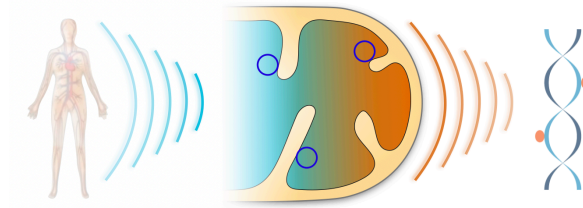


Energetic regulation of aging: What is the role of the brain?



Yale — Biology of Aging

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 OF
OPPORTUNITY. | **New York State
Psychiatric Institute**

PART 1

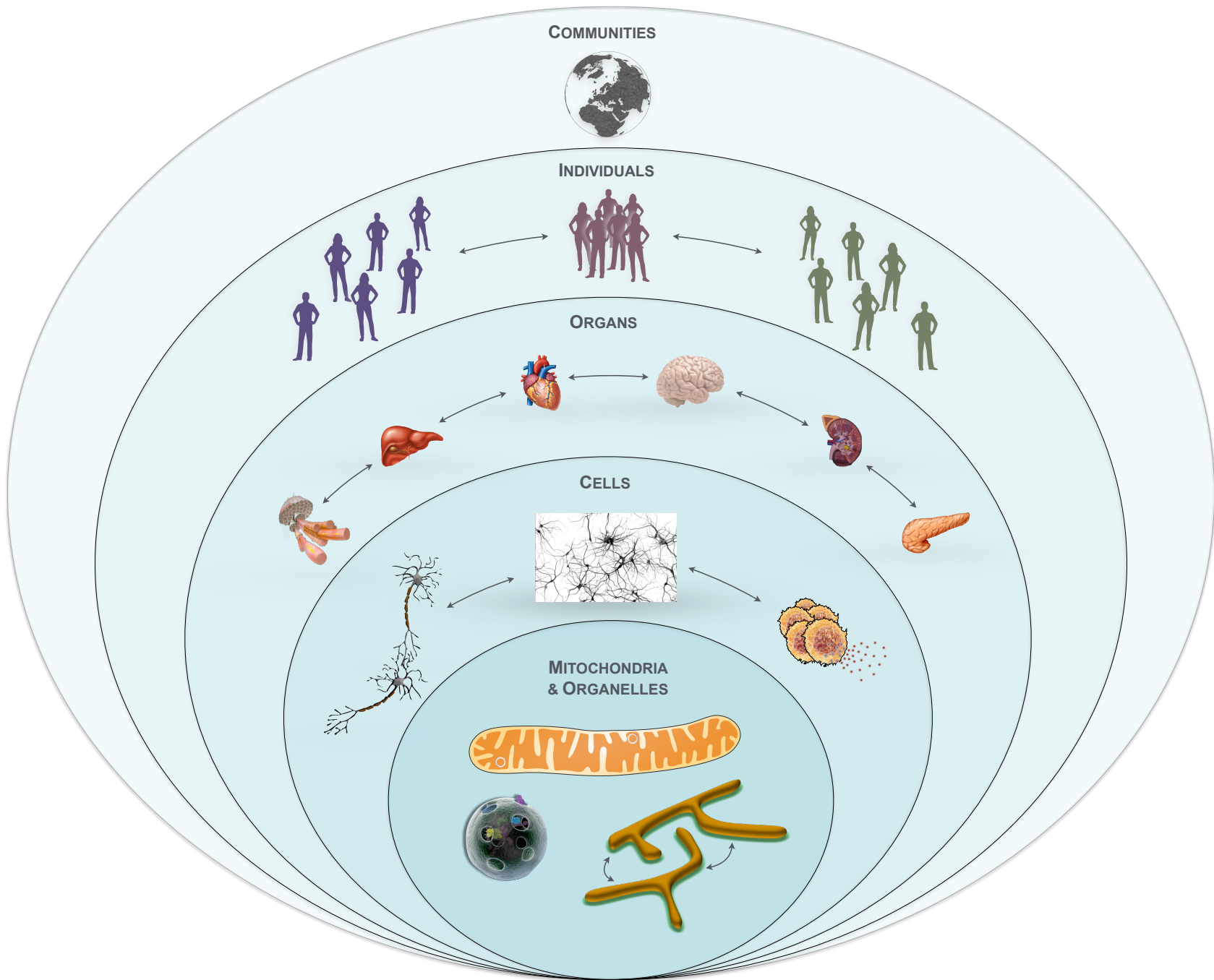
Mitochondrial Signal Transduction, Hair greying

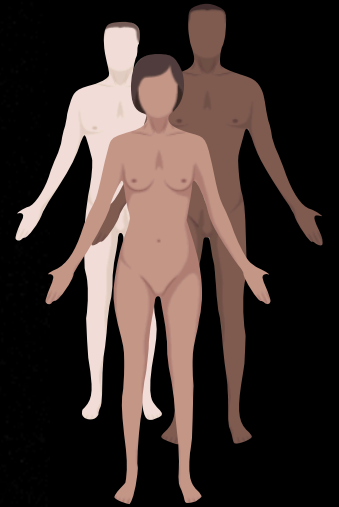
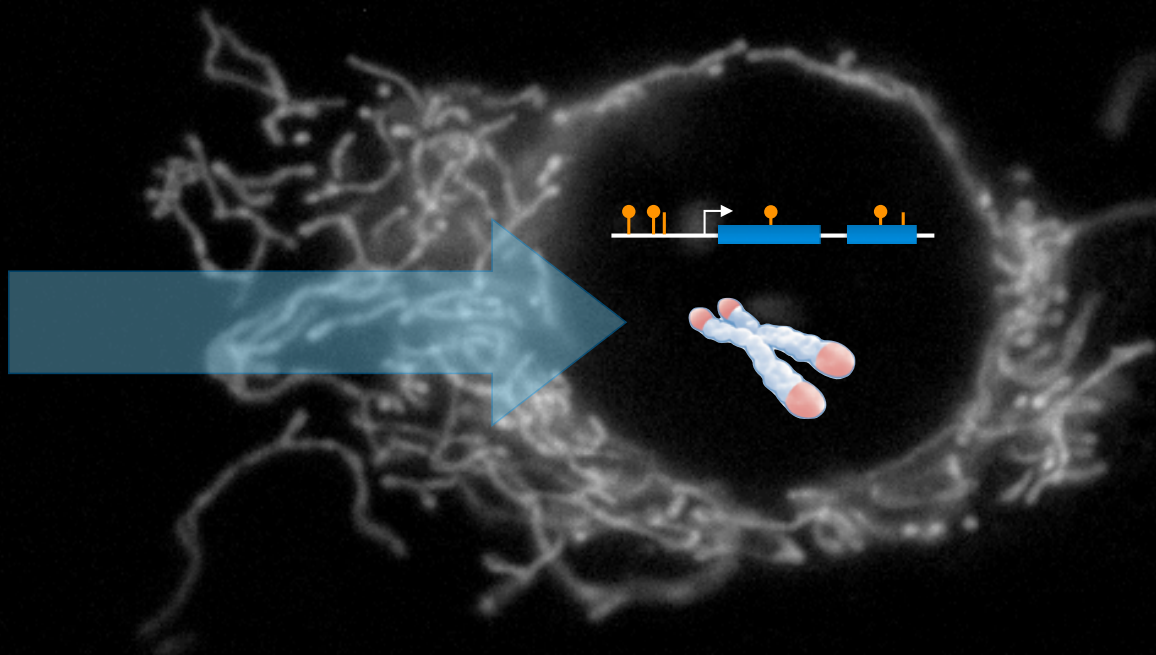
PART 2

Energy, OxPhos defects, Hypermetabolism, Rate of aging

PART 3

Somato-cognitive Energy Conservation — SEC





Environment

X

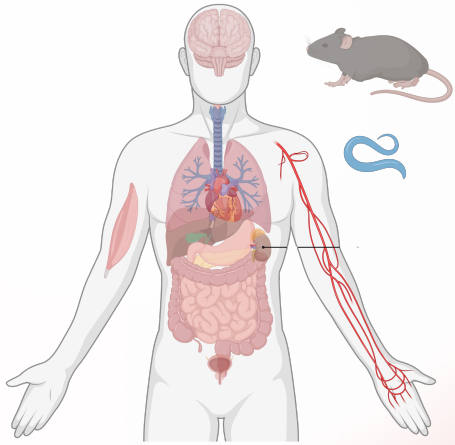
Gene



Health
Lifespan

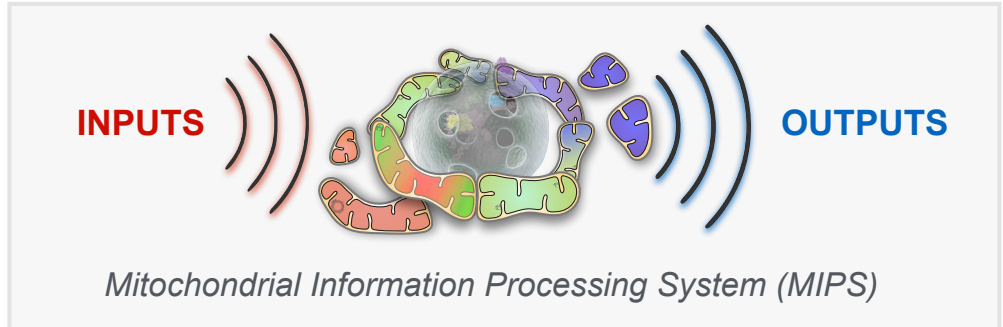
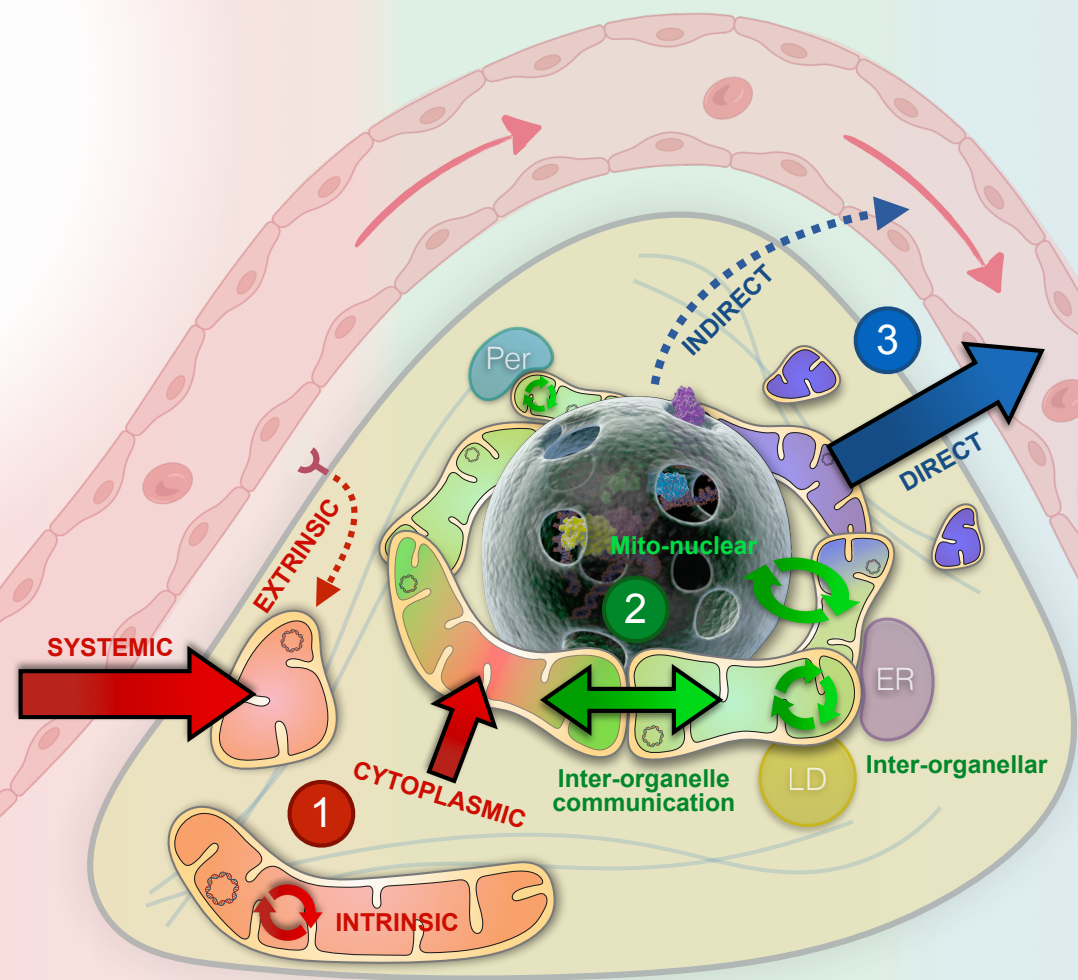
“Are mitochondria the X factor?”

