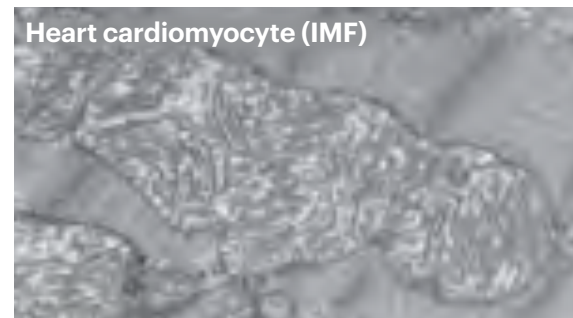
Pancreas



Testicular Leydig cells



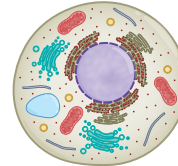
The many faces of mitochondria



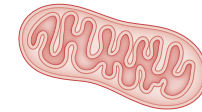
Analogous levels of biology



Organism



Cell types and subtypes

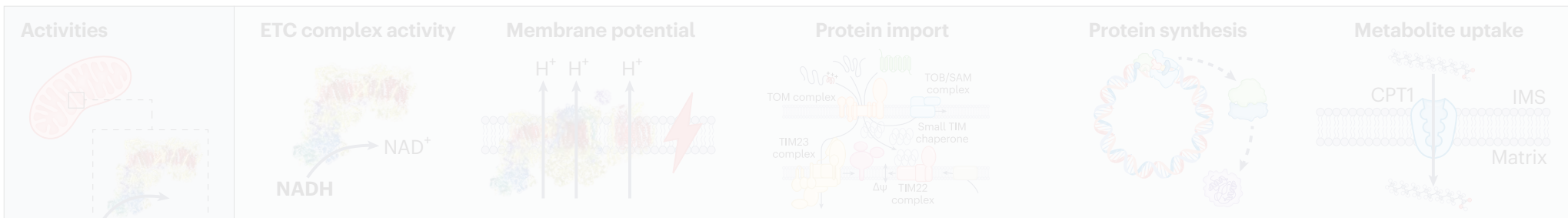
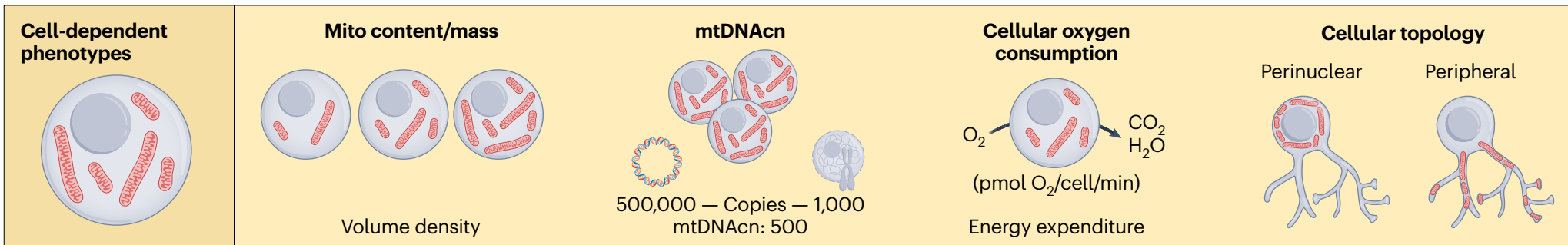


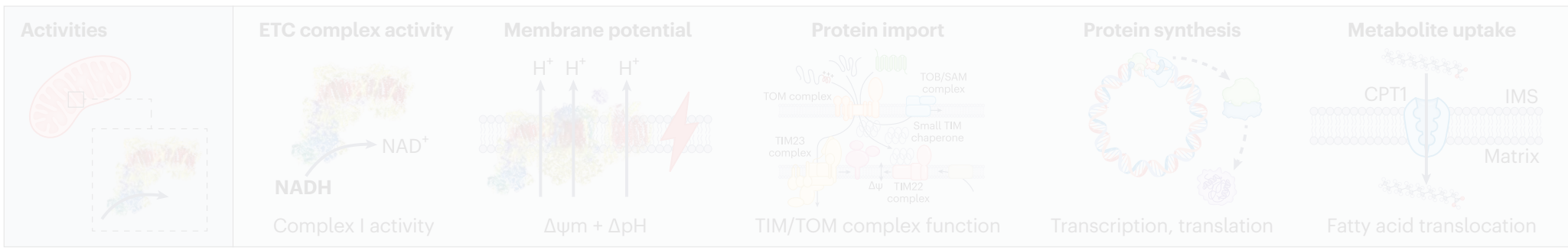
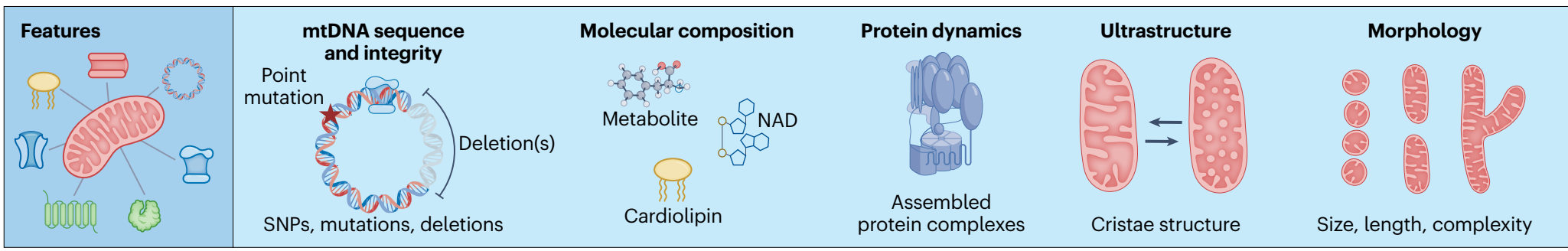
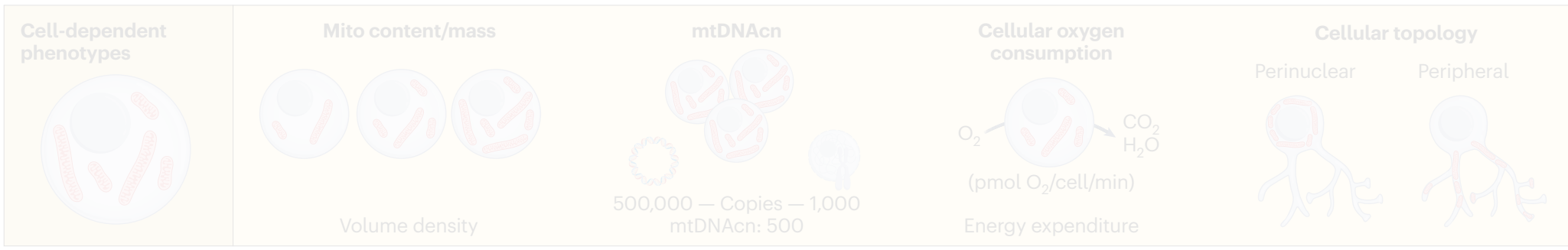
Mitochondrial phenotypes

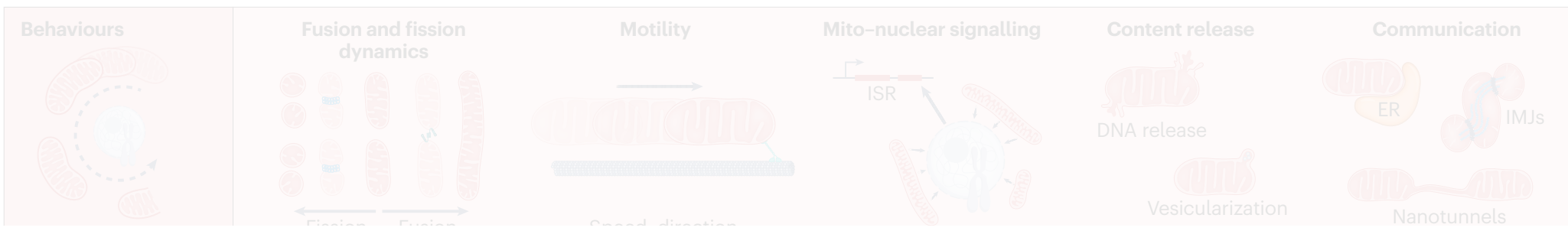
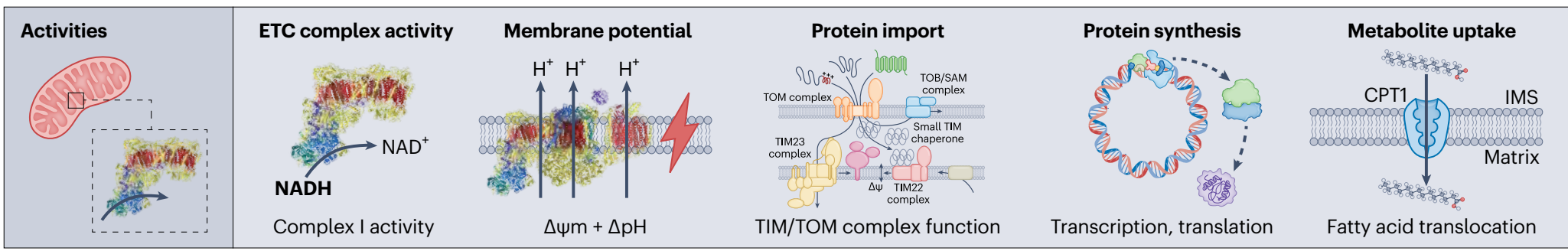
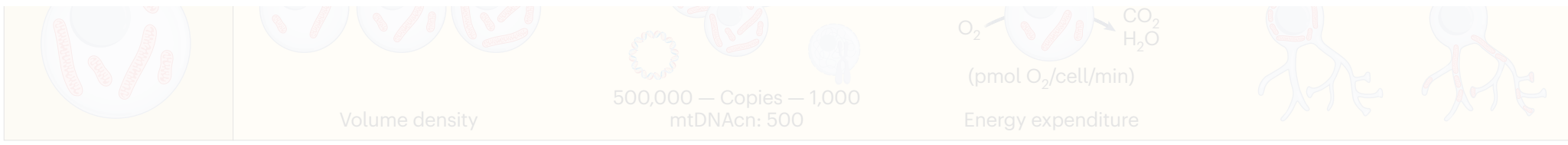
| | | | |
|-------------------------------------|---|--|--|
| Features | Body characteristics Height, body mass index, hydration level, muscle mass, biological sex | Molecular components that define cell types and subtypes Cell surface receptors, gene expression patterns, DNAm epigenetic marks | Static, molecular characteristics that define mitochondrial phenotypes RNA, proteins, OxPhos subunits, mtDNA integrity, morphology, lipid composition |
| Activities | Organ-level processes Skeletal muscle contraction, insulin secretion, cardiac output, peristalsis | Sub-cellular processes Transcription, translation, autophagy, receptor-mediated signal transduction | Processes of individual molecular components ETC enzyme kinetics, other enzymes, Fe buffering, DNA repair |
| Functions | Physiological processes Glycemic control, blood pressure, digestion, wound healing, circadian rhythms, sleep | Integrated cellular processes Specific cytokines release, phagocytotic activity, cell migration, contraction | Integrated processes of mitochondria requiring multiple individual activities OxPhos, Fe/S cluster synthesis, ROS production, steroidogenesis, anaplerosis |
| Behaviours | Goal-directed complex set of functions Social behaviours, reproduction, thinking and feeling, walking and running, ageing | Goal-directed processes involving the cell as a whole Differentiation, extravasation, developmental apoptosis | Goal-directed complex processes involving the mitochondrion as a whole Fusion, fission, motility, inter-organelle signalling |
| Context-dependent phenotypes | Physiological states driven by social and environmental demands Homeostasis, allostasis and allostatic load | Cellular characteristics relevant only at the organ level Hyperplasia, inflammation, elasticity | Mitochondrial characteristics relevant in the context of the host cell Mitochondrial content, mtDNAcn, cellular O ₂ consumption |

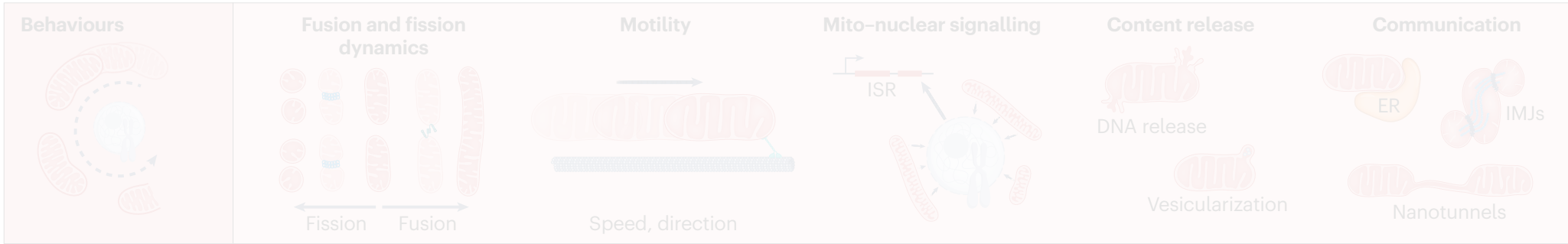
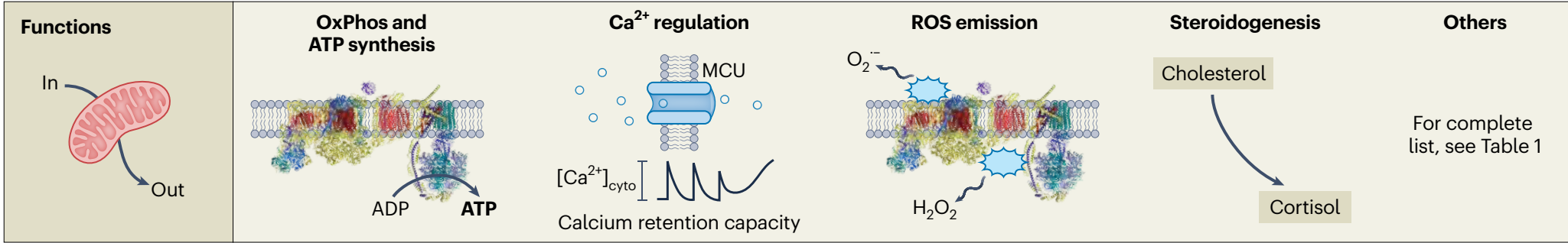
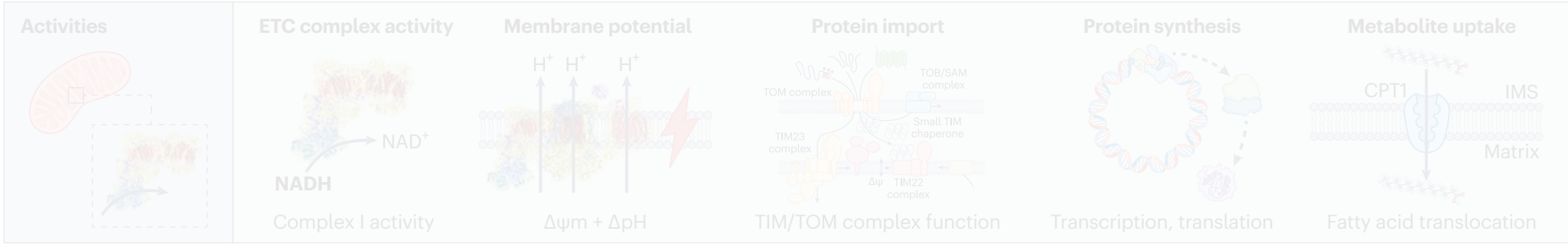
Analogous levels of biology