Multicellular organisms

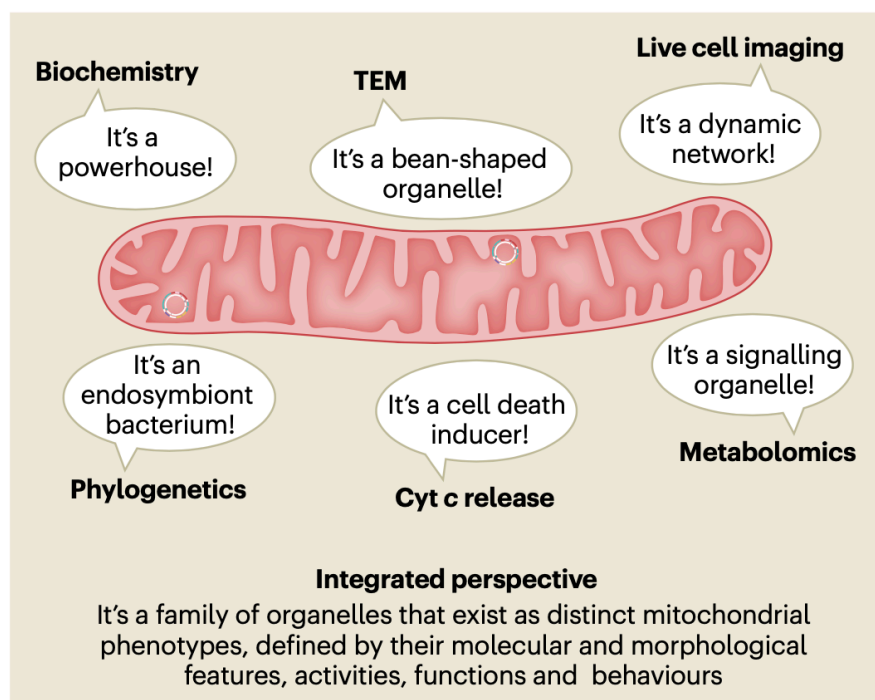
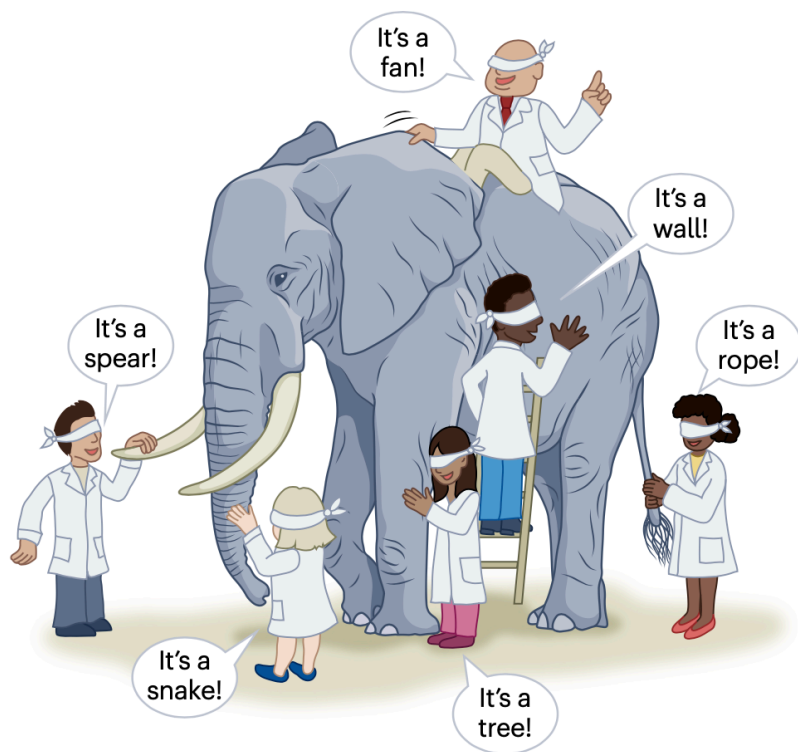


- INPUTS**
- Peptide hormones
 - Steroid & other hormones
 - Nutrients levels
 - Metabolites
 - Ions
 - Gases (e.g., O₂, NO)
 - ATP/ADP (ΔG_p)
 - NAD(P)⁺/NAD(P)H ratio
 - mtDNA variations
 - Others

- 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

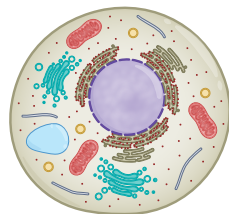
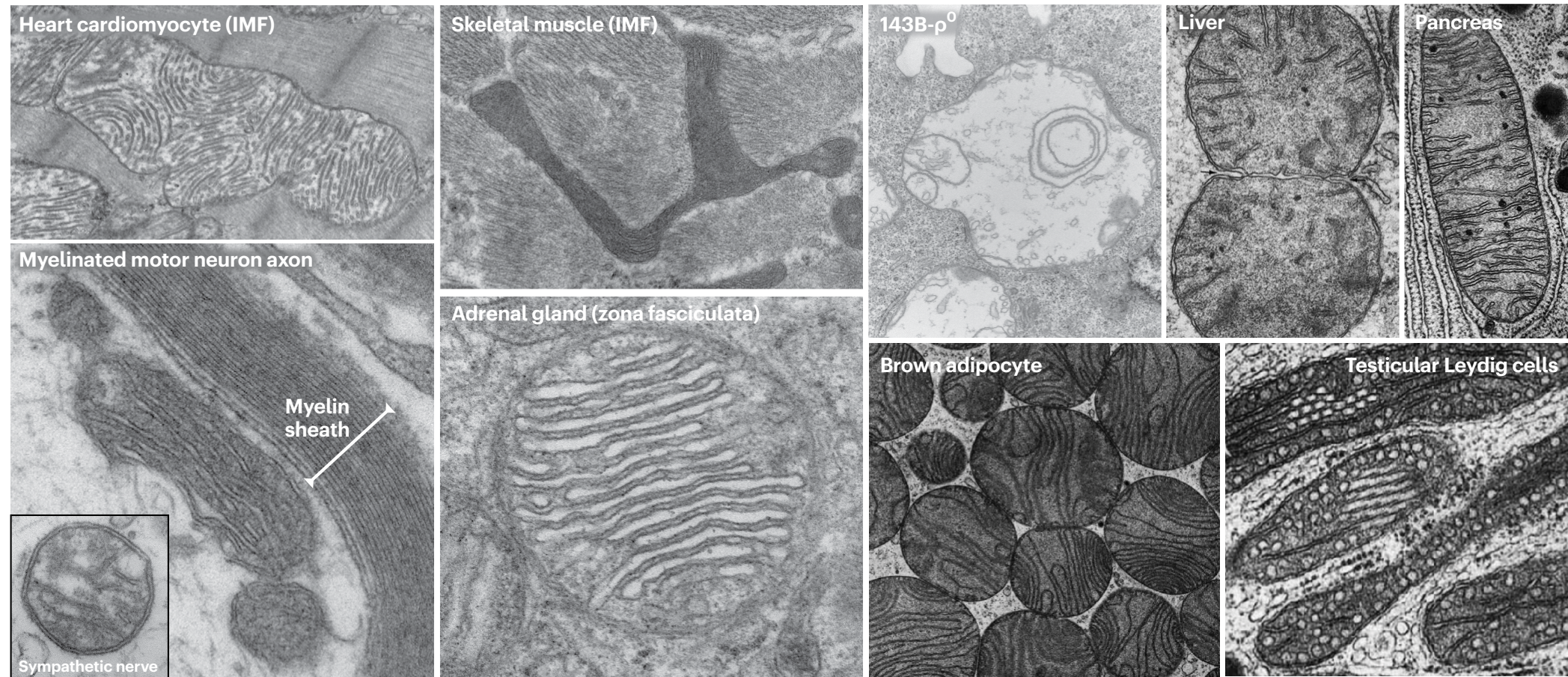


Multifaceted mitochondria: moving mitochondrial science beyond function and dysfunction

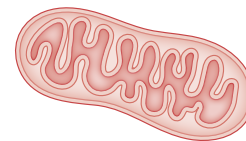


A catalogue of mitochondrial *functions*

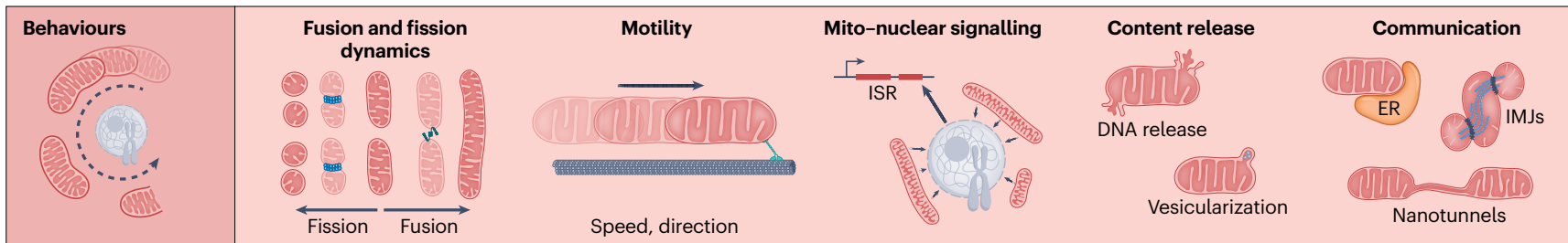
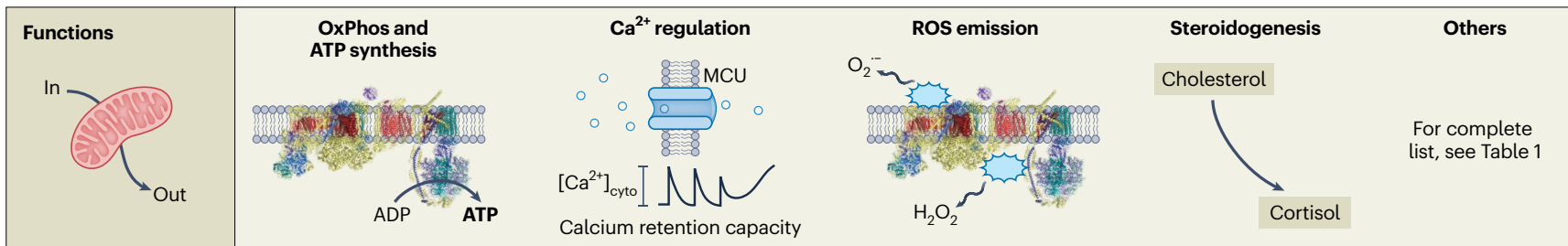
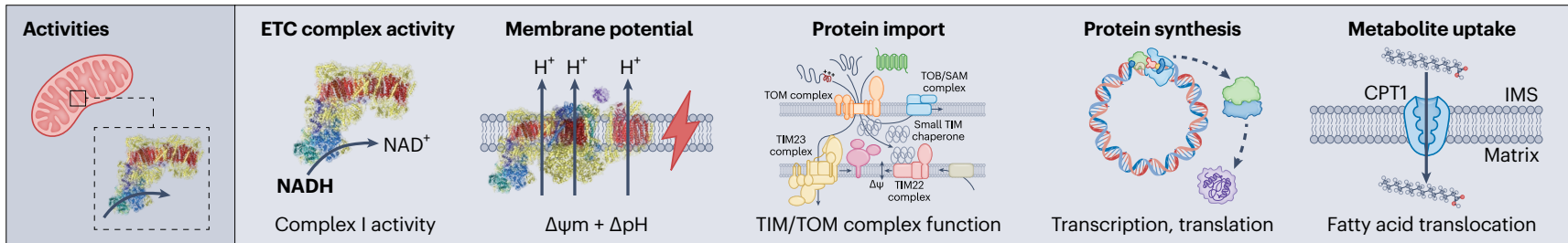
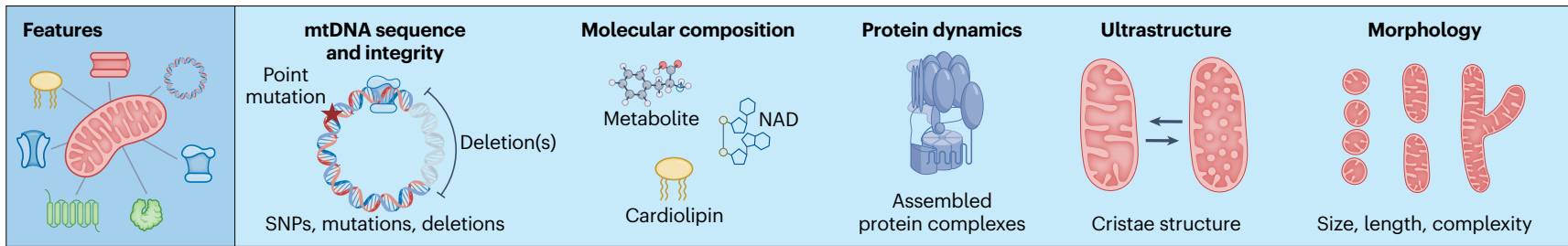
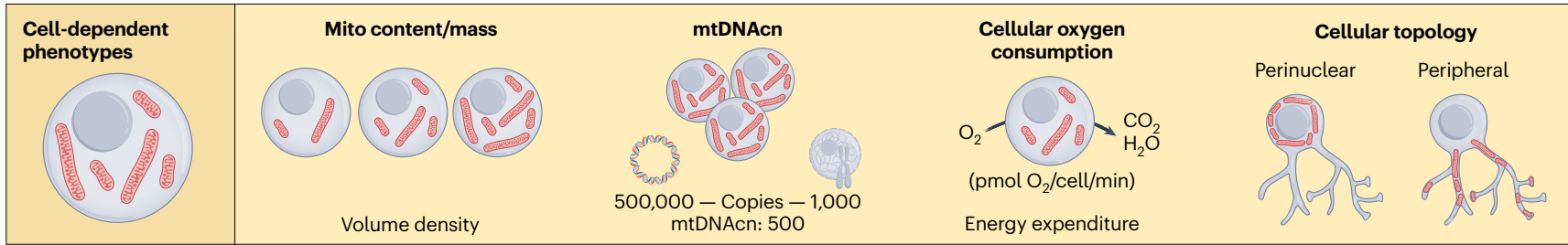
Different mitochondria types (mitotypes)



Cell types and subtypes

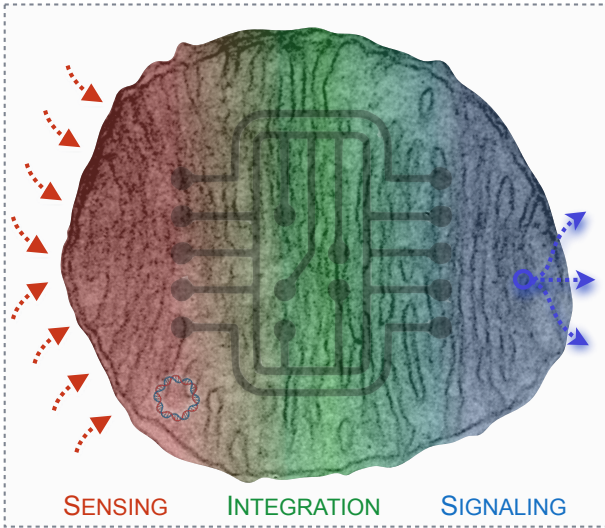


Mitochondrial phenotypes

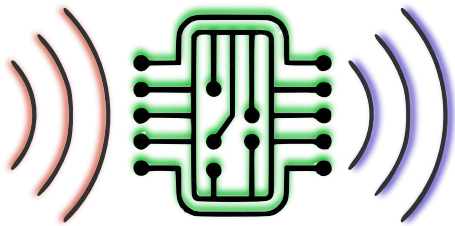


Mitochondrial Information Processing System — MIPS

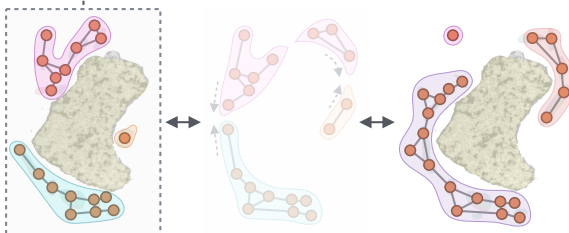
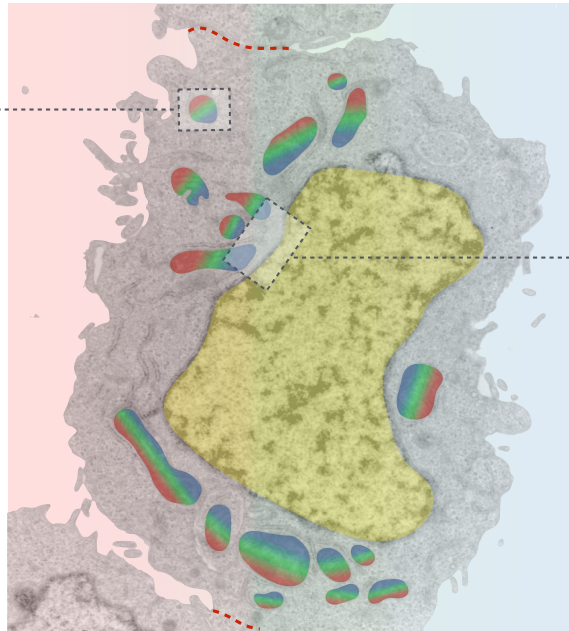
Signal transducing mitochondrion



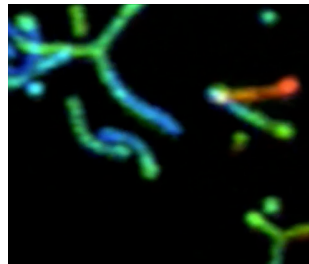
Incoming data  Outgoing data



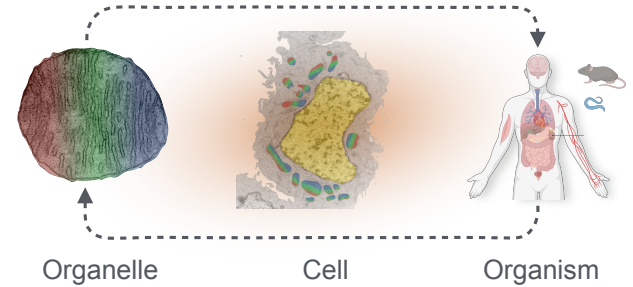
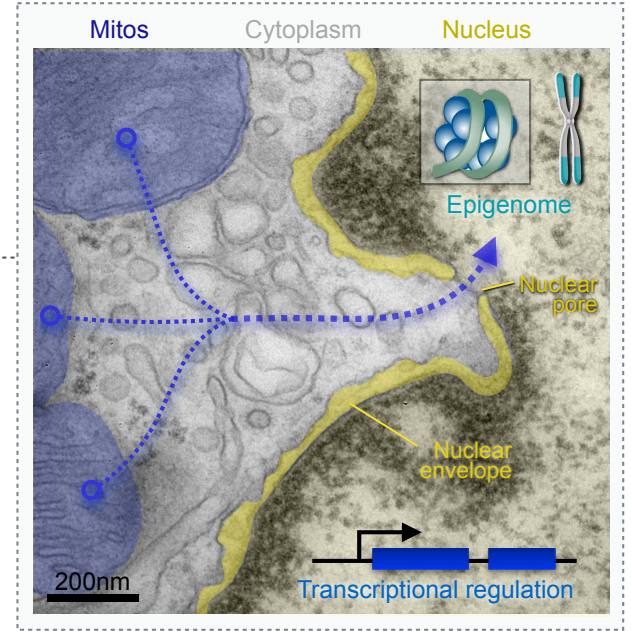
"Mitochondria are the processor of the cell"



Dynamic remodeling of mito networks



Mito-nuclear unit



SEC theory of aging

PART 1

Mitochondrial Signal Transduction, Hair greying

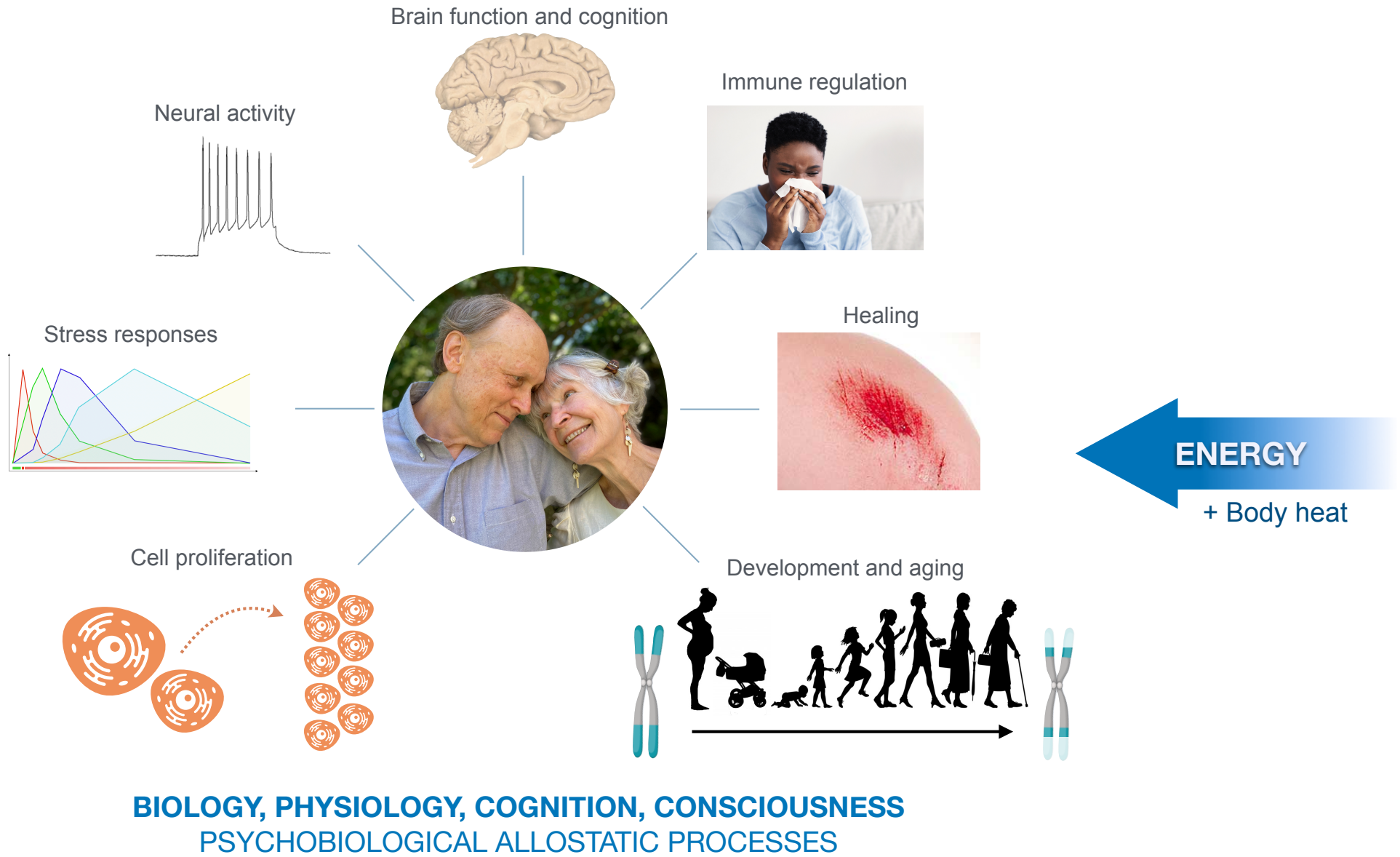
PART 2

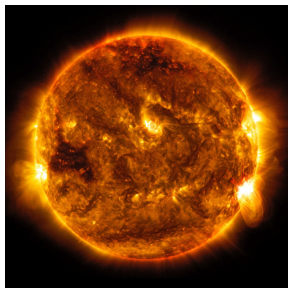
Energy, OxPhos defects, Hypermetabolism, Rate of aging

PART 3

Somato-cognitive Energy Conservation — SEC

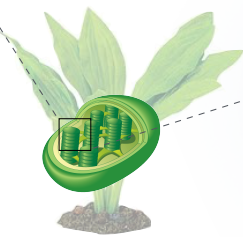
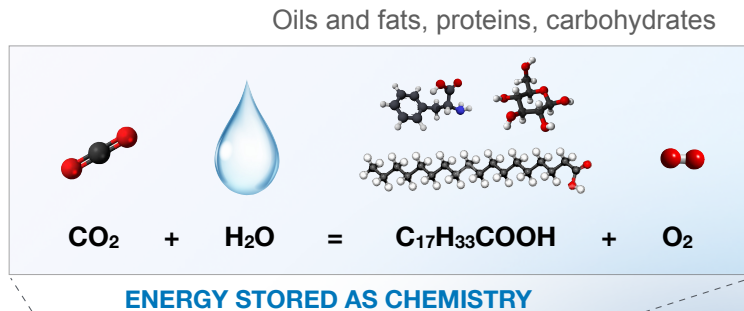
What costs energy?



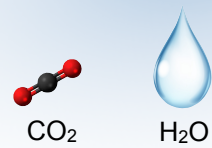
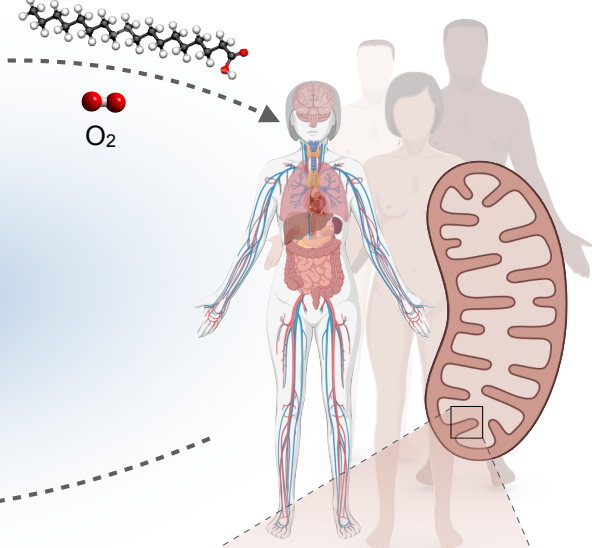


Nuclear fusion
Quantum
electrodynamics

Photons and heat



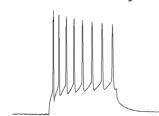
Food **EATING and BREATHING**



Brain function and cognition



Neural activity



Immune regulation



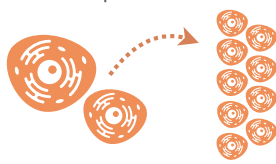
Stress responses



Healing



Cell proliferation



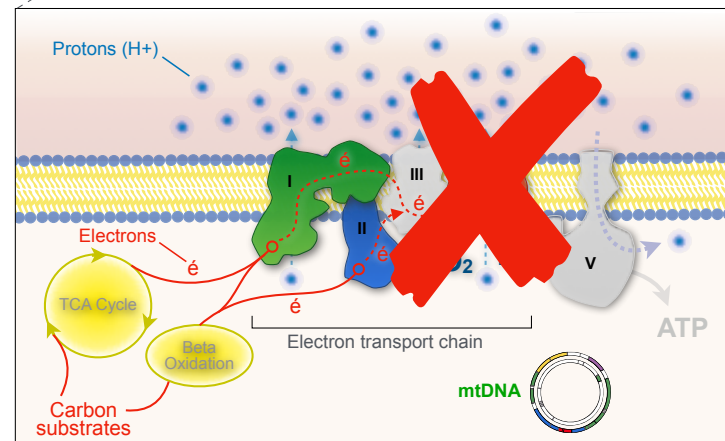
Development and aging



PHYSIOLOGY, COGNITION, CONSCIOUSNESS
PSYCHOBIOLOGICAL ALLOSTATIC PROCESSES

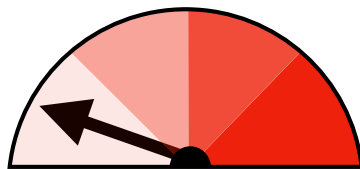
ENERGY
+ Body heat

Electricity and Chemiosmosis
 $\Delta\Psi_m + \Delta\rho\text{H}$

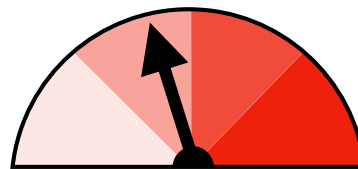


CHEMICAL ENERGY TRANSFORMED INTO
ELECTROCHEMICAL FORCE

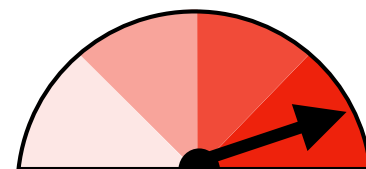
How do OxPhos defects influence cellular energy demand / expenditure?



HYPOmetabolism

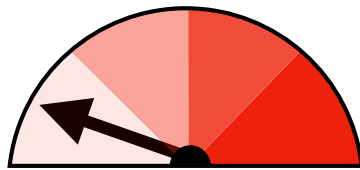


NORMOmetabolism

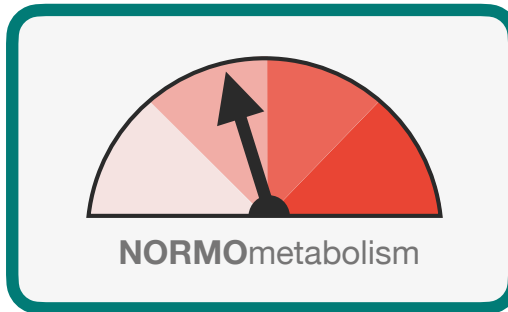


HYPERmetabolism

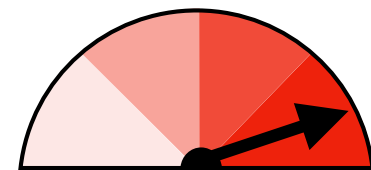
How do OxPhos defects influence cellular energy demand / expenditure?



HYPOmetabolism

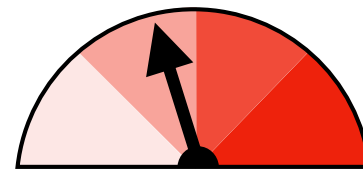
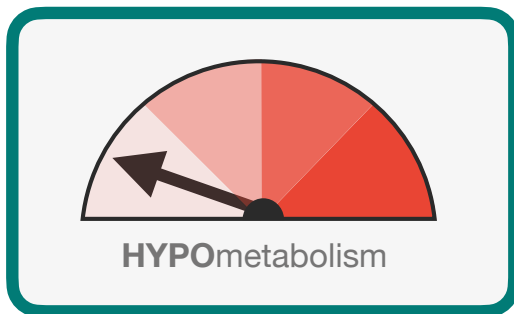


NORMOmetabolism

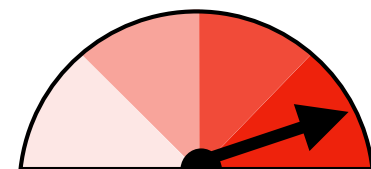


HYPERmetabolism

How do OxPhos defects influence cellular energy demand / expenditure?

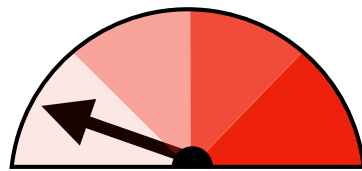


NORMOMETABOLISM

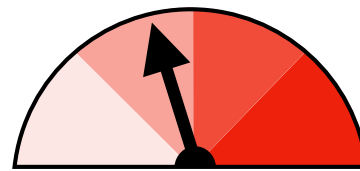


HYPERMETABOLISM

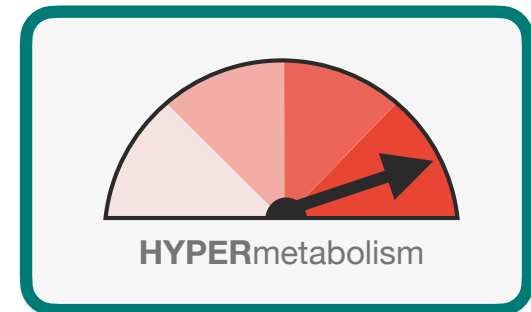
How do OxPhos defects influence cellular energy demand / expenditure?