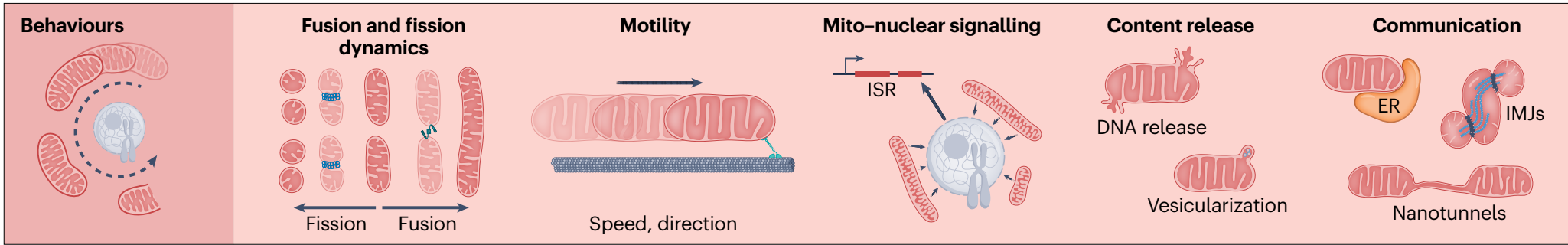
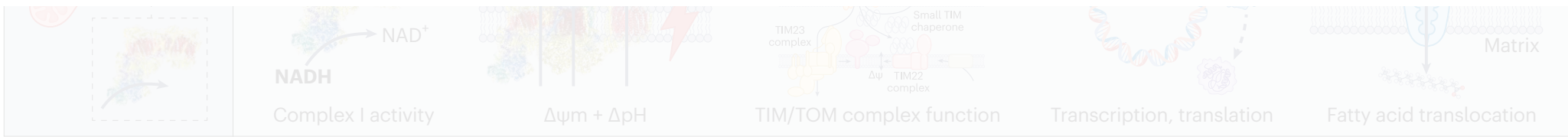
| | Organism | Cell types and subtypes | Mitochondrial phenotypes |
|------------------------------|---|--|--|
| Features | Body characteristics—height, body mass index, hydration level, thyroid mass, biological sex | Molecular components that define cell types and subtypes Cell surface receptors, gene expression patterns, DNA sequence marks | Basic, molecular characteristics that define mitochondrial phenotypes mtDNA sequence, Cristae pattern, mtDNA integrity, morphology, lipid composition |
| Activities | Organ-level processes Blood muscle contraction, insulin secretion, cardiac output, peristalsis | Sub-cellular processes Transcription, translation, autophagy, receptor-mediated signal transduction | Processes of individual molecular components E.C. activity, metabolic flux, apoptosis, FA trafficking, DNA repair |
| Functions | Physiological processes Climate control, blood pressure, digestion, wound healing, circadian rhythms, sleep | Integrated cellular processes Gene/protein control, energy homeostasis, activity, cell migration, contraction | Integrated processes of mitochondria Handling multiple individual activities OxPhos, FA β oxidation, synthesis, ROS production, peroxisomal interactions |
| Behaviours | Goal-directed complex set of functions Social behaviours, reproduction, thinking and feeling, working and learning, ageing | Goal-directed processes involving the cell as a whole Differentiation, adaptation, developmental apoptosis | Goal-directed complex processes involving the mitochondrion as a whole Energy balance, mobility, inter-organellar signalling |
| Context-dependent phenotypes | Physiological status driven by social and environmental demands Homeostasis, allostasis and allostasis load | Cellular characteristics relevant only at the organ level Proliferation, inflammation, elasticity | Mitochondrial characteristics relevant in the context of the host cell Mitochondrial content, mtDNA, cellular O $_2$ consumption |











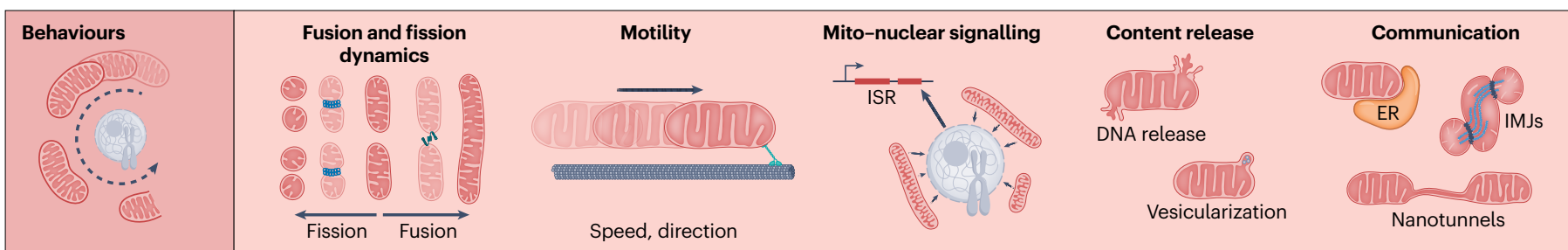
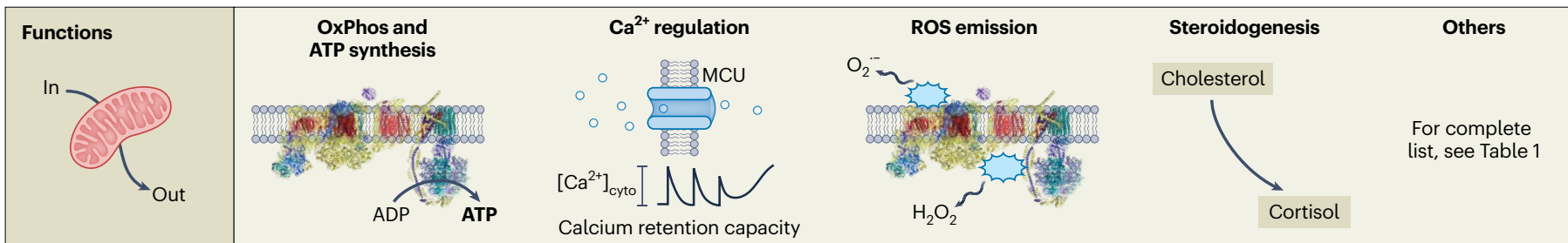
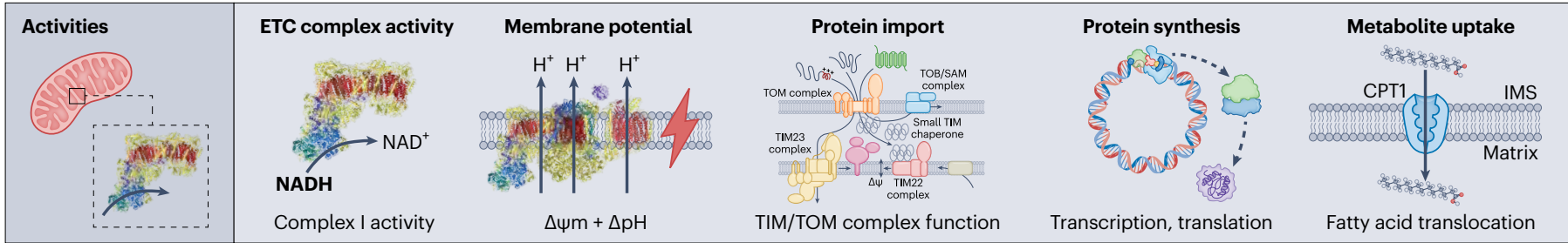
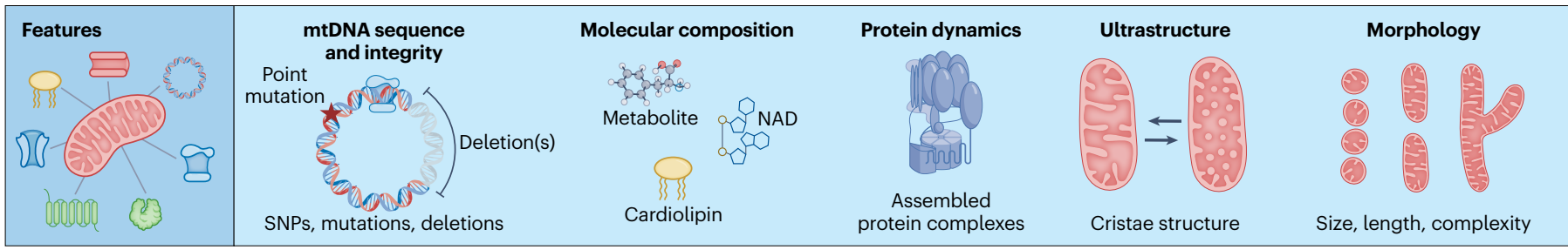
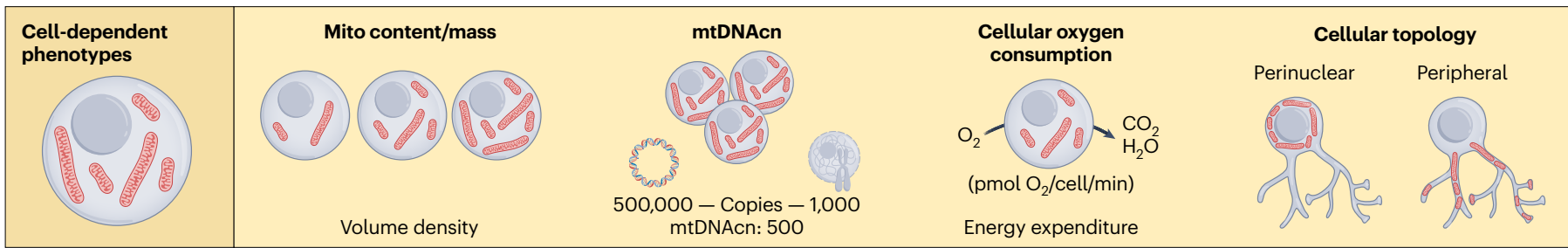


Table 1 | Mitochondrial functions and behaviours

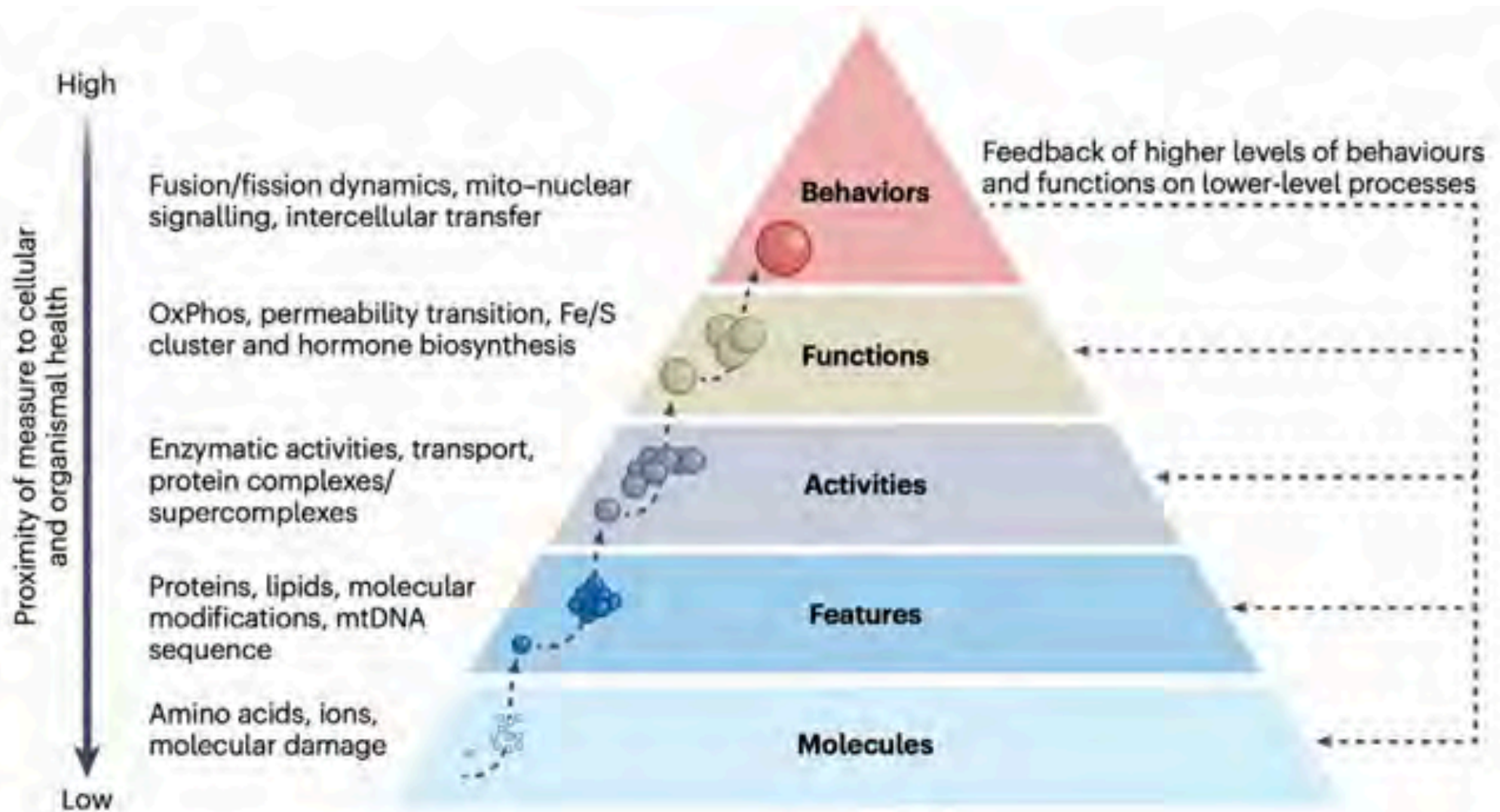
| | Description | Reviewed in ref(s). | Methods described in ref(s). |
|--|--|---------------------|------------------------------|
| Functions | | | |
| ^aMembrane potential generation | Formation of the electrochemical gradient ($\Delta\Psi_m + \Delta pH$) across the IMM, usually by the electron pumping capacity of the respiratory complexes I, III and IV, but also by other processes including through ATP hydrolysis by the F_0F_1 ATP synthase (complex V). | 104 | 105,106 |
| Amino acid metabolism | Lysine metabolism (lysine- α -ketoglutarate reductase, encoded by <i>AASS</i>). Electrogenic malate–aspartate shuttle system, which is important for balancing pyridine dinucleotide redox states across subcellular compartments. Branched-chain keto and amino acids. Choline and derivatives as structural precursors for lipoproteins, membrane lipids and the neurotransmitter acetylcholine. Betaine as osmoregulator and an intermediate in the cytosolic transsulfuration pathway. | 107–111 | 112–119 |
| Ascorbate metabolism | L-ascorbate (vitamin C) biosynthesis in many plants and animals, but not in primates, which serves as osmoregulator and antioxidant. Mitochondria may recycle oxidized (dehydro)ascorbic acid. | 120 | 121,122 |
| Bicarbonate metabolism | Production of bicarbonate (HCO_3^-) by mitochondrial carbonic anhydrase V (encoded by <i>CA5A</i>), used as a cofactor for anaplerotic reactions (for example, ureagenesis and gluconeogenesis) and acid–base balance. The TCA cycle is an important contributor to cellular/extracellular acidification due to CO_2 production. | 123 | – |
| Calcium uptake and extrusion | Uptake of cytoplasmic Ca^{2+} via the mitochondrial calcium uniporter in a $\Delta\Psi_m$ -dependent manner; extrusion by the sodium/calcium exchanger NCLX (encoded by <i>SLC8B1</i>). | 124–126 | 127,128 |
| Hydrogen sulfide detoxification | Mitochondrial sulfide quinone oxidoreductase (encoded by <i>SQOR</i>) oxidizes hydrogen sulfide to glutathione persulfide by reducing CoQ. | 129–132 | 133 |
| Heat production | Heat generation is stimulated by uncoupling $\Delta\Psi_m + \Delta pH$ from ATP synthesis (thereby increasing electron flux and respiration) by UCP1 (encoded by <i>UCP1</i>), the ADP/ATP carrier (<i>AAC</i> , also <i>ANT1</i>), or by creatine-dependent substrate cycling and other futile cycles. | 134–137 | 138 |
| Intermediate metabolism | Enzymatic interconversion of metabolic intermediates to enable the synthesis of specific macromolecules, including five major anaplerotic ones. This includes the conversion of pyruvate into oxaloacetate by pyruvate carboxylase (encoded by <i>PC</i>), a critical step for de novo glucose synthesis (gluconeogenesis); citrate export to the cytoplasm where it is used for lipid synthesis or converted to acetyl-CoA for acetylation reactions; synthesis of itaconate, a derivative of <i>cis</i> -aconitate; succinate, α -ketoglutarate and others that participate in a variety of signalling | 25,139,140 | 141,142 |