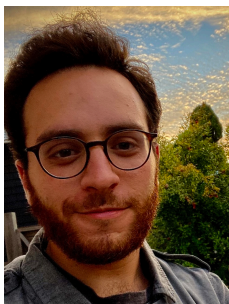
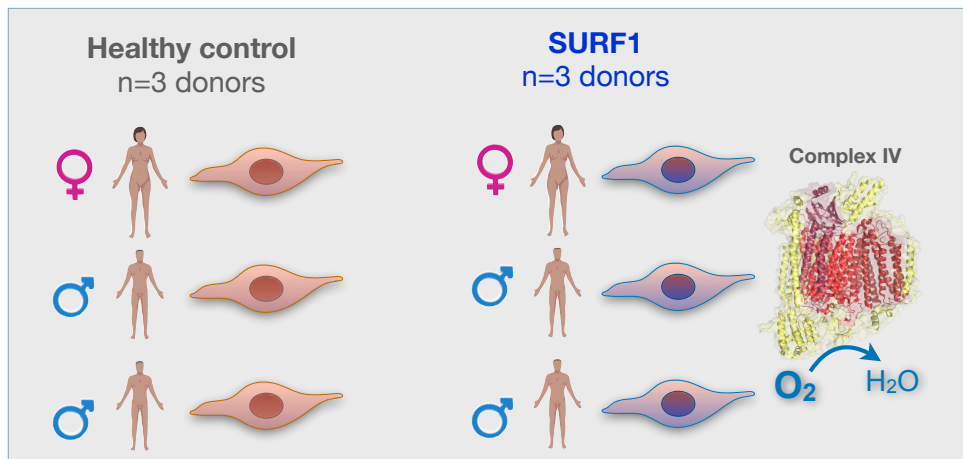
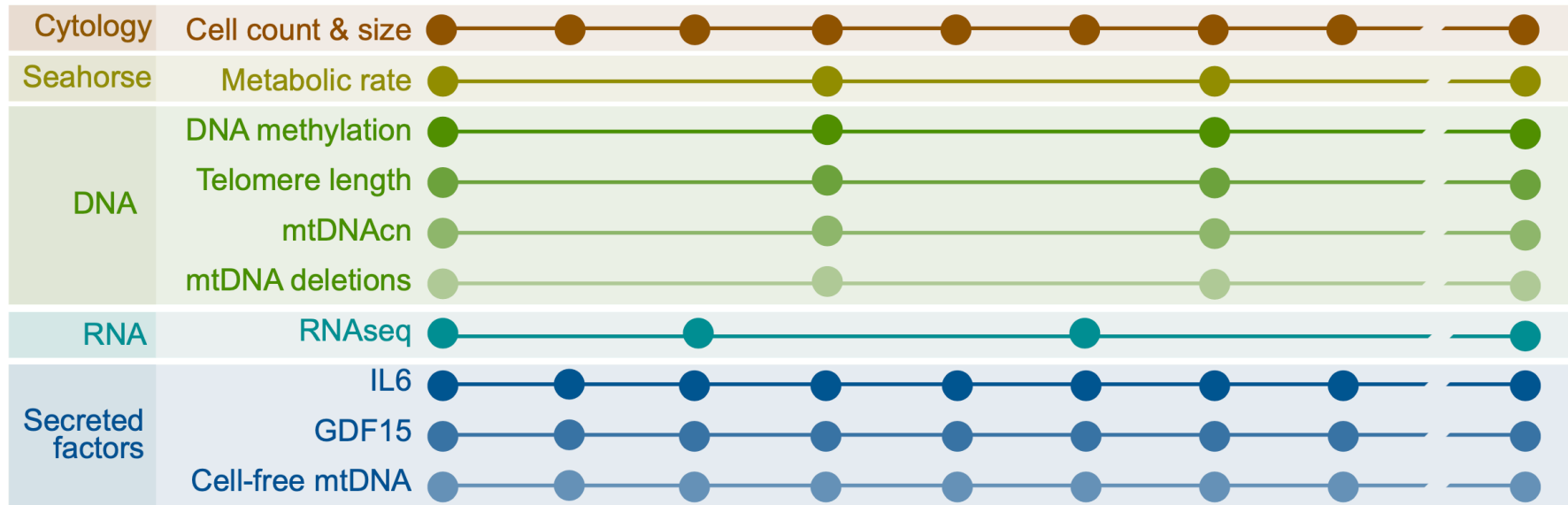
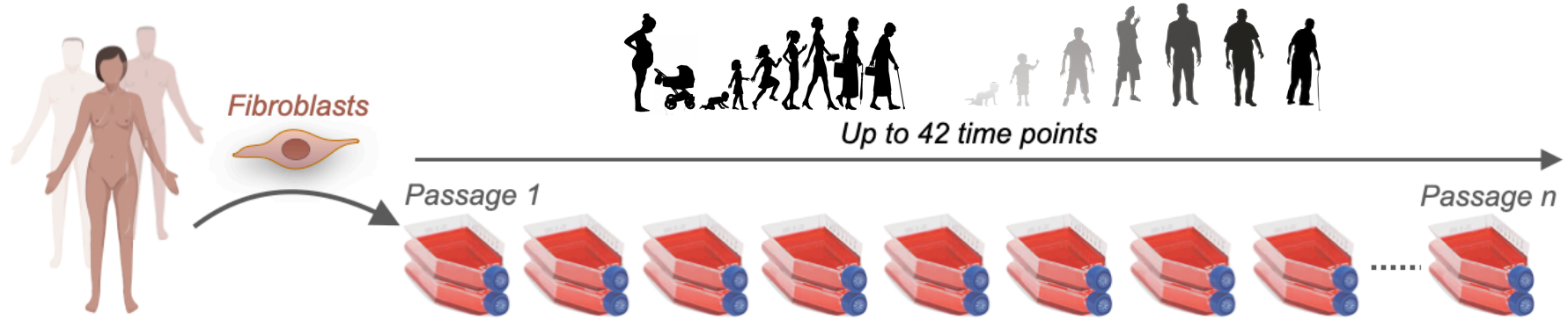
HYPOmetabolism



NORMOmetabolism

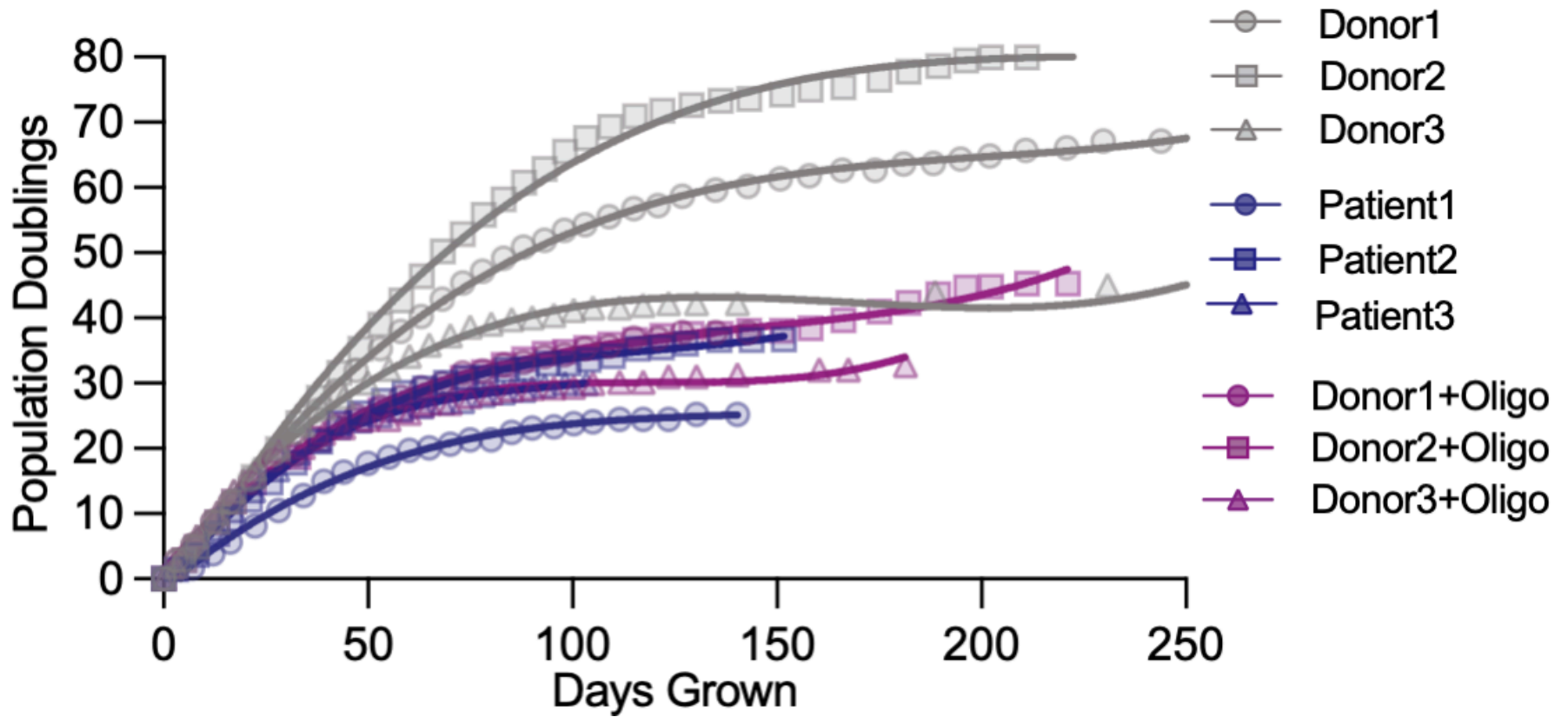


HYPERmetabolism



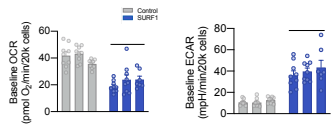
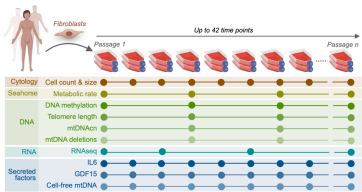
Gabriel Sturm

OxPhos defects **reduce** cell division rate by 32-48%

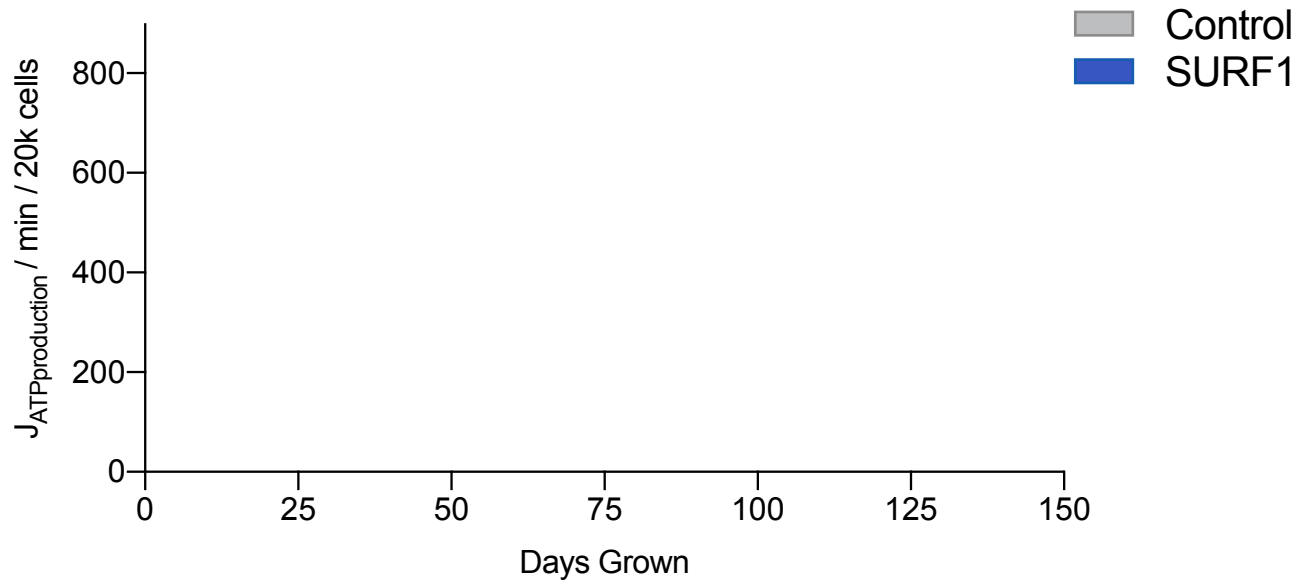
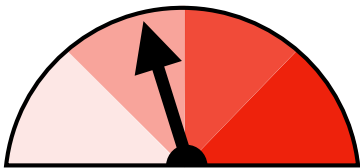


Slower division = less DNA replication, less protein synthesis, less telomerase activity, less mitochondrial biogenesis, ... **ENERGY SAVINGS?**

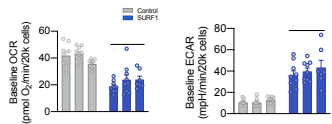
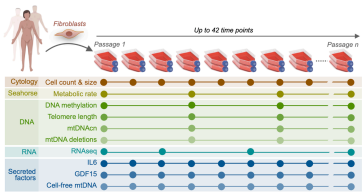
Lifespan trajectories of energy expenditure



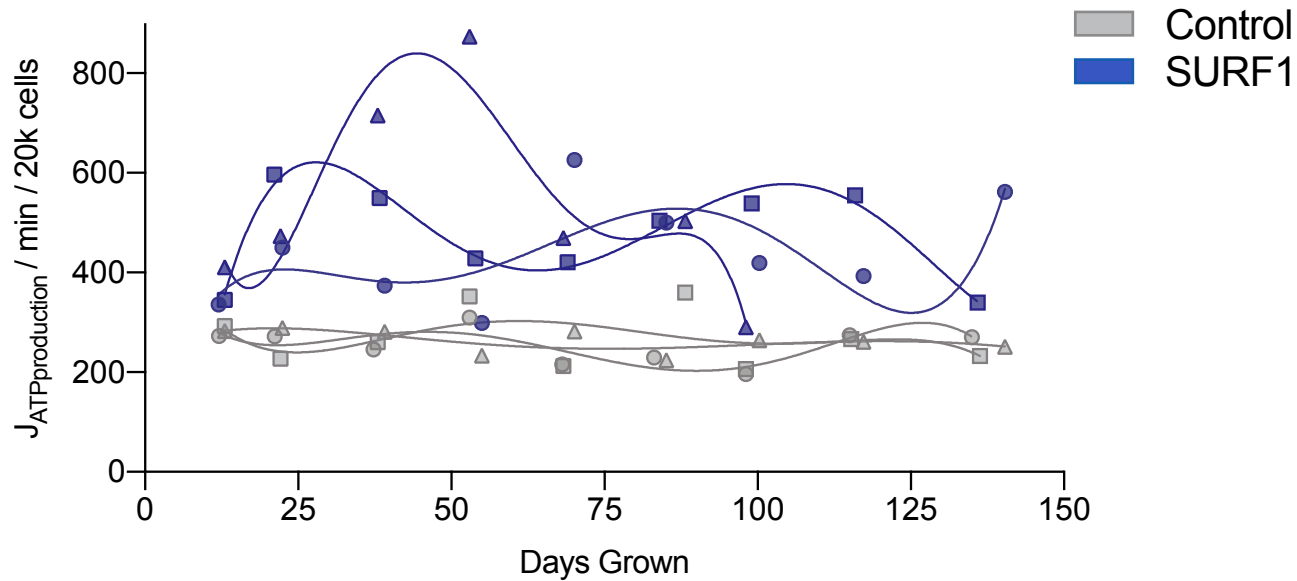
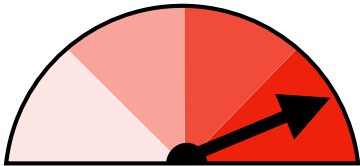
Total energy consumption



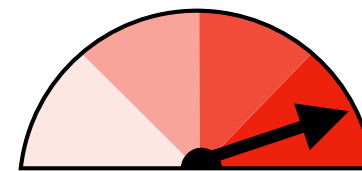
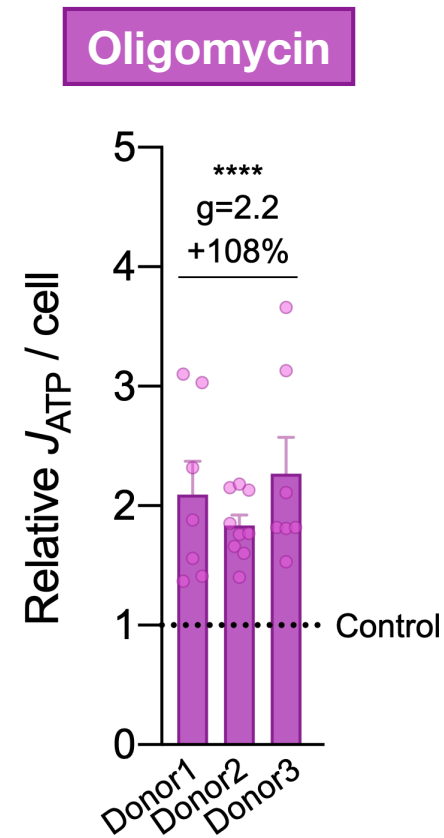
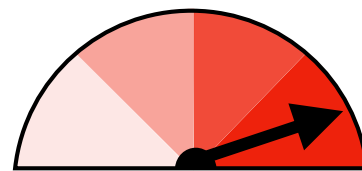
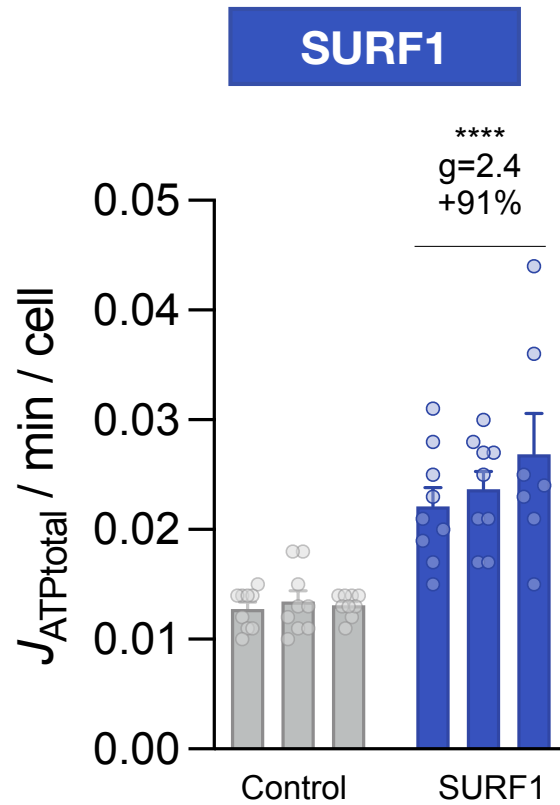
Lifespan trajectories of energy expenditure



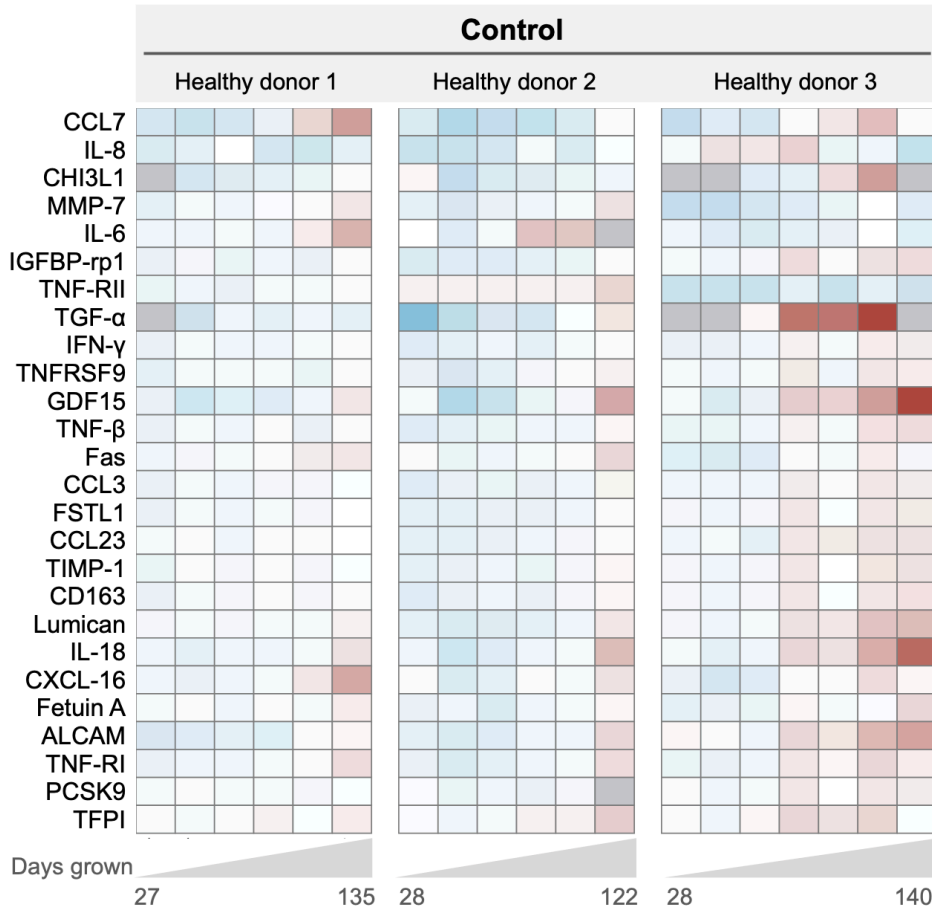
Total energy consumption



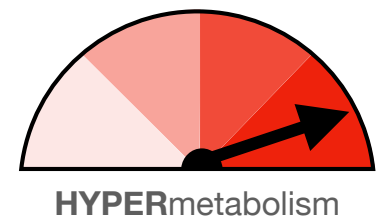
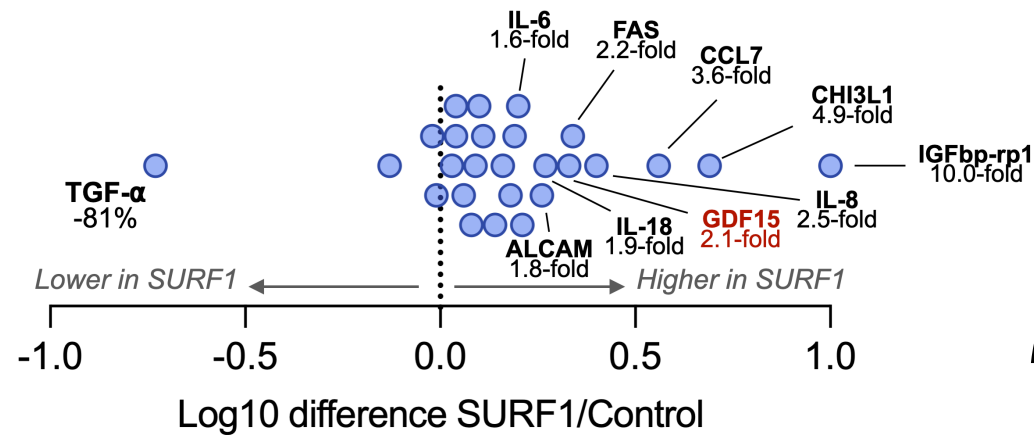
OxPhos-deficient cells are **hypermetabolic**



OxPhos defects increase **cytokine release**

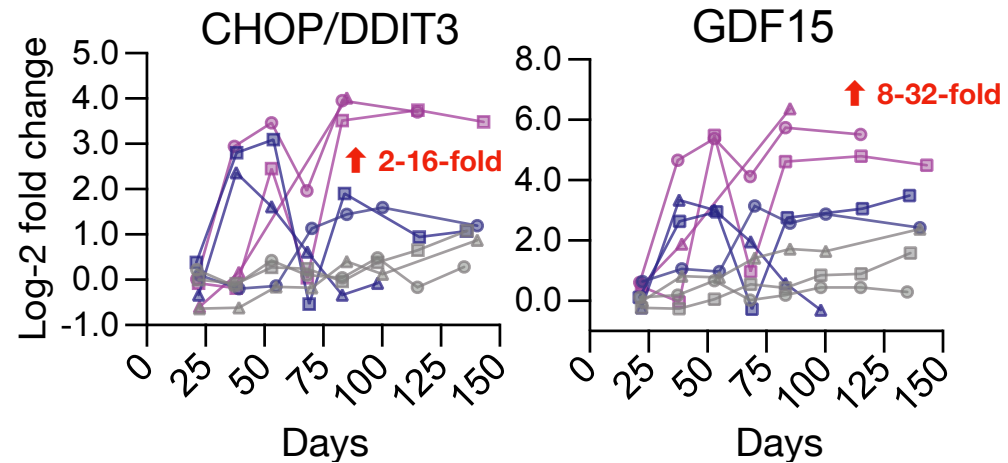
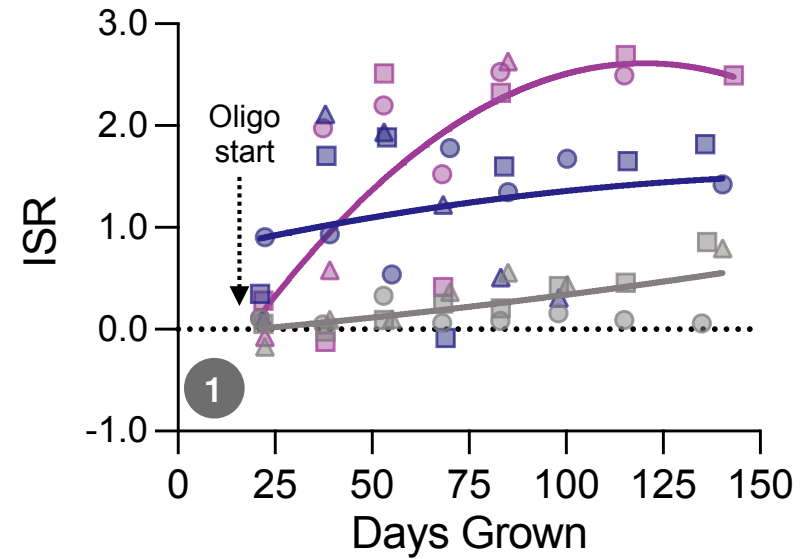
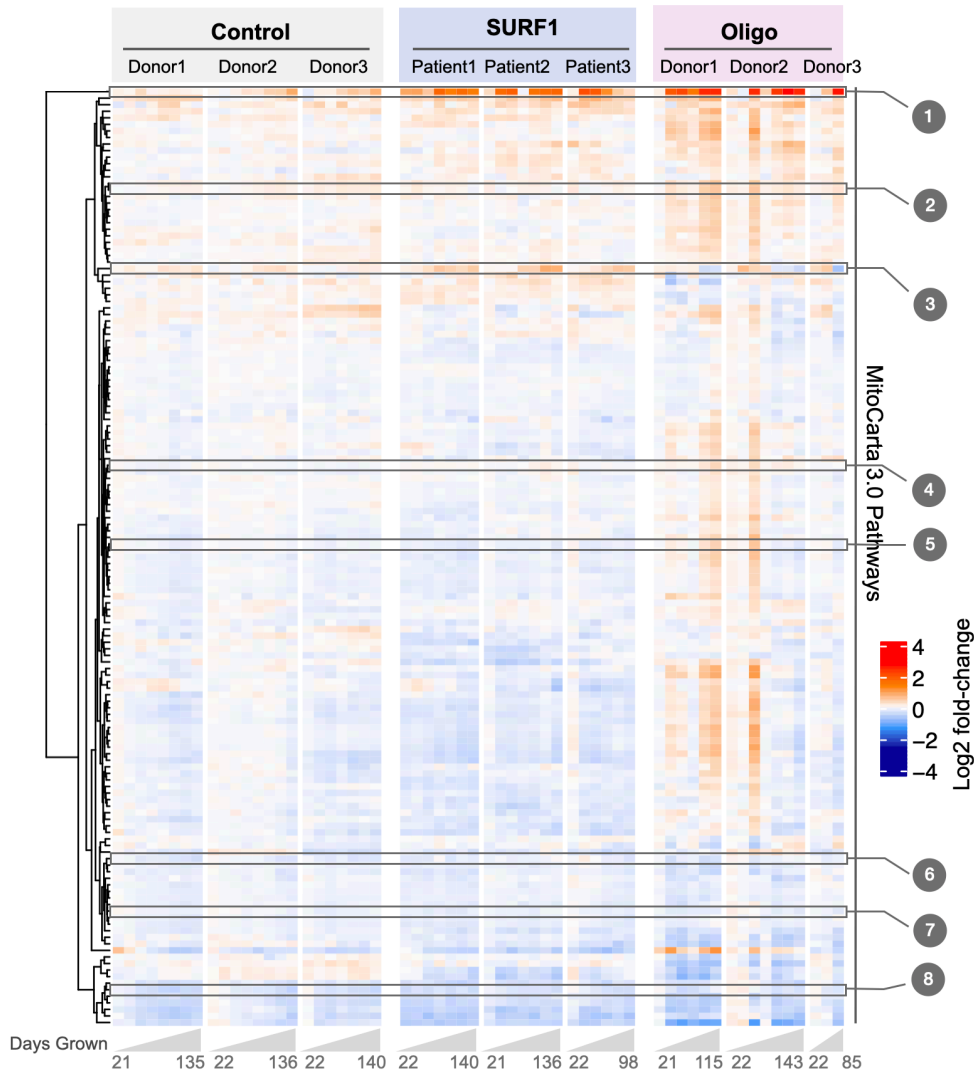