| | | | |
|---|--|---------------|---------|
| Lipid oxidation | Beta-oxidation of long-chain, medium-chain and short-chain fatty acids into acetyl-CoA. | 145 | 146 |
| Lipid synthesis | Synthesis of cardiolipin and phosphatidylethanolamine from ER precursors in the IMM. | 147-150 | - |
| mtDNA maintenance and expression | mtDNA replication, transcription, protein synthesis and assembly of the OxPhos system. | 151,152 | 153,154 |
| Na⁺import/export | Sodium (Na ⁺) uptake and release against cytoplasmic Ca ²⁺ by the sodium/calcium exchanger protein NCLX (encoded by <i>SLC8B1</i>) or by Na ⁺ /H ⁺ antiporter (molecular identity pending). | 124,155 | 156 |
| Neurotransmitter synthesis and degradation | Synthesis of the cofactor BH4 (tetrahydrobiopterin), used by hydrolase enzymes to synthesize catecholamines and neurotransmitters (serotonin, melatonin, norepinephrine and epinephrine) and nitric oxide. Mitochondria with OMM-anchored monoamine oxidases (encoded by <i>MAOA</i> and <i>MAOB</i> , donate electrons and contribute to electron flow in the ETC) also degrade catecholamines. Mitochondria also participate in GABA metabolism. | 9,157 | 158,159 |
| One-carbon metabolism and pyrimidine synthesis | The one-carbon metabolism connects the synthesis of nucleotides (purine and pyrimidine), amino acids (methionine, serine and glycine), S-adenosyl-methionine and folate. Ubiquinone-mediated oxidation of dihydroorotate to orotate by dihydroorotate dehydrogenase (encoded by <i>DHODH</i>) is a key step in pyrimidine synthesis. | 160-163 | 164 |
| OxPhos | Transduction of $\Delta\Psi_m + \Delta pH$ generated by the electron transport chain (ETC, also 'respiratory chain') into ATP synthesis by the F ₀ F ₁ ATP synthase (complex V), abbreviated as OxPhos. | 165 | 166 |
| Oxygen sensing | The electron transport and free-radical generation by ETC complexes I and III is modulated by the partial pressure of oxygen, which can limit respiration at very low partial pressures of O ₂ . | 167-170 | - |
| Permeability transition | Opening of the high-conductance permeability transition pore (PTP), which dissipates membrane potential and promotes the release of intracristae and matrix-located components into the cytoplasm. | 171,172 | 173-175 |
| Protein import | Import, processing and folding of nuclear-encoded polypeptides from the cytoplasm by the translocator of the inner membrane (TIM) and outer membrane (TOM) complexes and associated proteins. | 176 | - |
| Redox homeostasis | Re-oxidation of enzymes and/or their redox cofactors (involved in anabolic and catabolic reactions) by the electron acceptors CoQ and cytochrome c (encoded by <i>CYTC</i>) within the mitochondrial respiratory chain, and production of NADPH by <i>NNT</i> . | 177,178 | - |
| Respiration | Electrons stored in reducing equivalents NADH and FADH ₂ , or derived from diverse redox reactions are sequentially delivered to respiratory complex I and CoQ, or cytochrome c, respectively, to promote the reduction of molecular oxygen at cytochrome c oxidase (complex IV). | 179,180 | 181 |
| ROS production | Production and release of ROS (H ₂ O ₂ , O ₂ ⁻ , others) mainly at respiratory chain complexes I and III. | 182,183 | 184 |
| Steroidogenesis | Production of pregnanolone from cholesterol imported via IMM steroidogenic pathway intermediates (encoded by <i>STARD</i>) followed by enzymatic conversion | 33,34,185,186 | 187 |

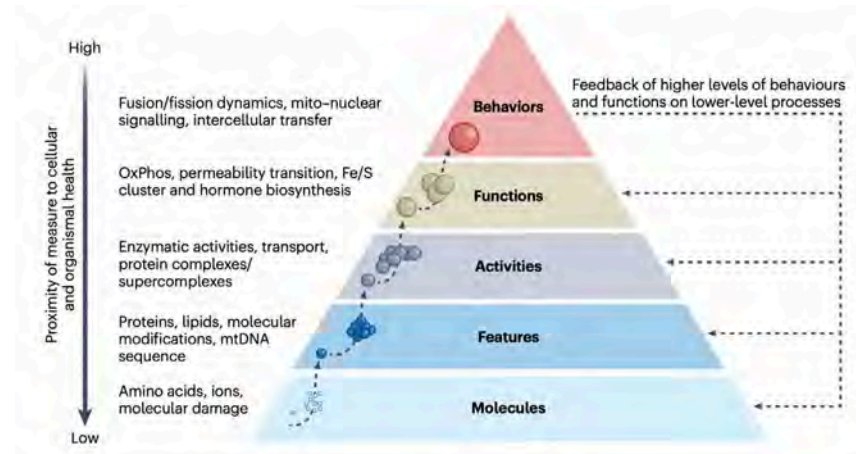
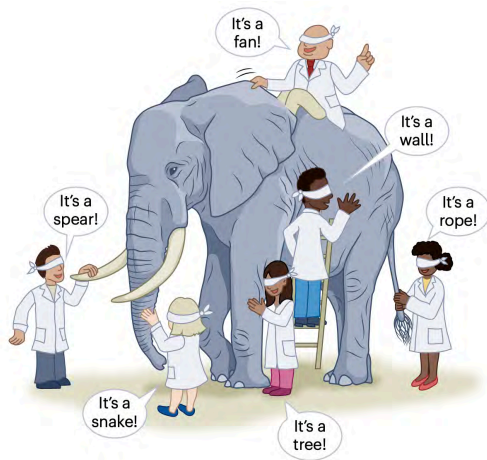
| | | | |
|--------------------------------------|---|-----------------|---------|
| ROS production | Production and release of ROS (H ₂ O ₂ , O ₂ ⁻ , others) mainly at respiratory chain complexes I and III. | 182,183 | 184 |
| Steroidogenesis | Production of pregnanolone from cholesterol imported via IMM steroidogenic acute regulatory protein (encoded by <i>STAR</i>) followed by enzymatic transformation by P450 _{ssc} (encoded by <i>CYP11A1</i>) in the matrix. Intermediate or terminal steps for some steroids occur in the ER. Cytochrome P450 family members participate also in xenobiotic metabolism as well as bile acid and vitamin D biosynthesis. | 33,34,185,186 | 187 |
| Behaviours | | | |
| Antiviral signalling | Assembly of the mitochondrial antiviral signal (encoded by <i>MAVS</i>) adaptor protein on the OMM to potentiate downstream signalling, and activation of nuclear interferon pathways in the nucleus by mtDNA release. | 39,188 | - |
| Apoptotic signalling | Release of cytochrome c (encoded by <i>CYCS</i>), apoptosis-inducing factor (encoded by <i>AIF</i>), and other proteins that trigger different forms of cell death by acting on cytoplasmic and nuclear effectors. | 189,190 | - |
| Cristae remodelling | Dynamic remodelling of IMM cristae junctions, cristae shape and distribution via the combined action of optic atrophy 1 (encoded by <i>OPA1</i>) and mitochondrial contact site and cristae organizing system (MICOS) proteins. | 103,191 | 95 |
| DNA signalling | mtDNA extrusion in the cytoplasm, particularly in the form of oxidized mtDNA fragments via proteinaceous pores forming across the IMM and OMM, which trigger inflammasome activation. | 189,190,192,193 | 175 |
| Epigenetic remodelling | Transduction of mitochondrial states into changes in epigenome via several functions including metabolic intermediates, DNA release, ROS production and others. | 30,194 | - |
| Inter-organelle communication | Exchange of information between mitochondria and other organelles, particular the ER, where mitofusin 2 (encoded by <i>MFN2</i>) plays a key role in tethering organelles. | 195,196 | 197,198 |
| Mitochondrial dynamics | Mitochondrial fusion and fission through OMM-anchored and IMM-anchored GTPase proteins capable of merging or constricting mitochondrial membranes to enact fragmentation of larger organelles into smaller ones. | 191,199–201 | 202 |
| Mito-mito communication | Exchange of information between mitochondria by soluble signals (for example, ROS-induced ROS release, RIRR), by complete membrane fusion, or by physical extensions of thin protein-carrying OMM and IMM membrane protrusions (that is, nanotunnels) and trans-mitochondrial cristae alignment between energized mitochondria. | 203–206 | 207–209 |
| Motility | Movement of energized mitochondria across the cytoplasm via the combined action of motor and adaptor proteins interacting with cytoskeletal elements. | 6,210 | 211 |
| Vesicle formation | Release of MDVs destined to different cellular fates by the action of motor and accessory proteins acting on the OMM and IMM. | 212 | 213,214 |

^aGeneration of mitochondrial membrane potential is the 'mother' of many other functions and behaviours, providing the driving force for the movement of ions, solutes and proteins across the IMM, the driving force for key enzymes and processes, including the phosphorylation of ADP into ATP (OxPhos). Mitochondrial features (that is, molecular components) and activities (individual enzyme and non-enzymatic activities) are too numerous to be comprehensively listed, so only functions and behaviours are included. CoQ, coenzyme Q.

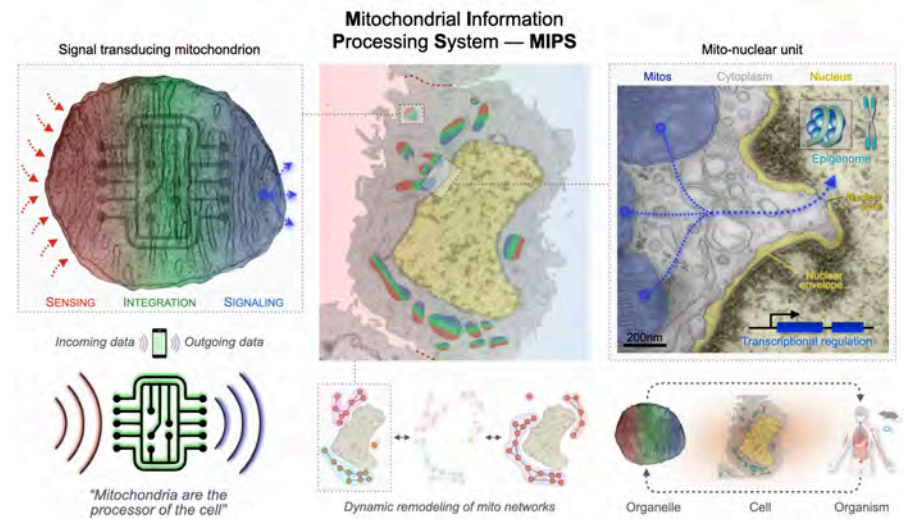
Hierarchy of mitochondrial needs



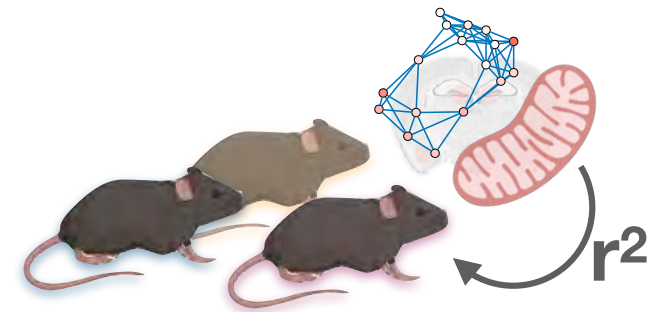
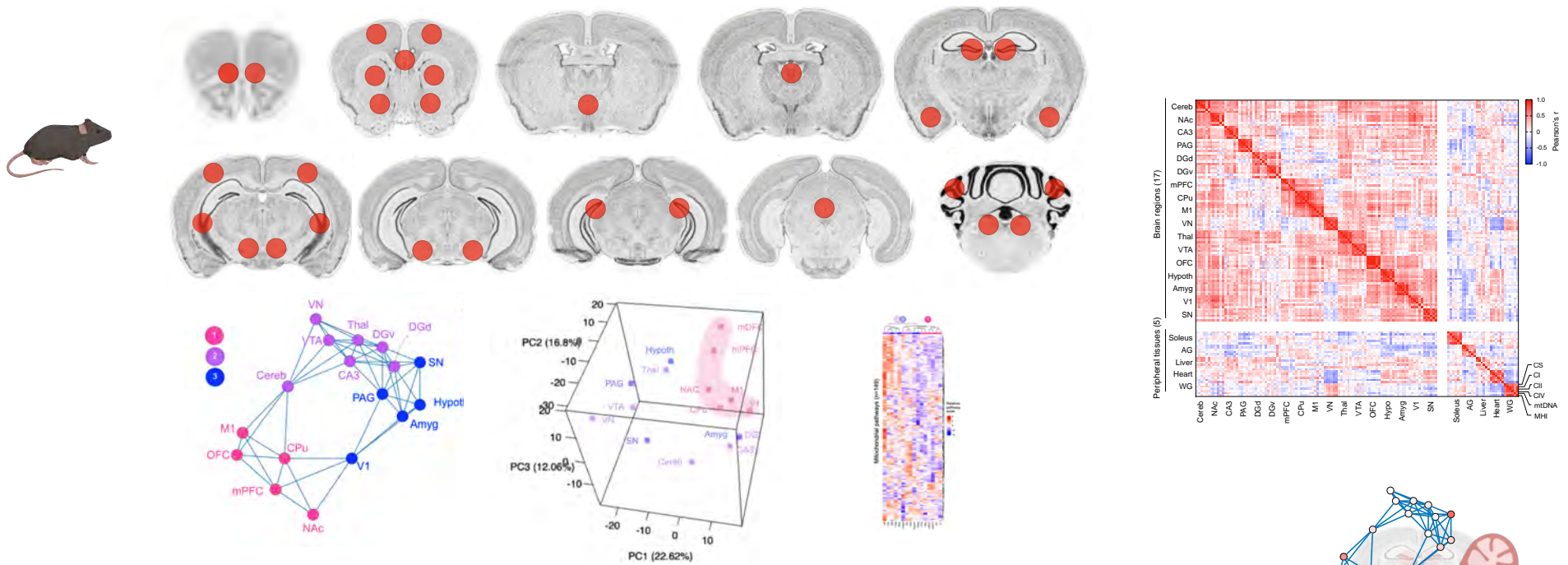
Mitochondria are diverse, multifunctional organelles that transduce information