Maximum cytokine levels across the lifespan

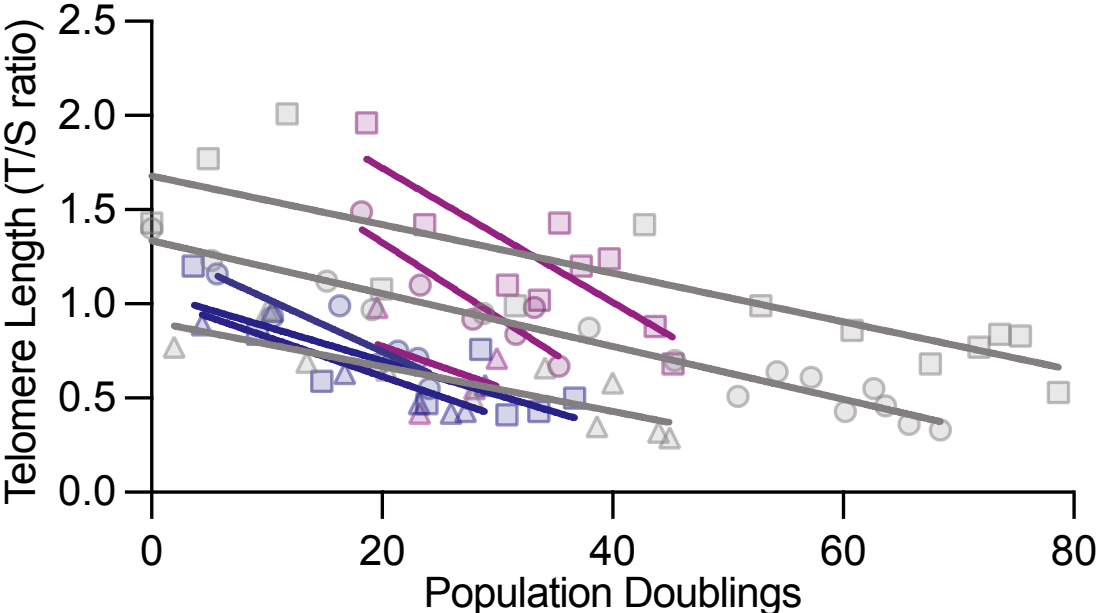
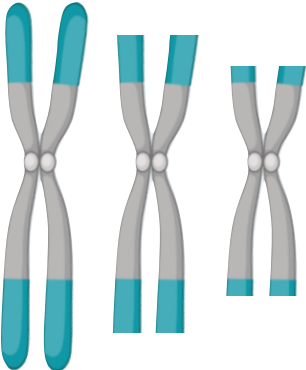


OxPhos defects cause a time-dependent activation of the integrated stress response (ISR)

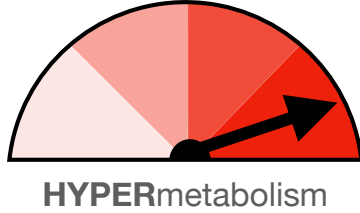
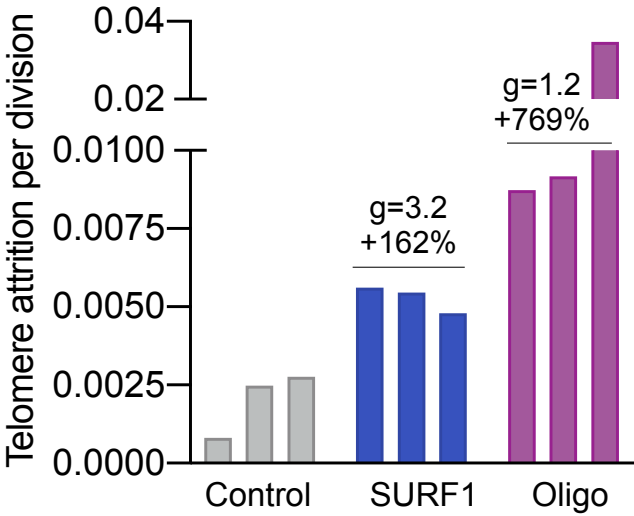
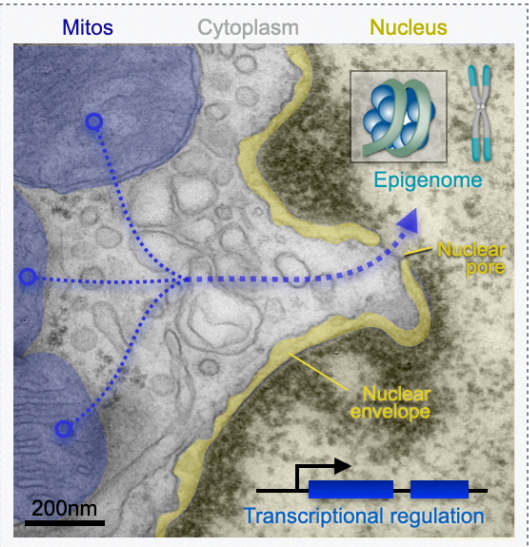
Mitochondrial gene expression



OxPhos defects accelerate telomere shortening rate

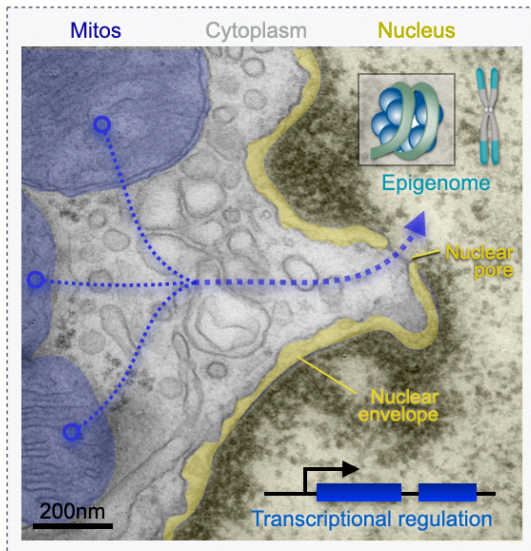
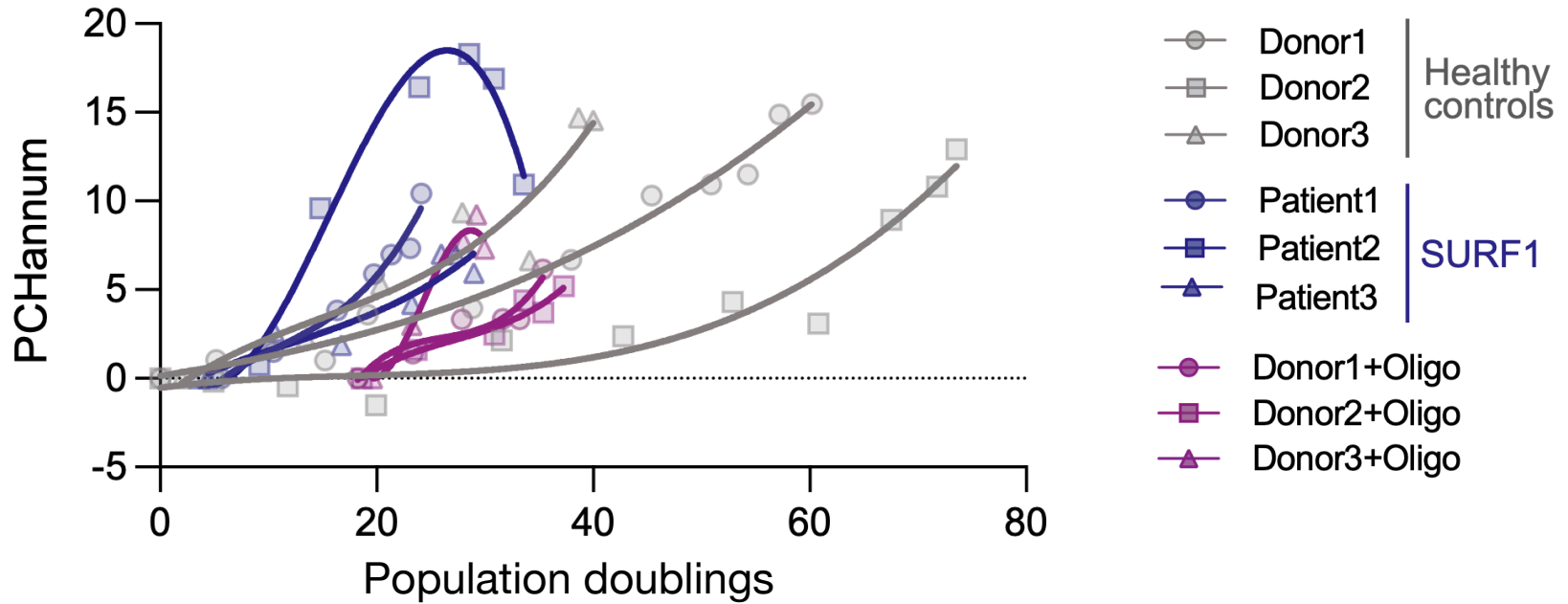


- Donor1 | Healthy controls
- Donor2 | Healthy controls
- △ Donor3 | Healthy controls
- Patient1 | SURF1
- Patient2 | SURF1
- ▲ Patient3 | SURF1
- Donor1+Oligo
- Donor2+Oligo
- ▲ Donor3+Oligo

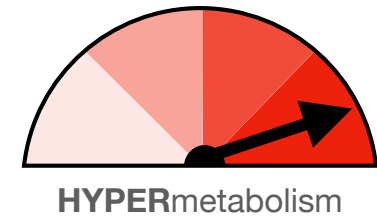
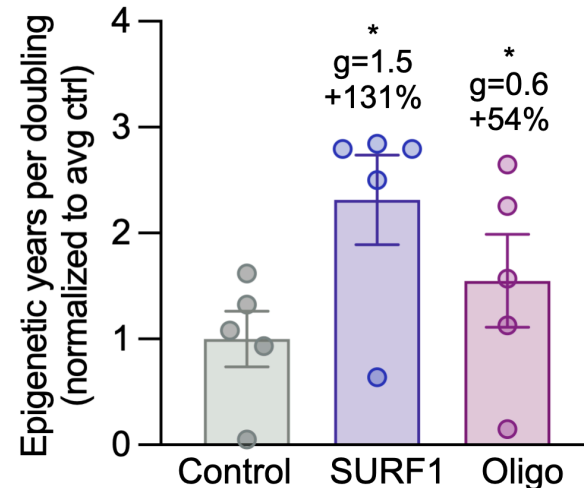


Jue Lin, Elissa Epel
Gabriel Sturm

OxPhos defects accelerate epigenetic aging DNA methylation clocks

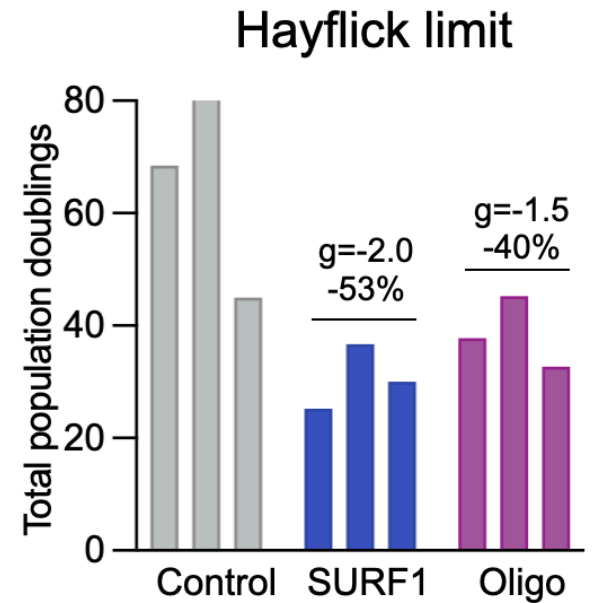
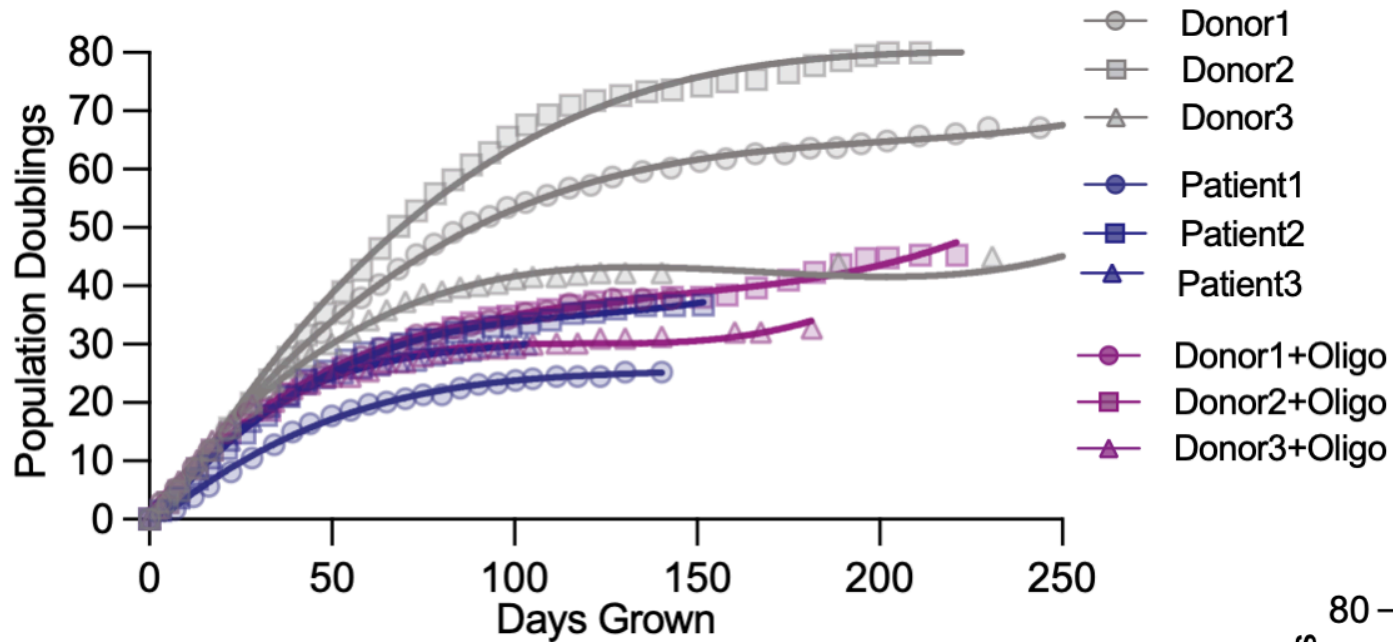


Average rate of epigenetic aging



Steve Horvath, Morgan Levine.
Albert Higgins-Chen
Gabriel Sturm

Hypermetabolic cells have a reduced Hayflick limit



scientific data



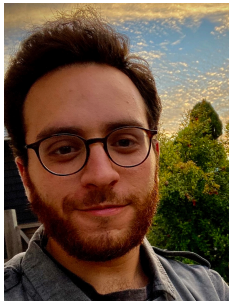
OPEN

DATA DESCRIPTOR

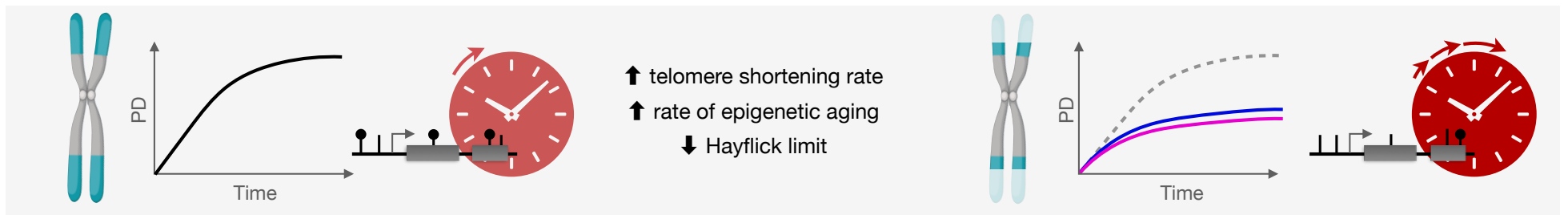
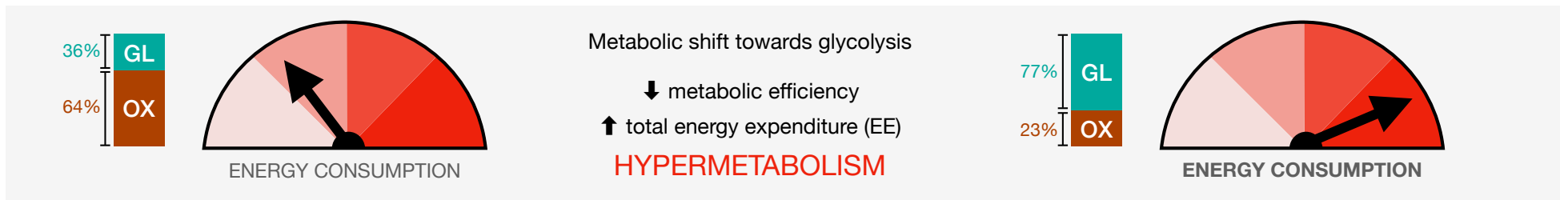
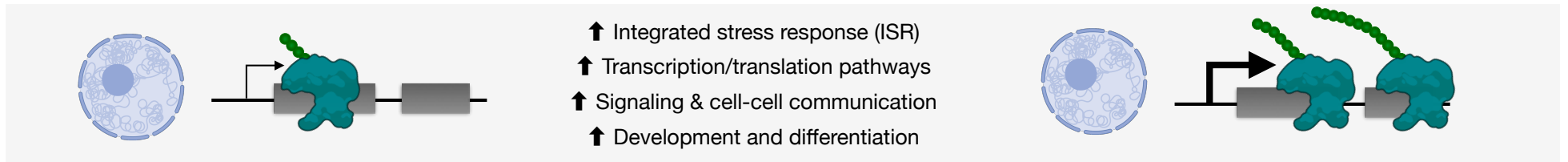
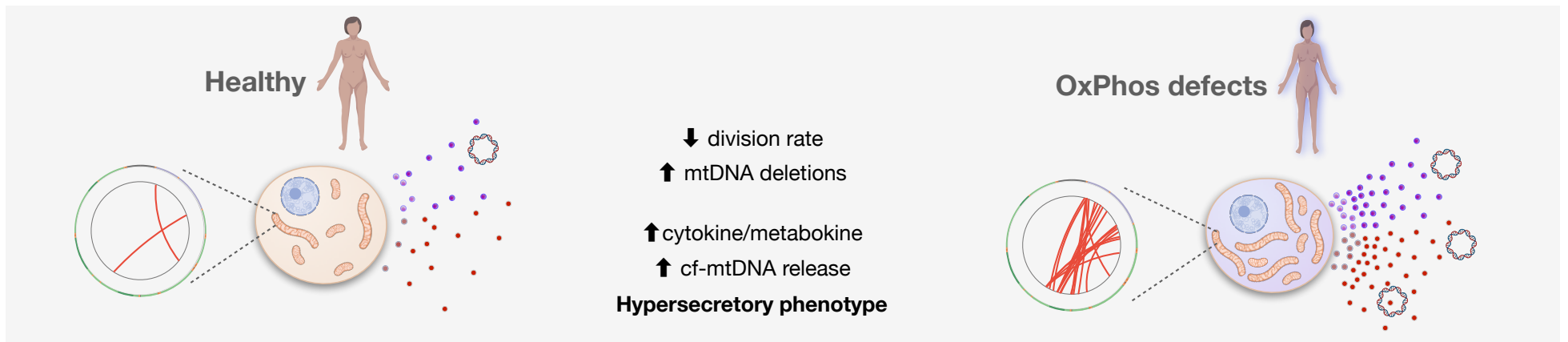
A multi-omics longitudinal aging dataset in primary human fibroblasts with mitochondrial perturbations

Gabriel Sturm^{1,2}, Anna S. Monzel¹, Kalpita R. Karan¹, Jeremy Michelson¹, Sarah A. Ware³, Andres Cardenas⁴, Jue Lin², Céline Bris^{5,6}, Balaji Santhanam⁷, Michael P. Murphy⁸, Morgan E. Levine^{9,10}, Steve Horvath^{10,11}, Daniel W. Belsky¹², Shuang Wang¹³, Vincent Procaccio^{5,6}, Brett A. Kaufman³, Michio Hirano¹⁴ & Martin Picard^{1,14,15} ✉

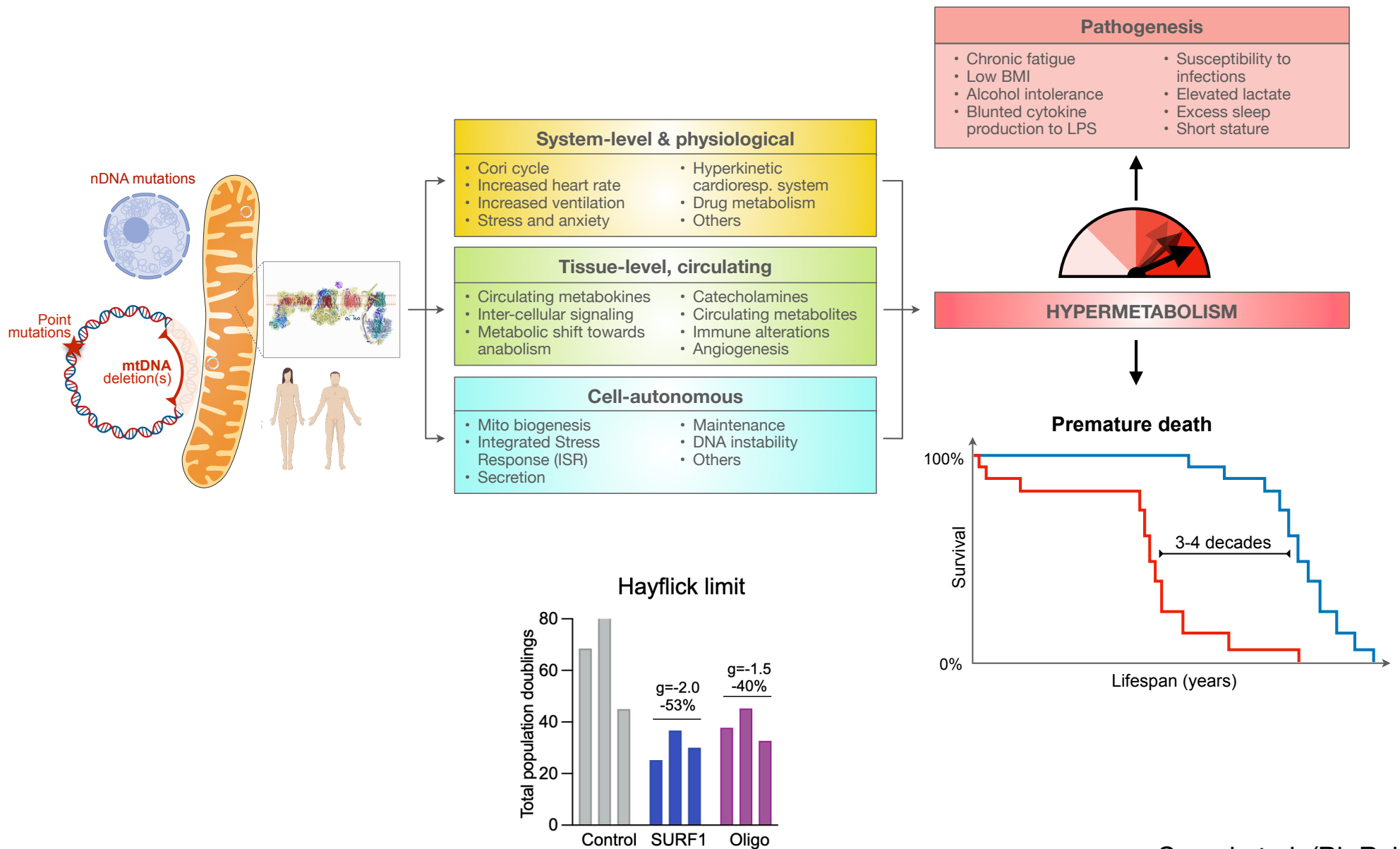
Aging is a process of progressive change. To develop biological models of aging, longitudinal datasets with high temporal resolution are needed. Here we report a multi-omics longitudinal dataset for cultured primary human fibroblasts measured across their replicative lifespans. Fibroblasts were sourced from both healthy donors (n = 6) and individuals with lifespan-shortening mitochondrial disease (n = 3). The dataset includes cytological, bioenergetic, DNA methylation, gene expression, secreted proteins, mitochondrial DNA copy number and mutations, cell-free DNA, telomere length, and whole-genome sequencing data. This dataset enables the bridging of mechanistic processes of aging as outlined by the “hallmarks of aging”, with the descriptive characterization of aging such as epigenetic age clocks. Here we focus on bridging the gap for the hallmark mitochondrial metabolism. Our dataset includes measurement of healthy cells, and cells subjected to over a dozen experimental manipulations targeting oxidative phosphorylation (OxPhos), glycolysis, and glucocorticoid signaling, among others. These experiments provide opportunities to test how cellular energetics affect the biology of cellular aging. All data are publicly available at our webtool: https://columbia-picard.shinyapps.io/shinyapp-Lifespan_Study/



Gabriel Sturm



In vivo – OxPhos defects trigger hypermetabolism and shorten lifespan in humans

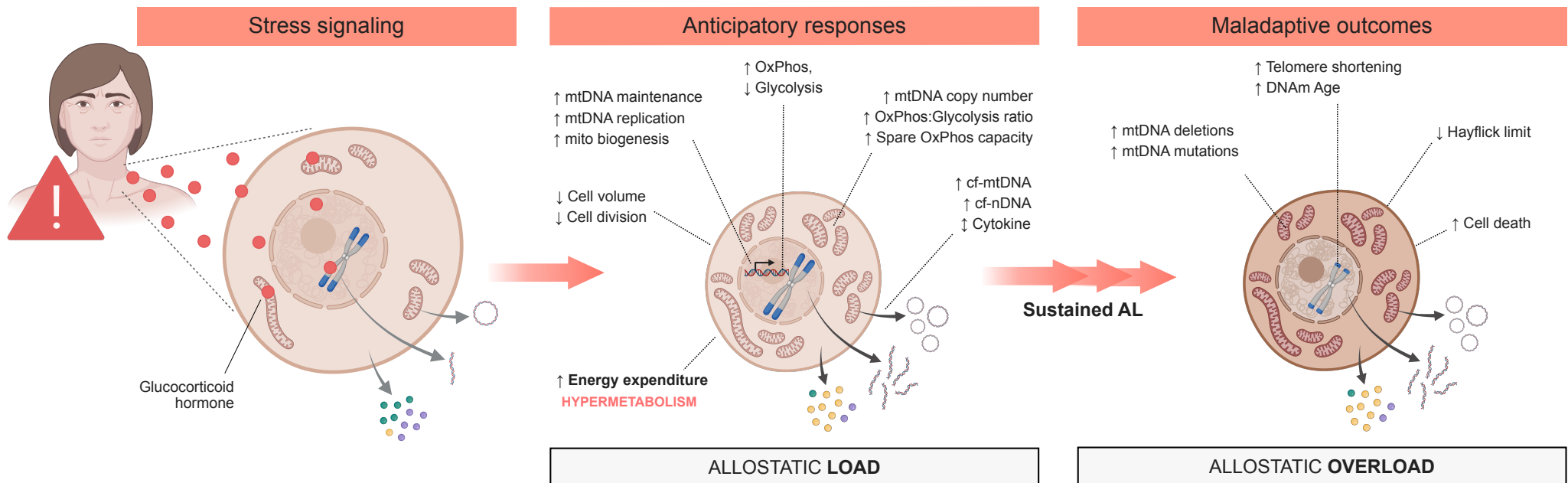




Natalia Bobba-Alves



Cellular allostatic load is linked to increased energy expenditure and accelerated biological aging



Glucocorticoid signaling increases energy expenditure by **60%**

And accelerates cellular aging by **10-40%**

Rate of living hypothesis?