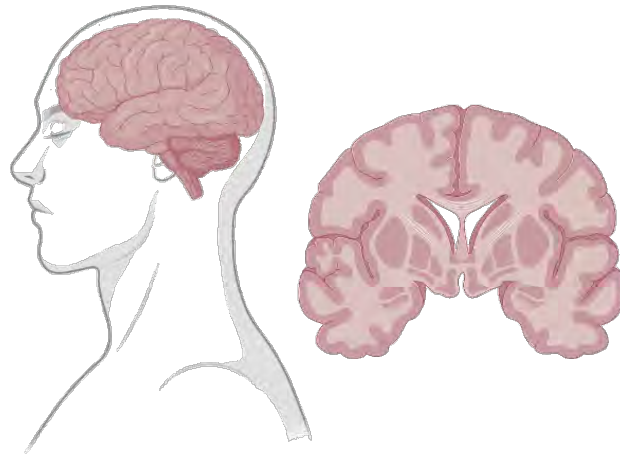
| | | | | | |
|----------------------------------|--|---|--|---|---|
| Cell-dependent phenotypes | Mito content/mass Volume density | mtDNAcn 500,000 – Copies – 1,000 mtDNAcn: 500 | Cellular oxygen consumption O ₂ → CO ₂ + H ₂ O (pmol O ₂ /cell/min) Energy expenditure | Cellular topology Perinuclear Peripheral | |
| Features | mtDNA sequence and integrity Point mutation Deletion(s) SNPs, mutations, deletions | Molecular composition Metabolite NAD Cardiolipin | Protein dynamics Assembled protein complexes | Ultrastructure Cristae structure | Morphology Size, length, complexity |
| Activities | ETC complex activity NADH Complex I activity | Membrane potential H ⁺ , H ⁺ Δμ _m → ΔpH | Protein import TIM/TOM complex function | Protein synthesis Transcription, translation | Metabolite uptake CPT1 IMS Matrix Fatty acid translocation |
| Functions | OxPhos and ATP synthesis ADP → ATP Calcium retention capacity | Ca²⁺ regulation [Ca ²⁺] _{low} Calcium retention capacity | ROS emission H ₂ O ₂ | Steroidogenesis Cholesterol Cortisol | Others For complete list, see Table 1 |
| Behaviours | Fusion and fission dynamics Fission Fusion | Motility Speed, direction | Mito-nuclear signalling ISR | Content release DNA release Vesicularization | Communication ER MJs Nanotunnels |



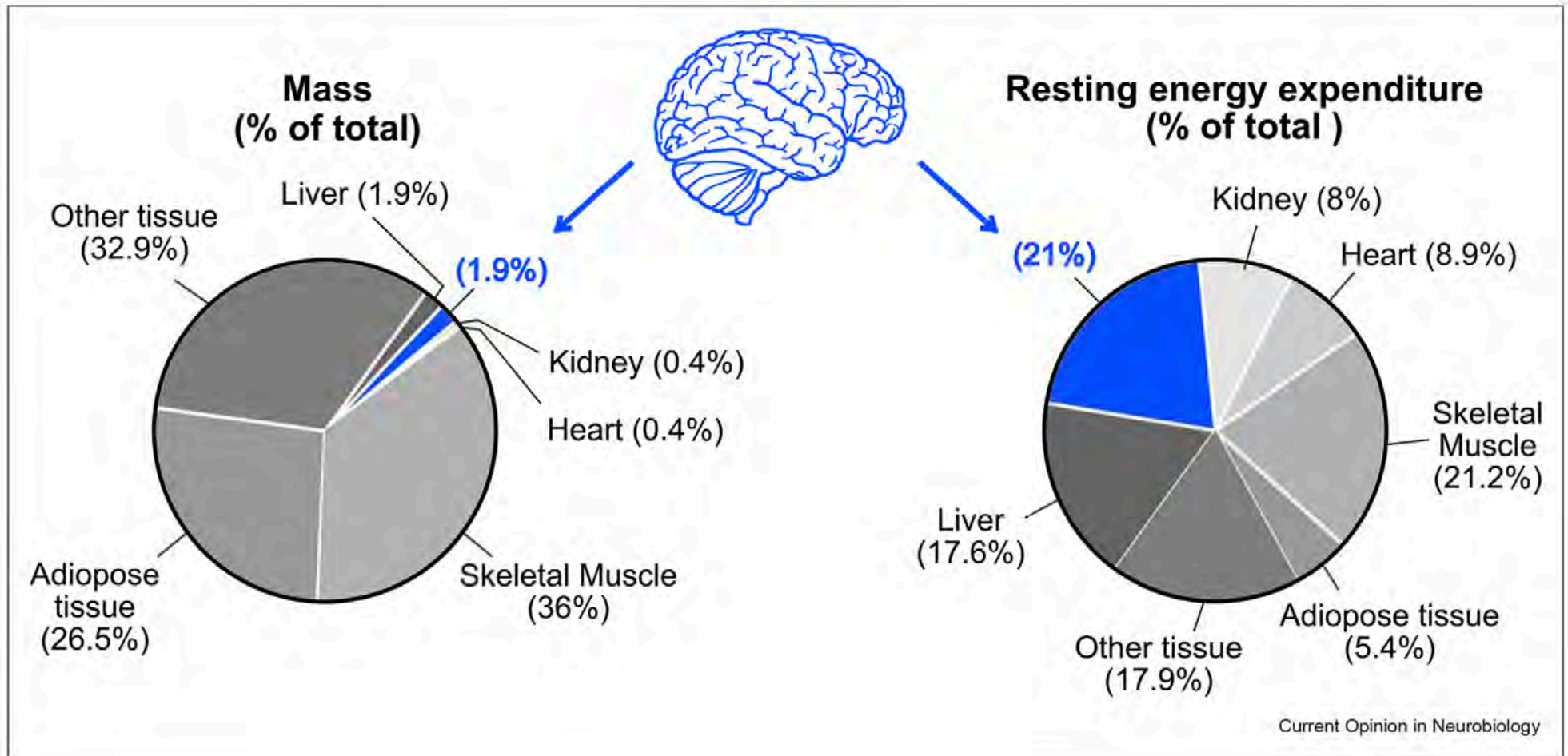
Brain mitochondrial diversity and network organization predict anxiety-like behavior in male mice



How are mitochondria distributed, and do they specialize across the *human* brain?



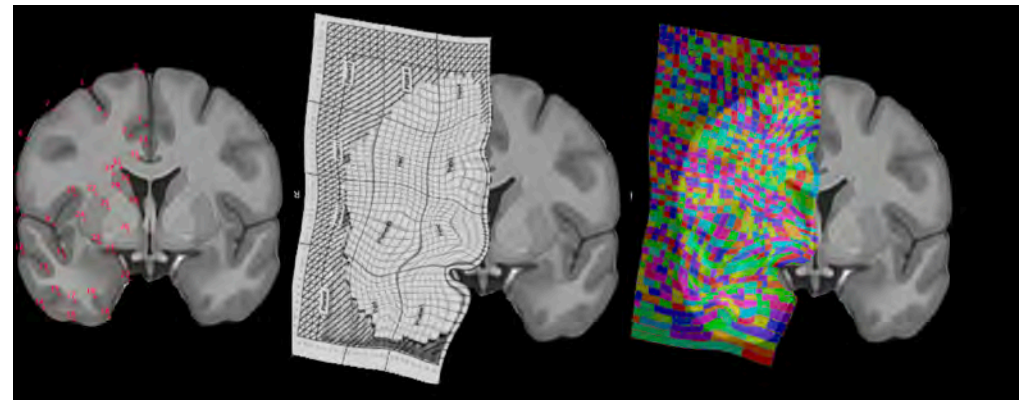
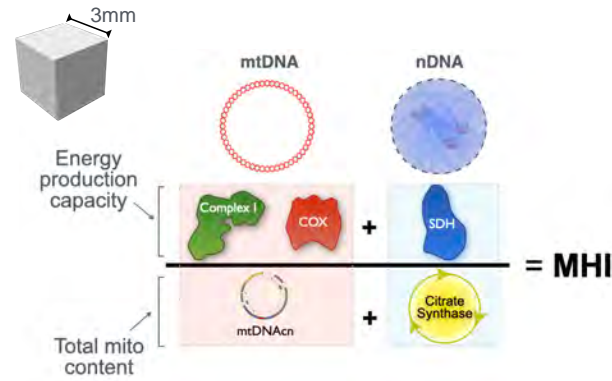
The brain's enormous, constant energy demand



Variations from rest to activity is ~5%

MitoBrainMap v1.0

A multi-function mitochondrial atlas of a single human coronal brain section at fMRI resolution



Closing the gap between organellar bioenergetic profiling and whole-brain neuroimaging modalities (fMRI, PET, CBV, DWI, etc)

Eugene Mosharov



a Right hemisphere slab at MNI -15.51



Anterior surface of the frozen brain slab



Cleaning (-1 mm)



Milling

d Collection



e Collection (-3 mm)



f Cleaning (-0.3 mm)





Cleaning (-1 mm)



Milling



e



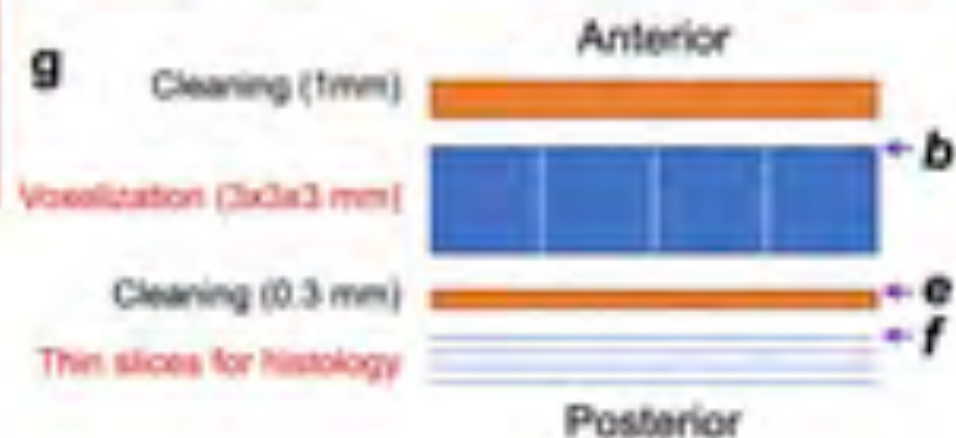
Collection (-3 mm)

f



Cleaning (-0.3 mm)

g



a Right hemisphere slab at MNI -15.51



Anterior surface of the frozen brain slab



Cleaning (-1 mm)



Milling

d Collection



Collection (-3 mm)



Cleaning (-0.3 mm)





Cleaning (-1 mm)



Milling



Collection (-3 mm)

e

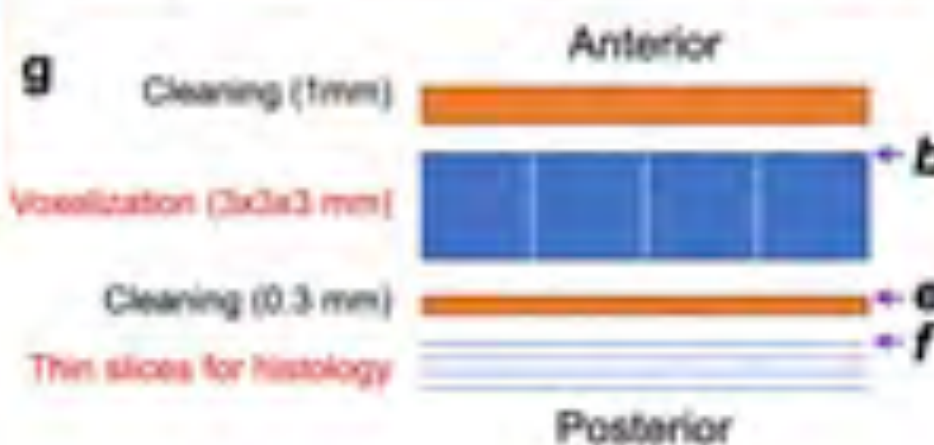


Cleaning (-0.3 mm)

f

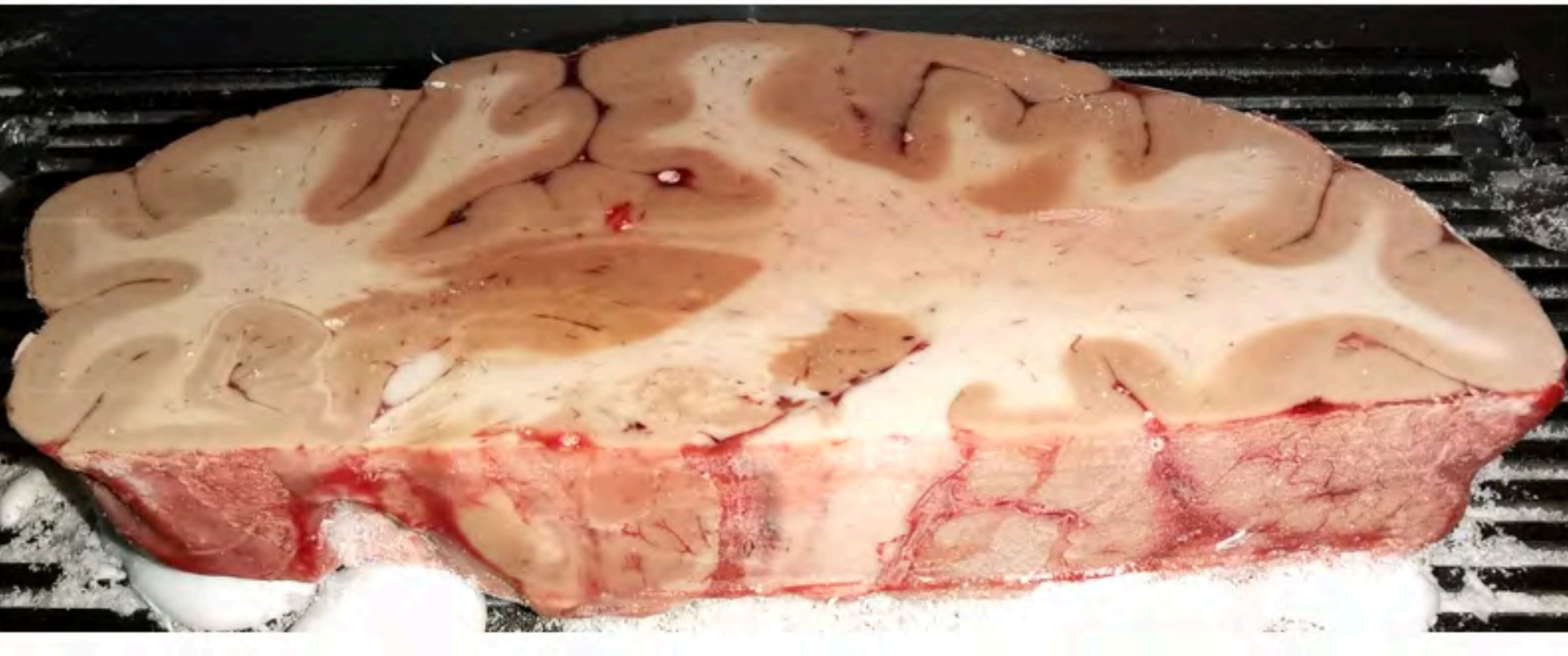


g



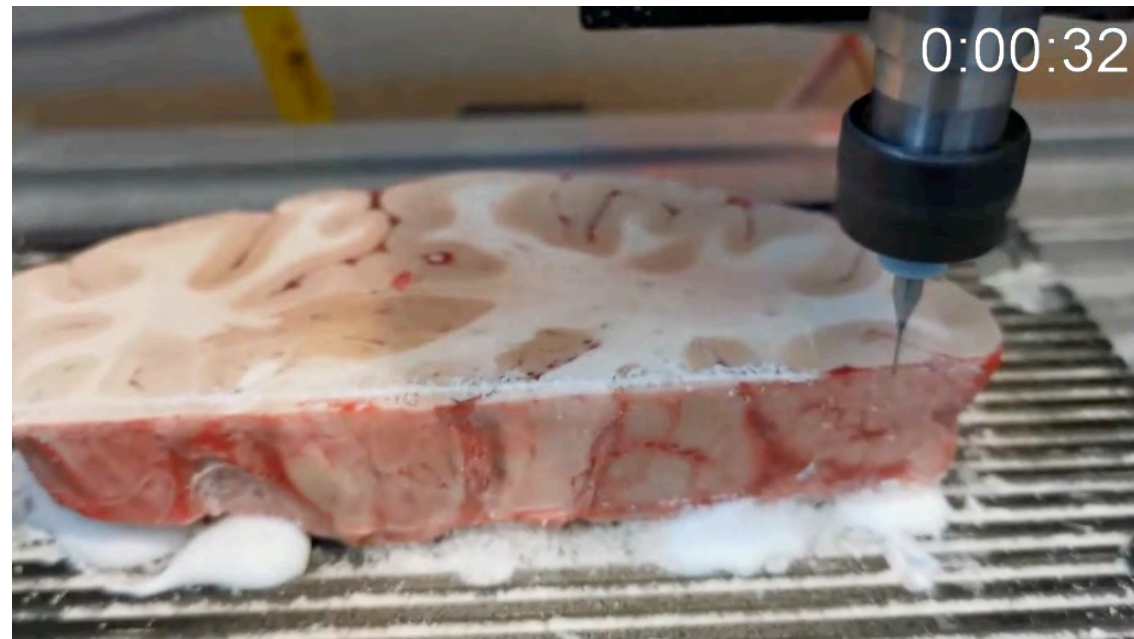
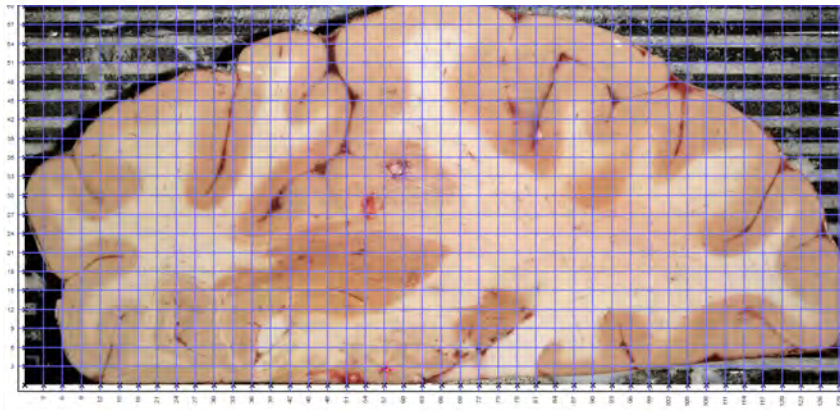
MitoBrainMap v1.0

A multi-function mitochondrial atlas of a single human coronal brain section at fMRI resolution

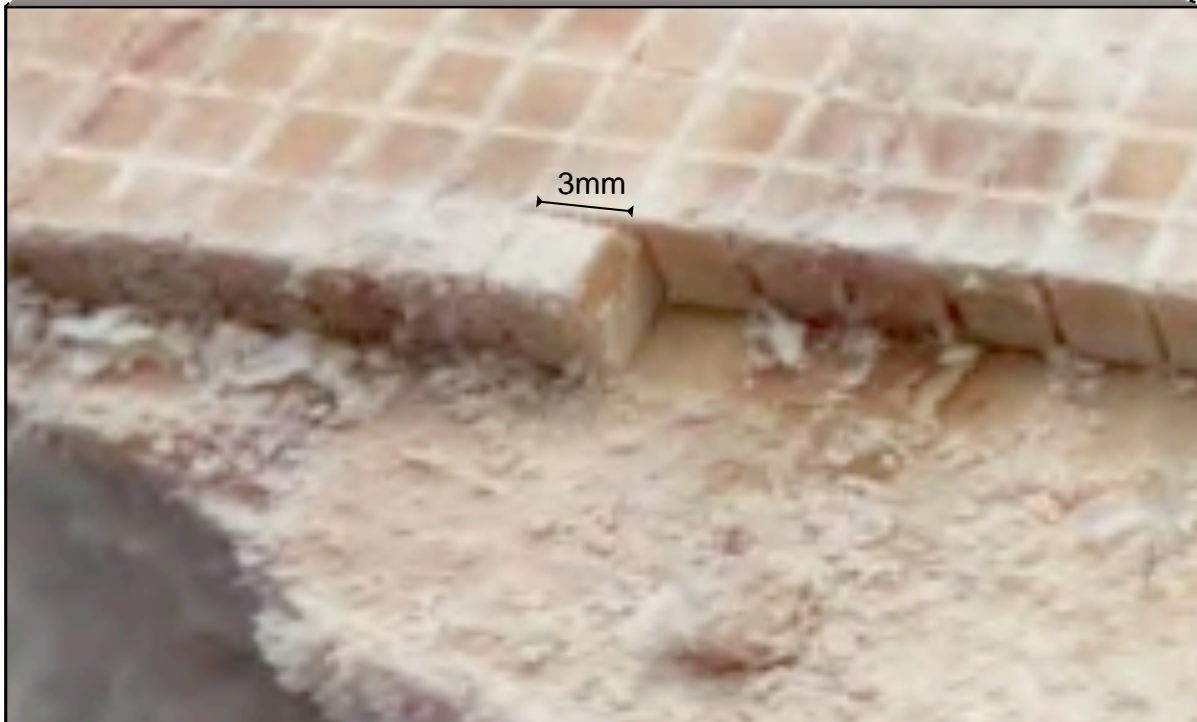
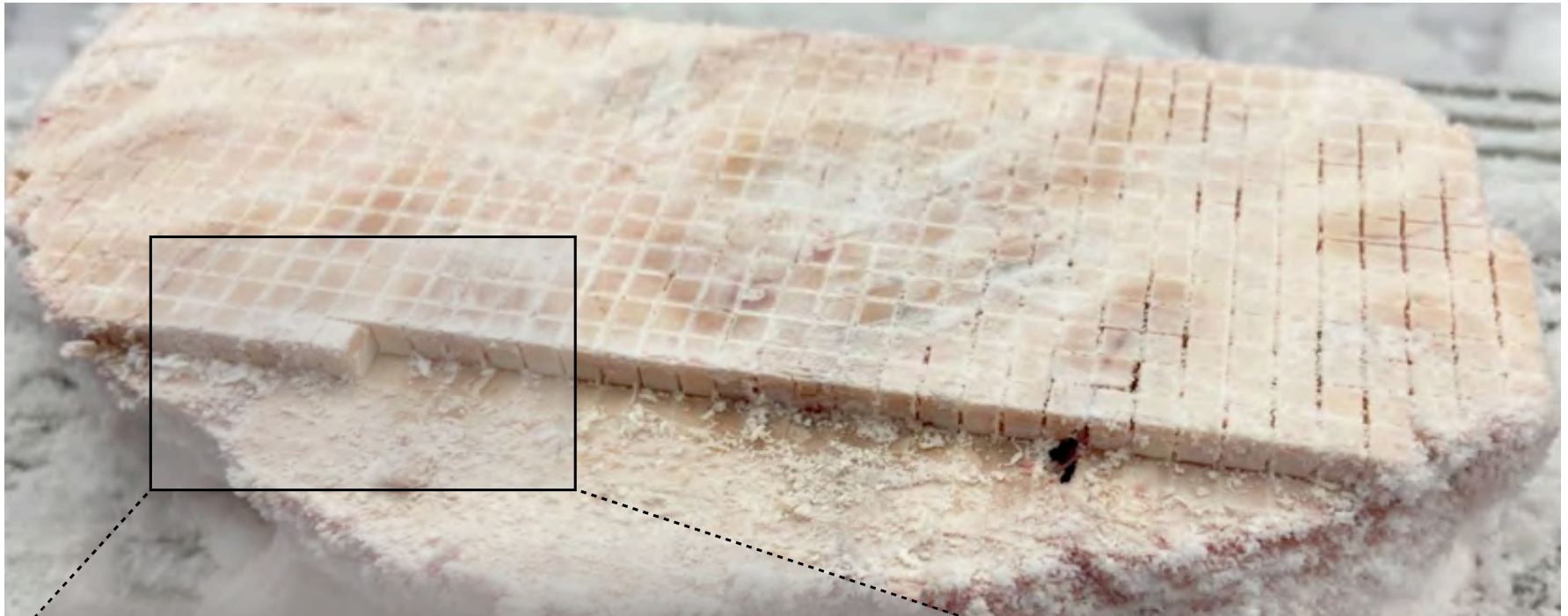


MitoBrainMap v1.0

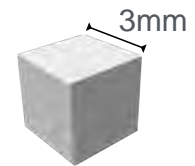
A multi-function mitochondrial atlas of a single human coronal brain section at fMRI resolution



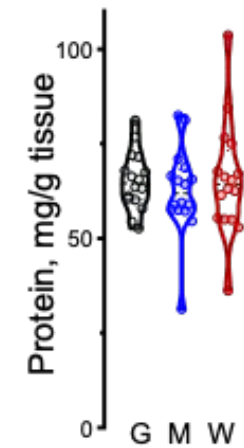
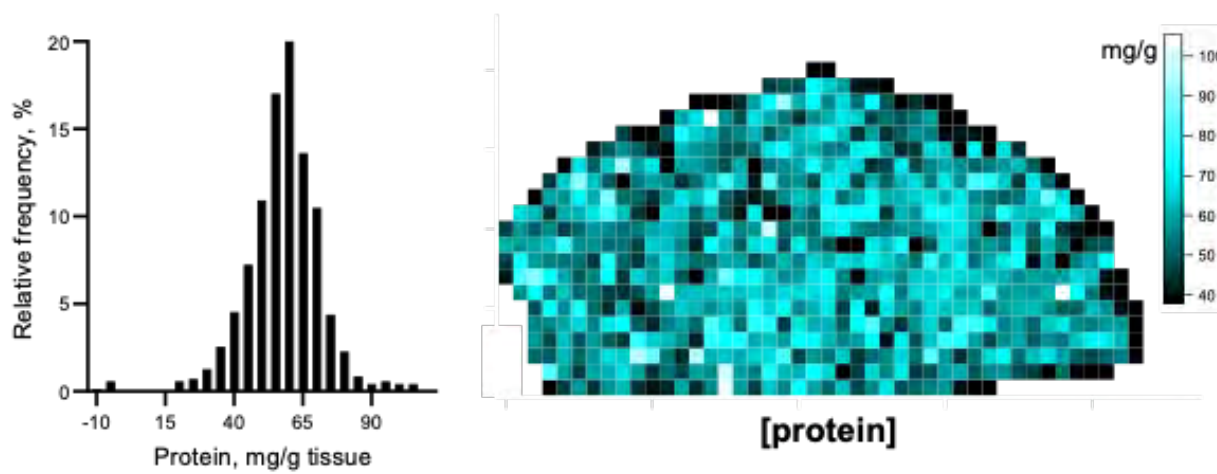
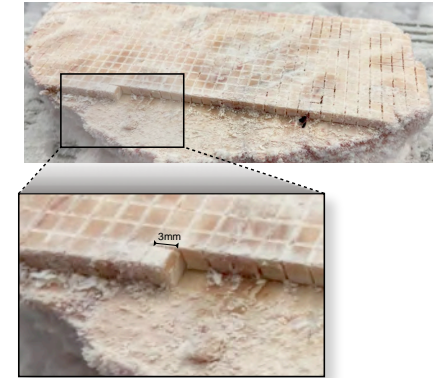
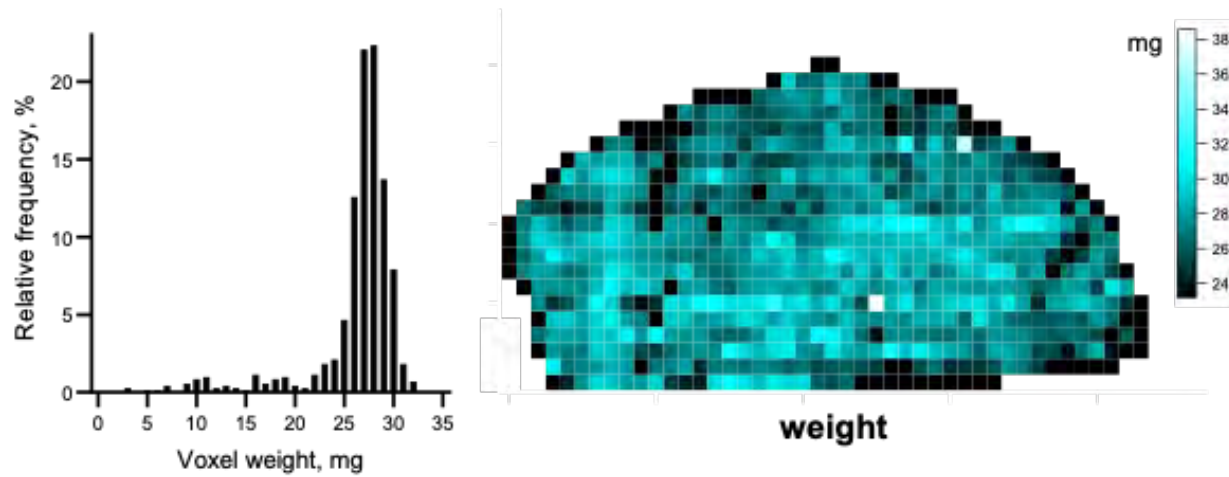
Eugene Mosharov

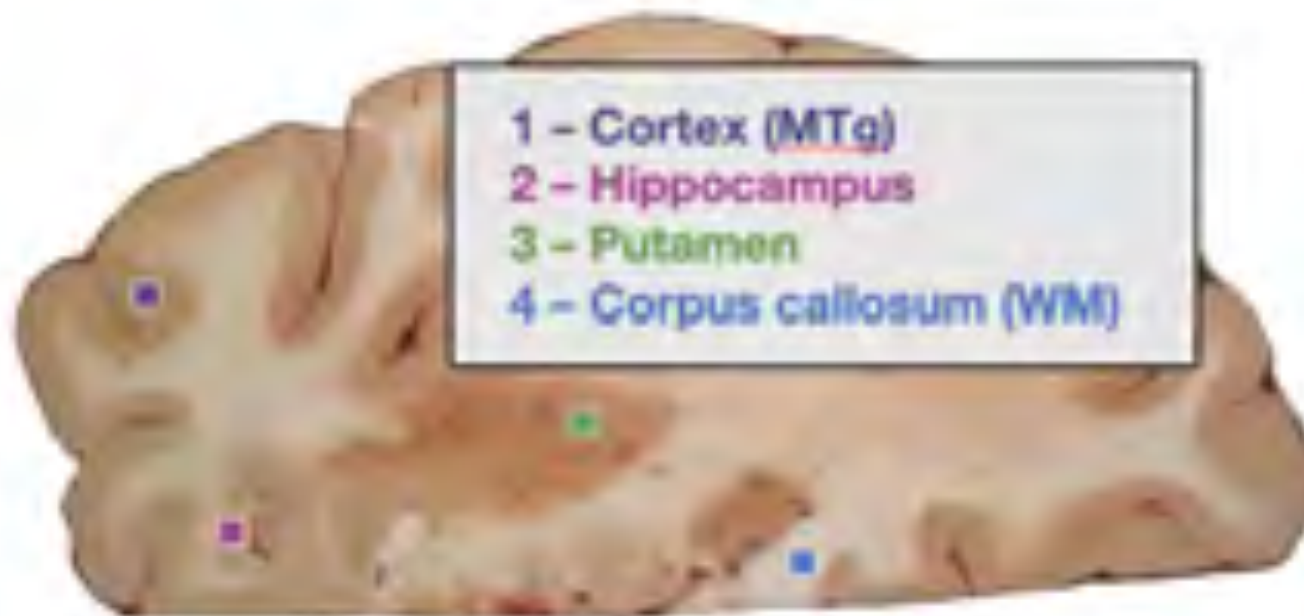


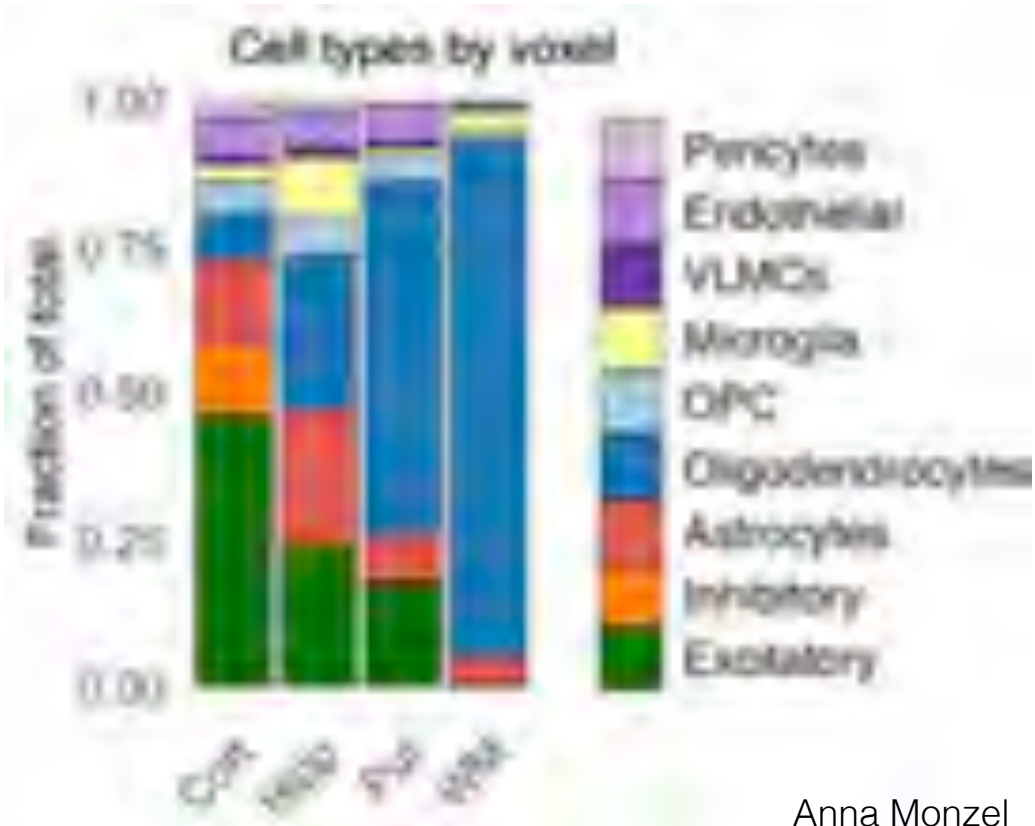
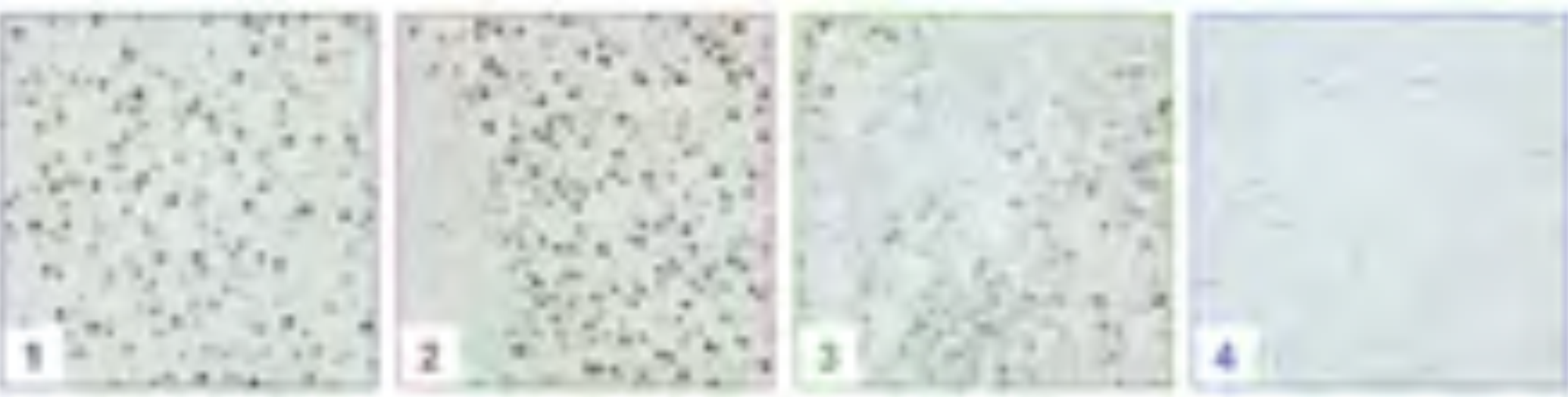
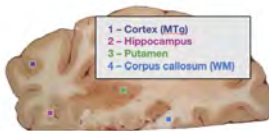
Physical *voxelization*
of the human brain
at fMRI resolution

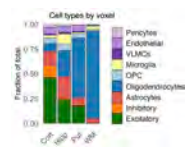
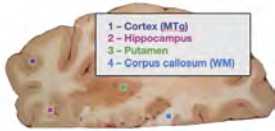
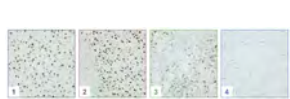


Quality control on 702 human brain voxels

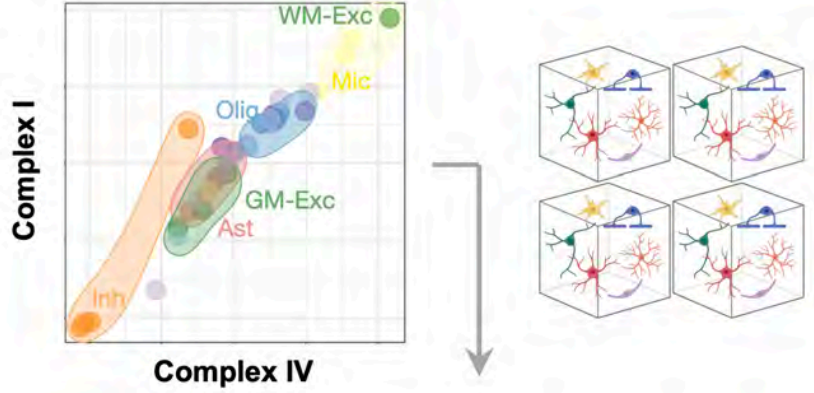




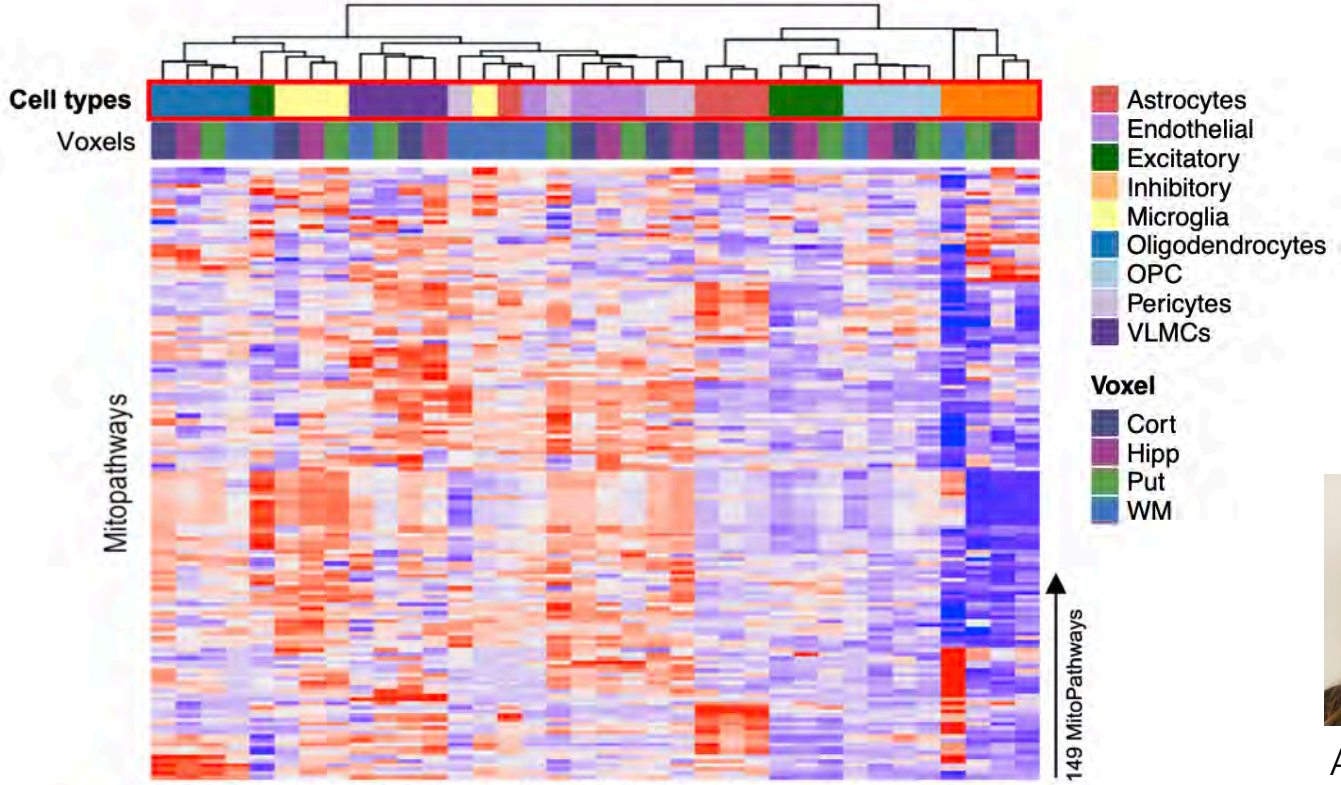




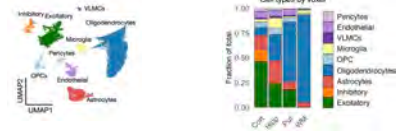
Normalized by voxel



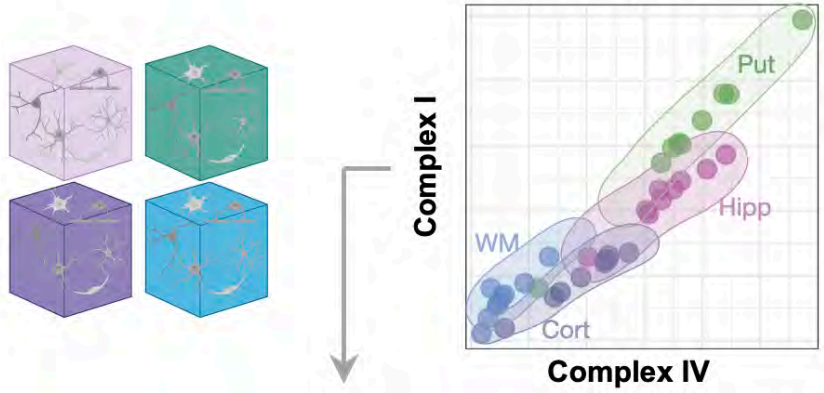
Cell type-specific mitotypes



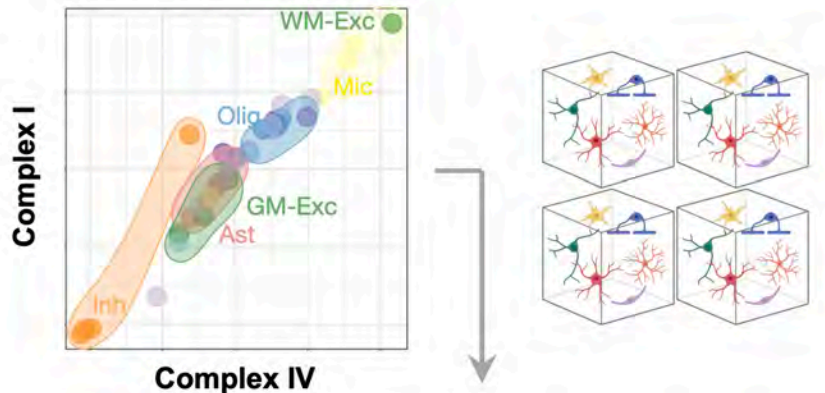
Anna Monzel



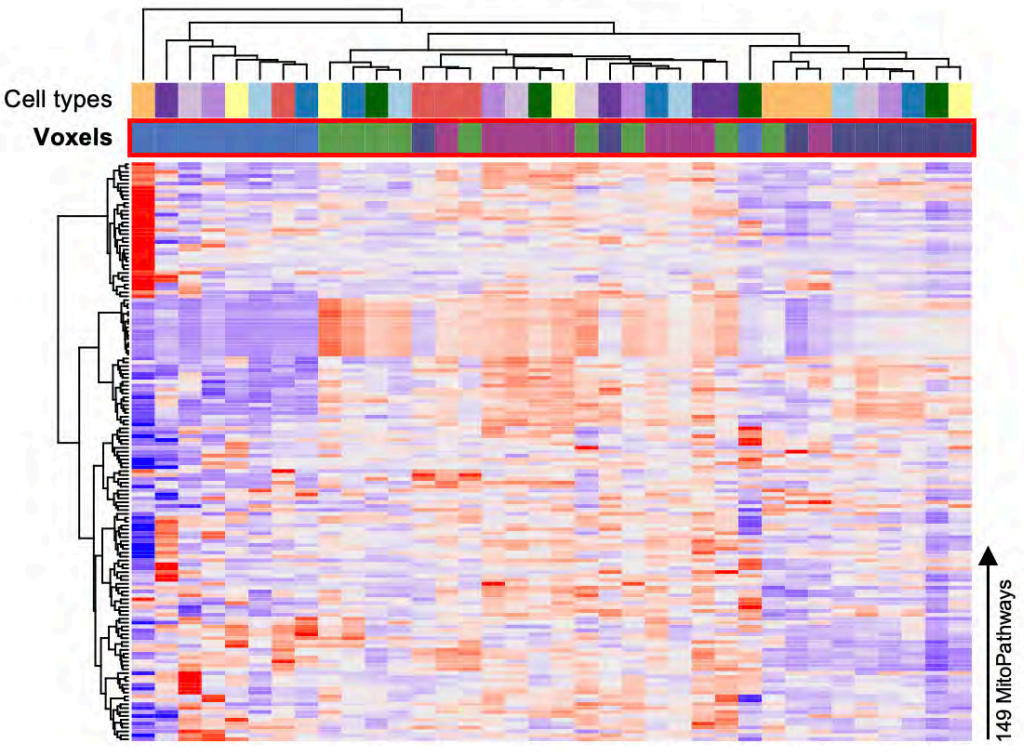
Raw expression



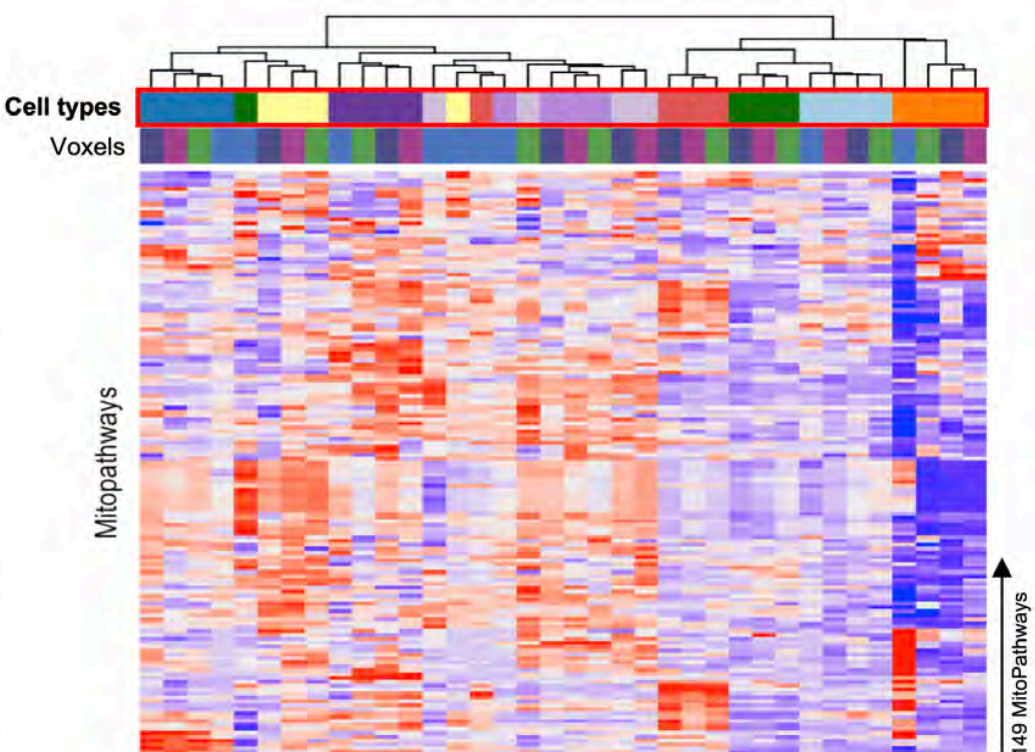
Normalized by voxel



Voxels-specific mitotypes



Cell type-specific mitotypes

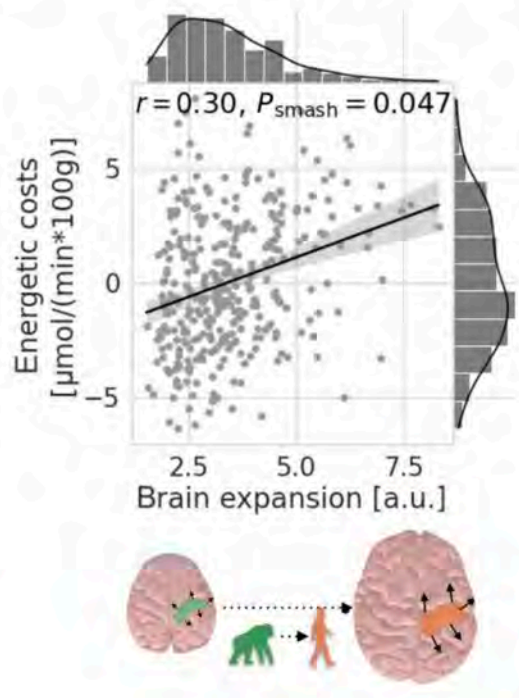


NEUROSCIENCE

An energy costly architecture of neuromodulators for human brain evolution and cognition

Gabriel Castrillon^{1,2,3}, Samira Epp^{1,4}, Antonia Bose^{1,4}, Laura Fraticelli^{1,4}, André Hechler^{1,4}, Roman Belenya^{1,4}, Andreas Ranft⁵, Igor Yakushev⁶, Lukas Utz¹, Lalith Sundar⁷, Josef P Rauschecker^{8,9}, Christine Preibisch^{1,10}, Katarzyna Kurcyus¹, Valentin Riedl^{1,3*}

B Human brain expansion

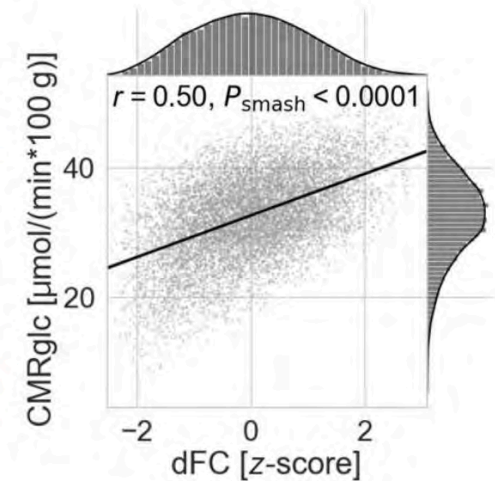
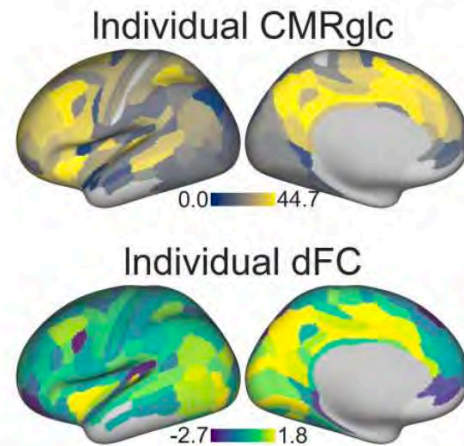


PET / MRI



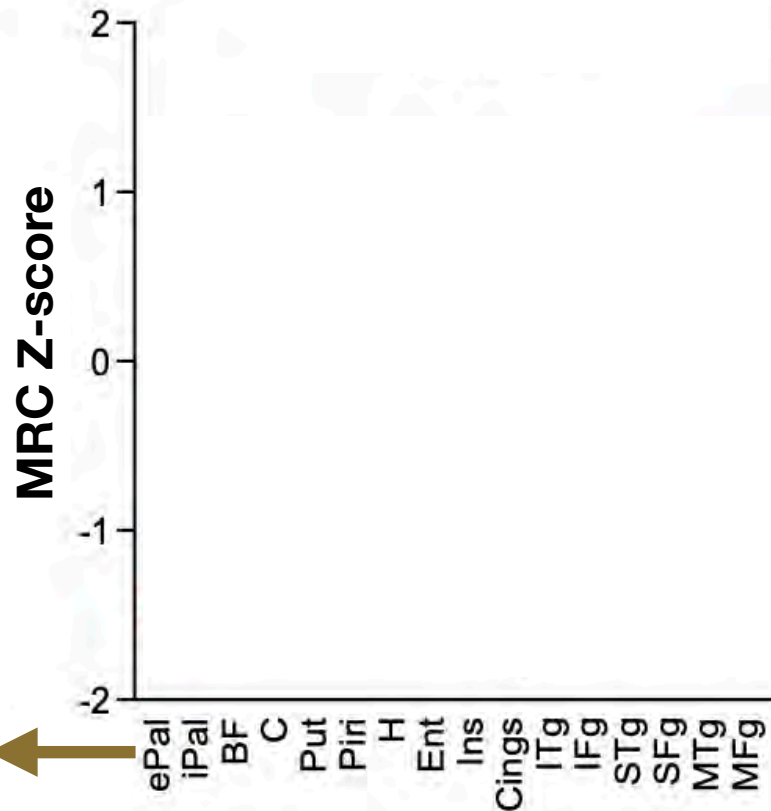
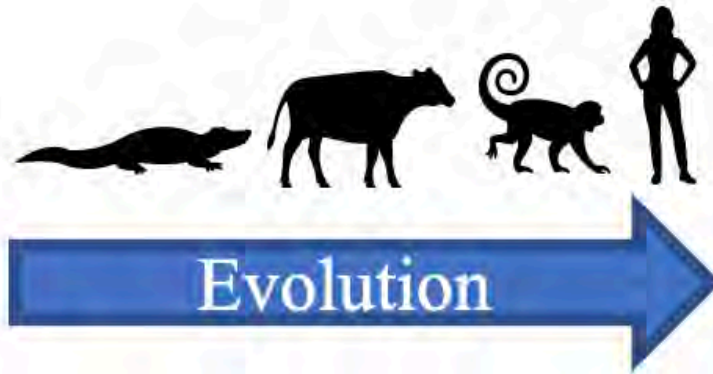
Arterial blood sampling

Subject space



G-coupled protein receptor expression correlates with glucose consumption

Evolutionary correlate of mitochondrial OxPhos specialization



Most "ancient" ←

→ Most "recent"



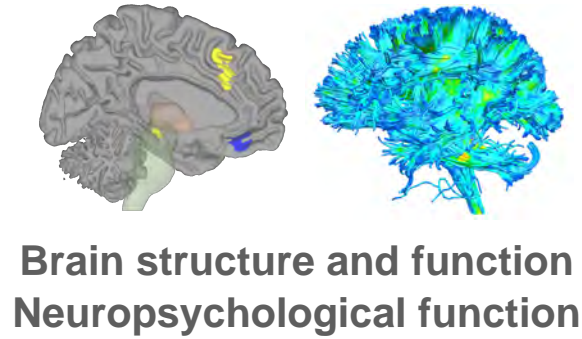
**MITO
BRAIN**

ROSMAP

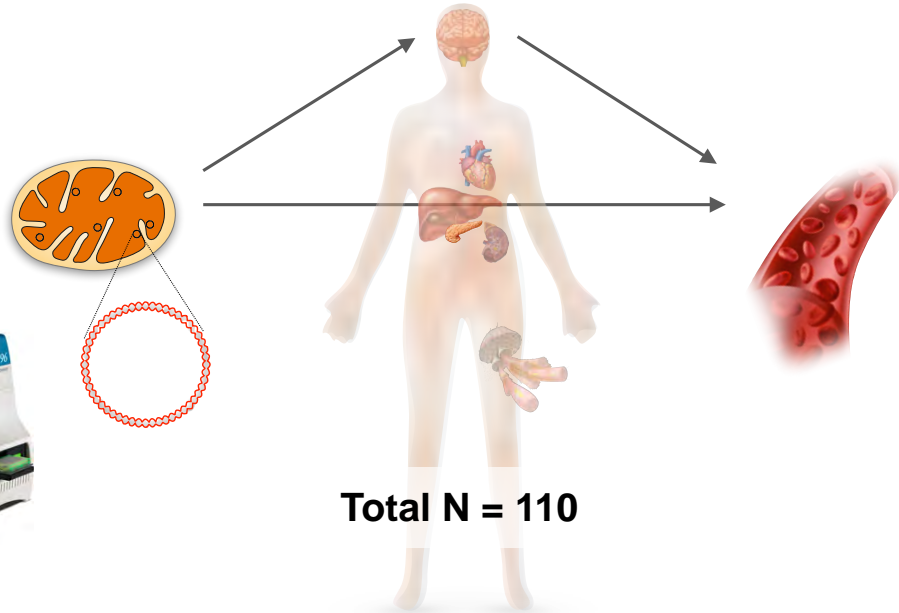
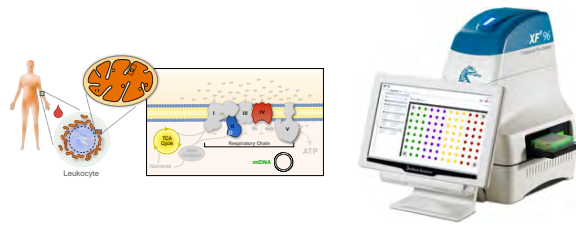


Cynthia Liu

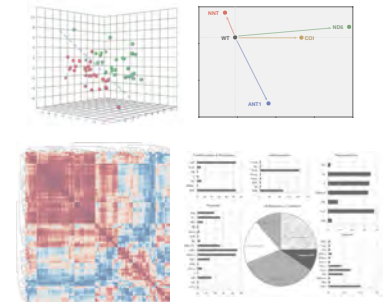
Mitochondrial Stress, Brain Imaging, and Epigenetics — MiSBIE



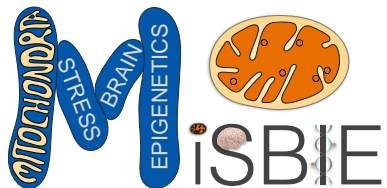
mtDNA heteroplasmy
Mitochondrial OxPhos
Lymphocytes, Monocytes,
Neutrophils, Platelets

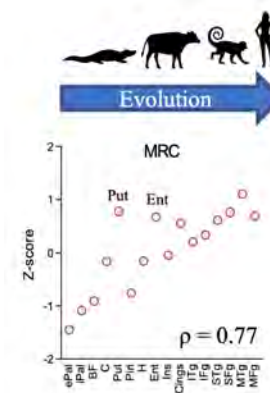
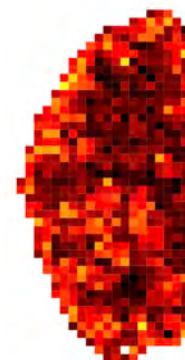
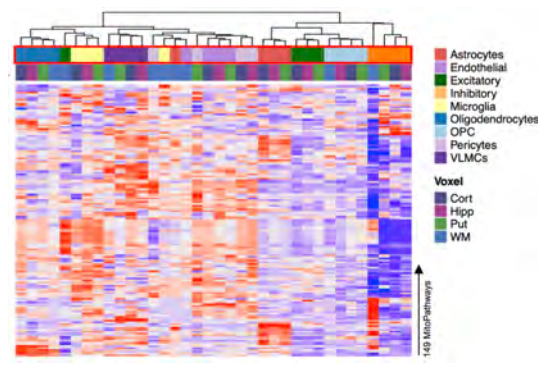
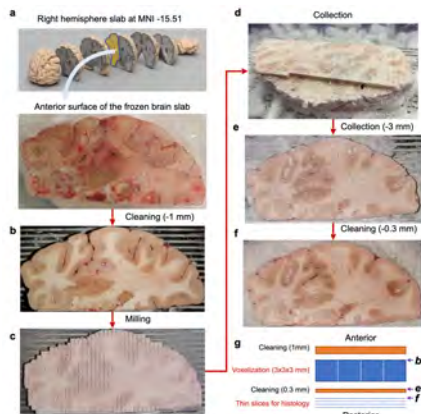
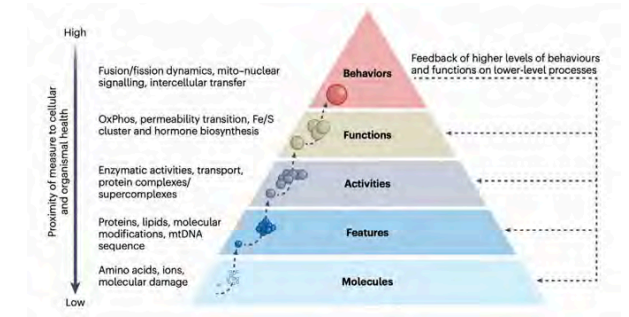
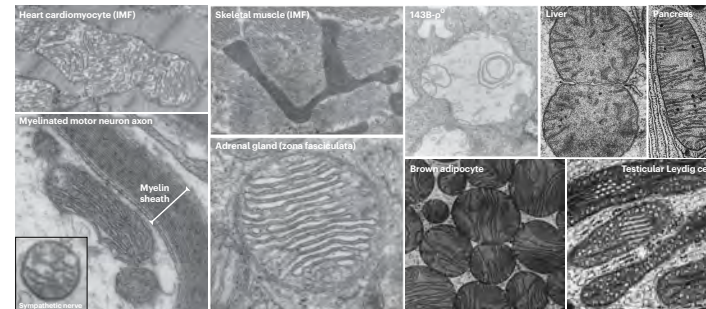
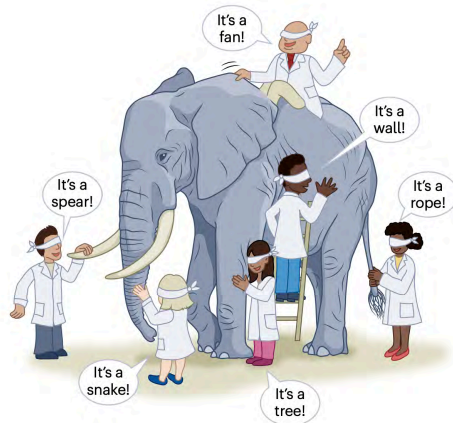
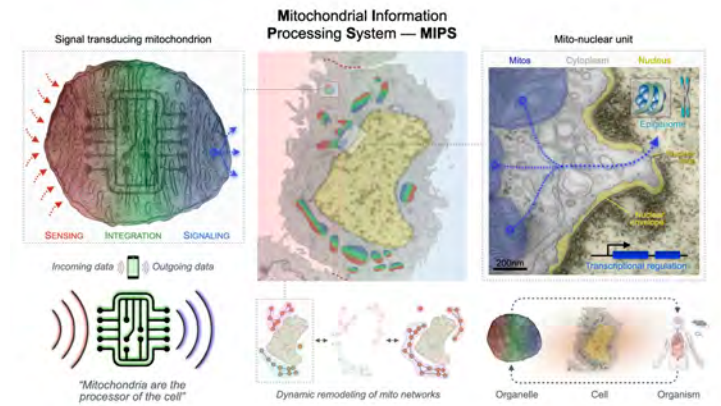
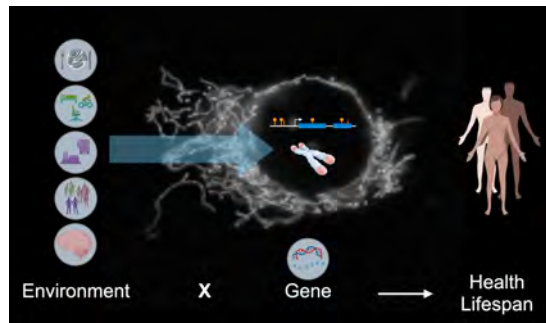
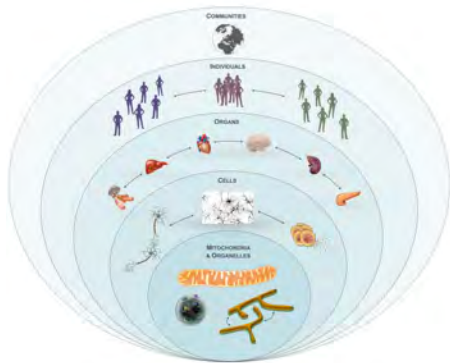


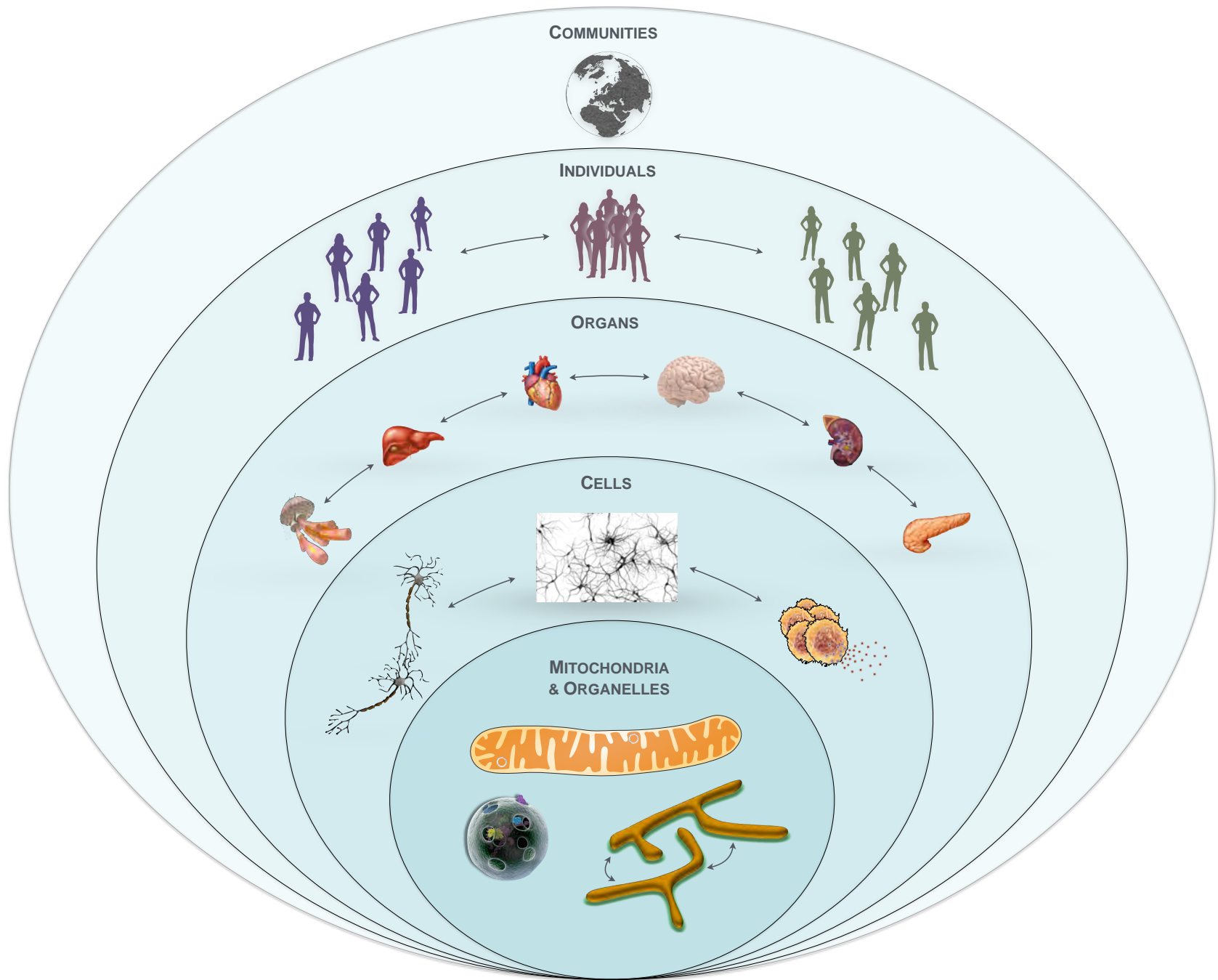
Disease biomarkers
Stress reactivity
Energy expenditure
e.g., GDF15, Lactate,
Pyruvate, Alanine, etc.



- **Healthy controls** (n = 70)
- **mtDNA defects**
 - 3243A>G (group A) (n = 20)
 - 3243A>G (group B) (n = 5)
 - Single deletion** (n = 15)



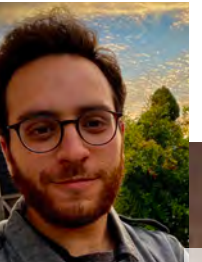




Mitochondrial PsychoBiology Lab

Linking molecular processes within mitochondria with the human experience

OUR RESEARCH



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Jack

Anna

Cynthia



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BASZUCKI
BRAIN RESEARCH FUND

The Nathaniel Wharton Fund 

 National Institute of Mental Health

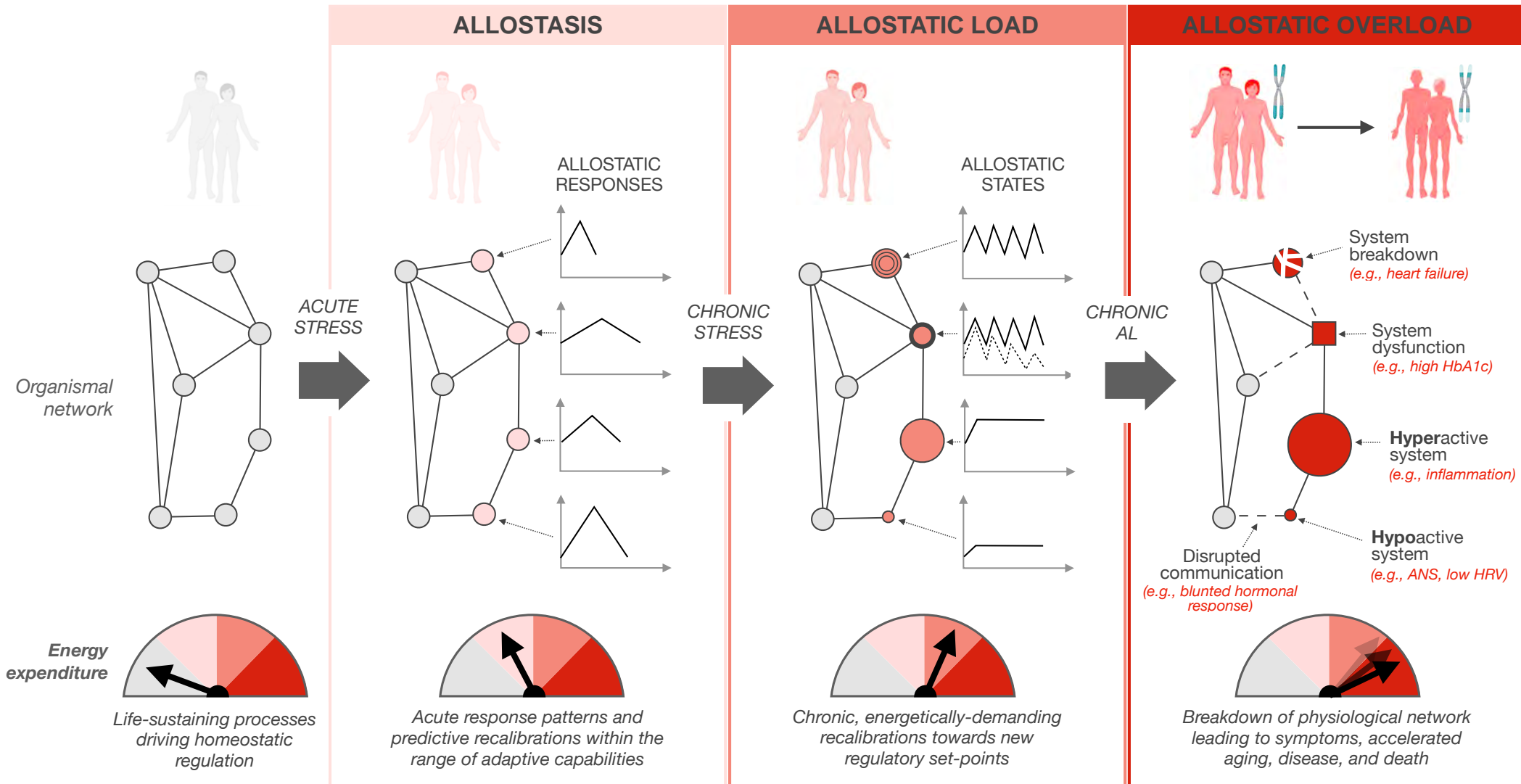
 National Institute of General Medical Sciences

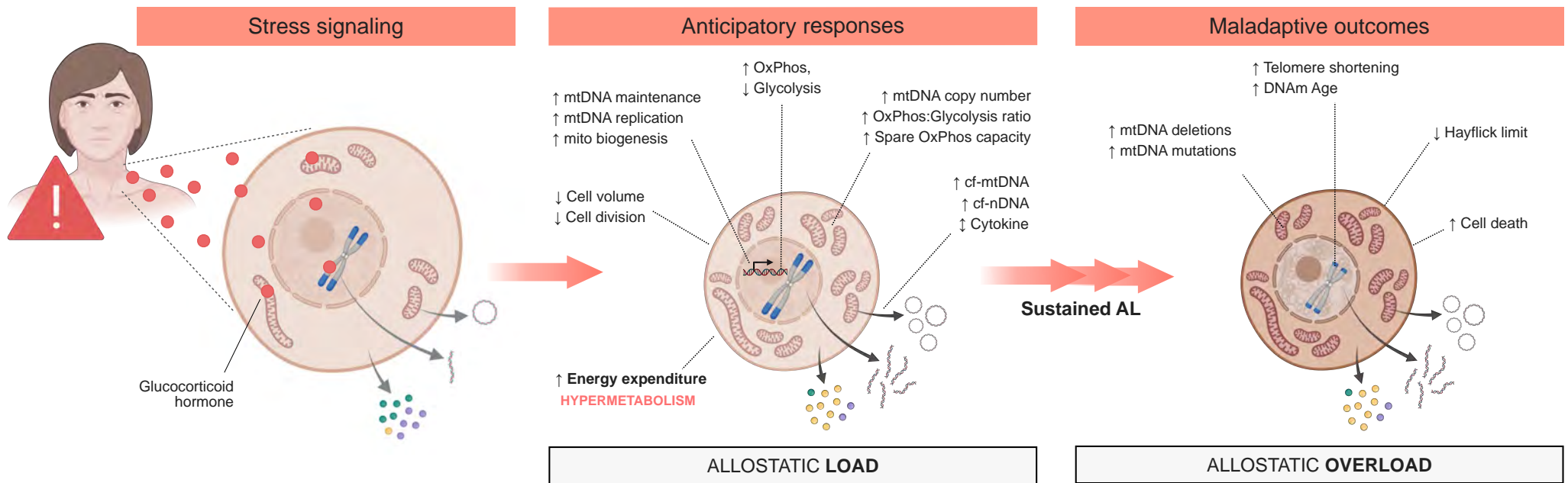
 National Institute on Aging



Downloadable
presentation slides

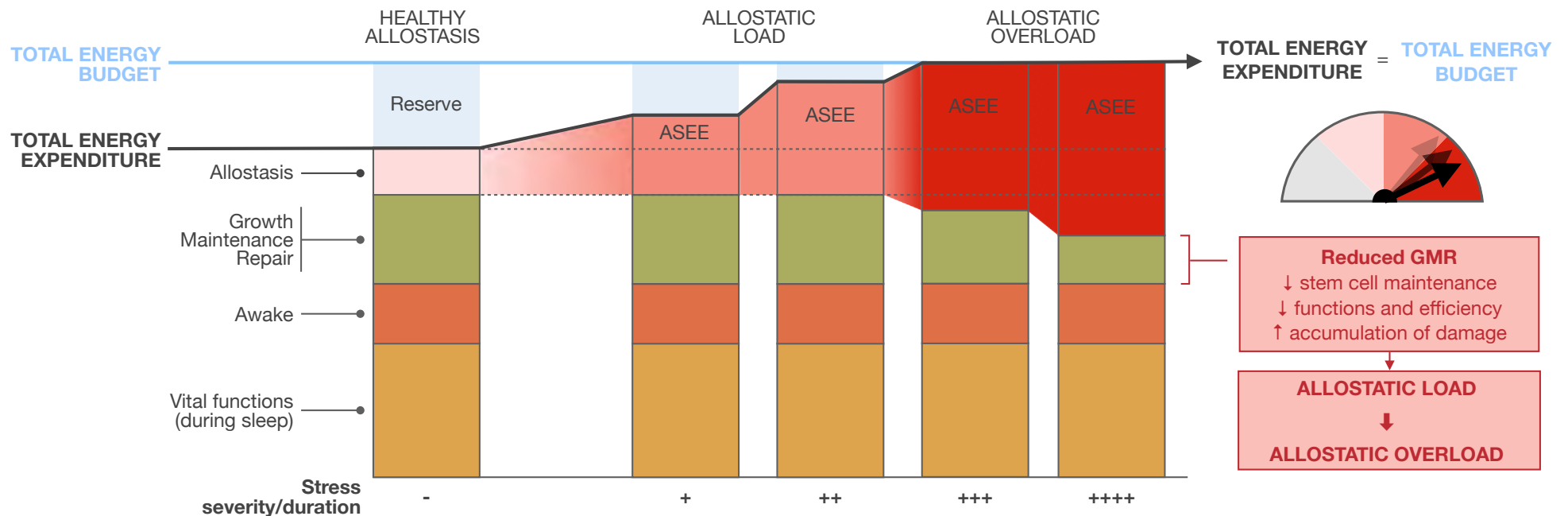
Energetic Model of Allostatic Load (EMAL)





Glucocorticoid signaling increases energy expenditure by **60%**
 And accelerates cellular aging by **10-40%**

Model: Aging and lifespan determined by the partitioning of limited energetic resources



Brain-body Energy Conservation (BEC)