2nd law of thermodynamics

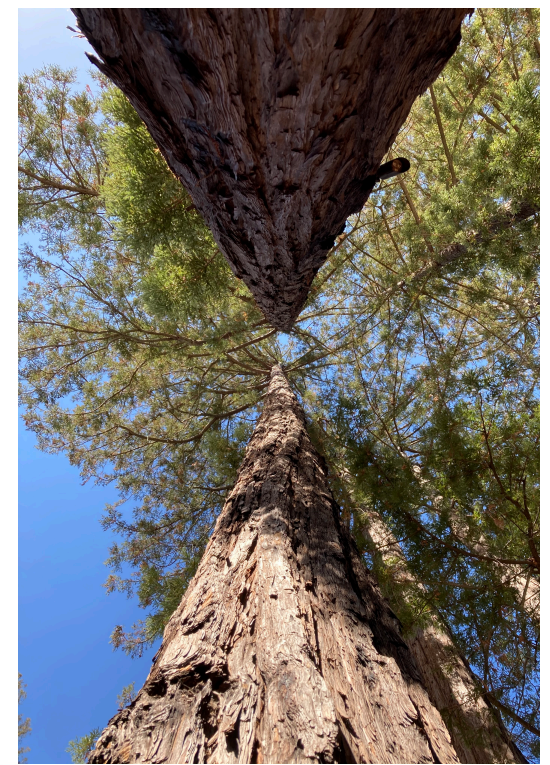
HYPERMETABOLISM



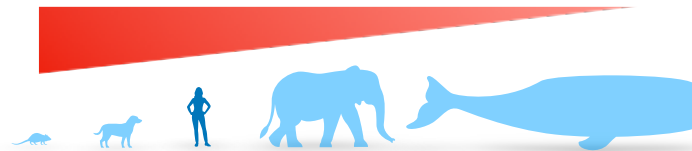
Seconds



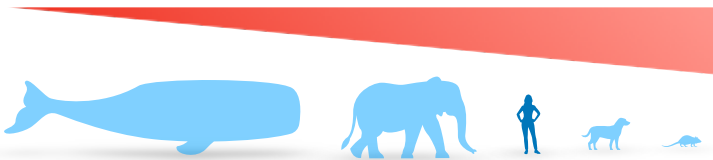
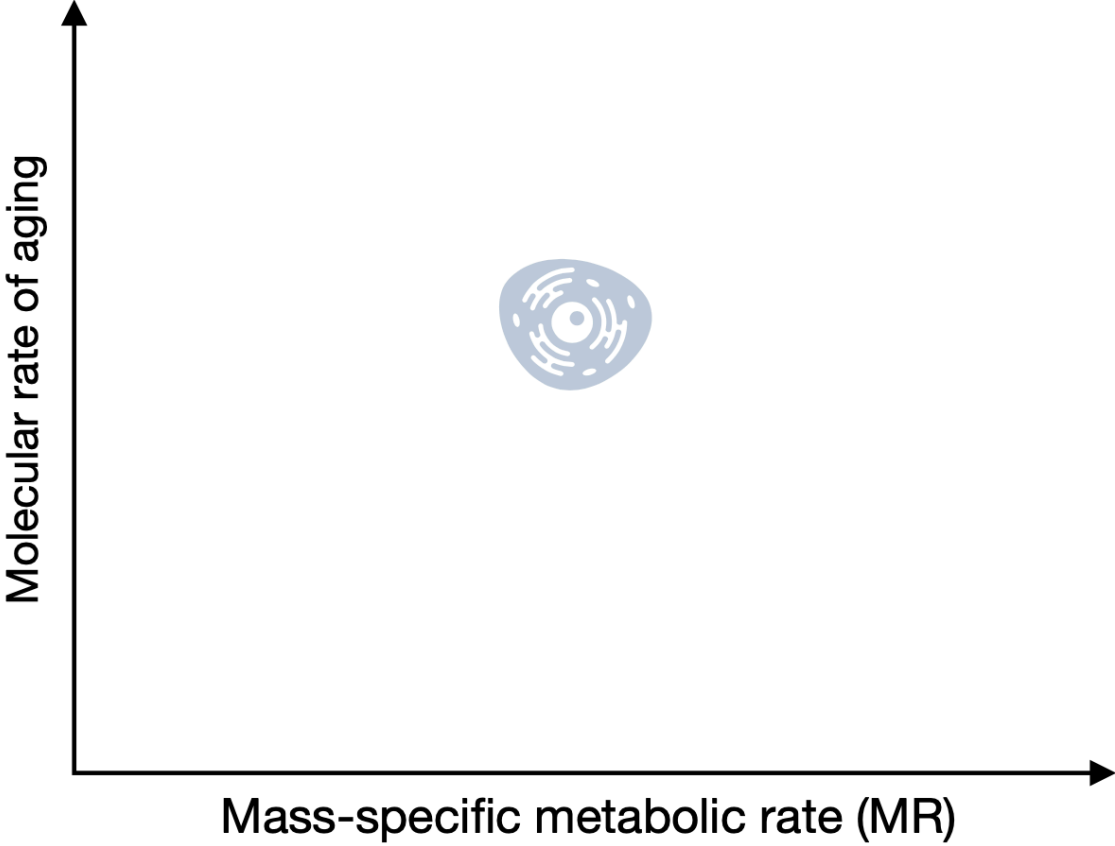
Years / Decades



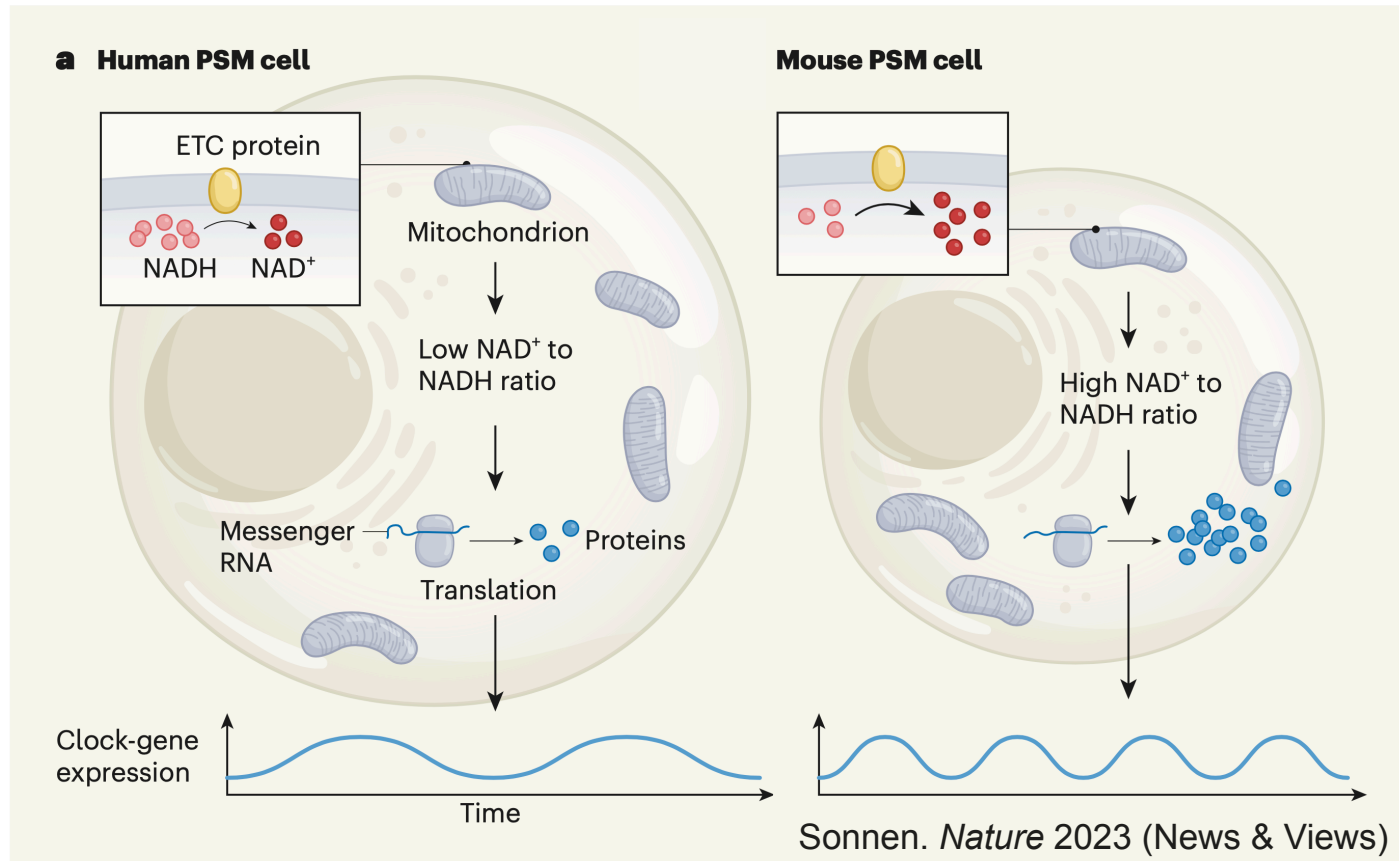
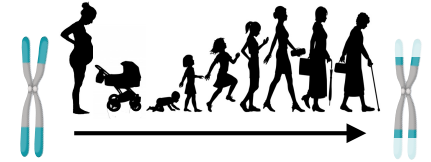
Centuries



Scaling of metabolic rate (MR) & rate of aging



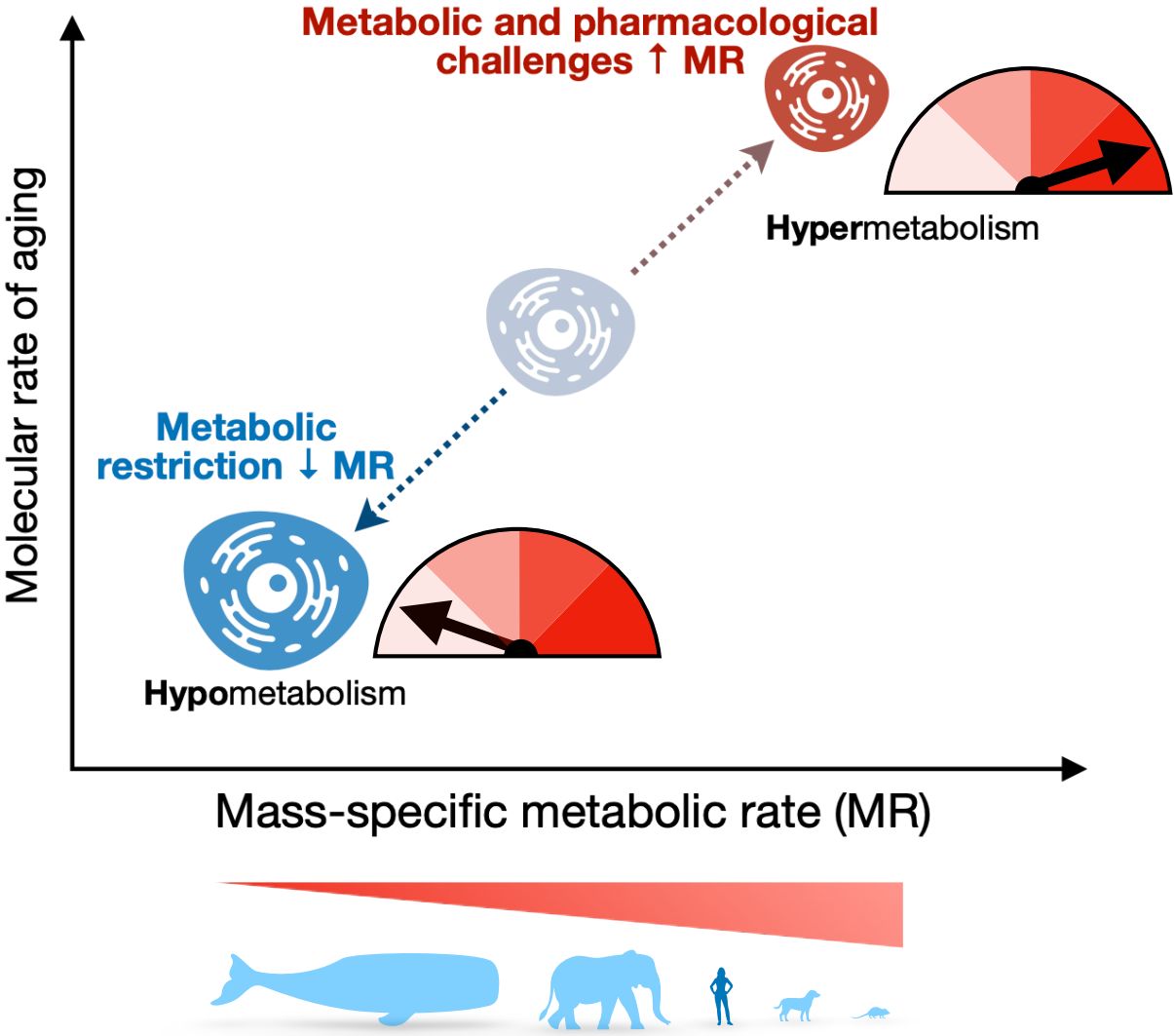
What sets the pace of aging ?

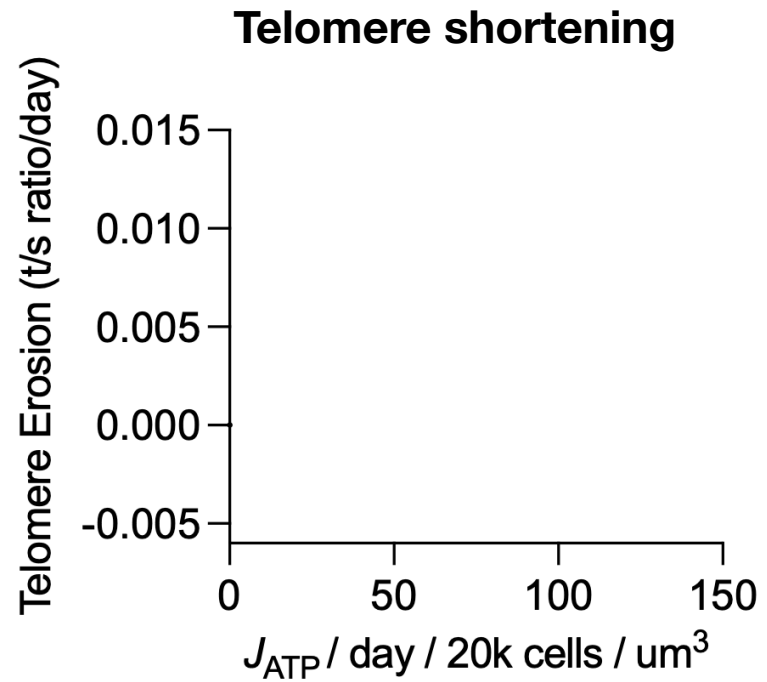
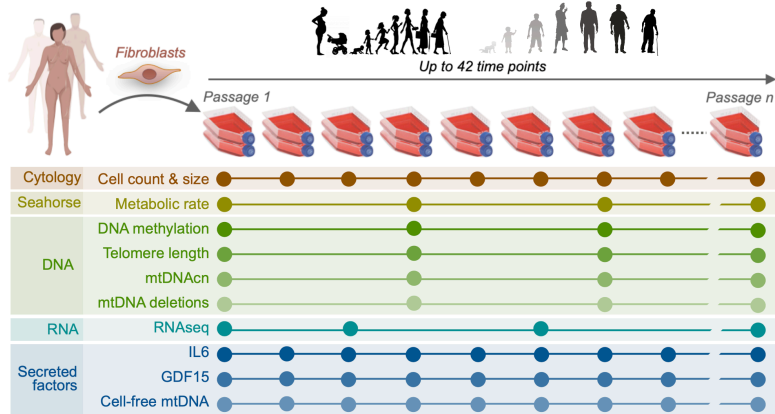
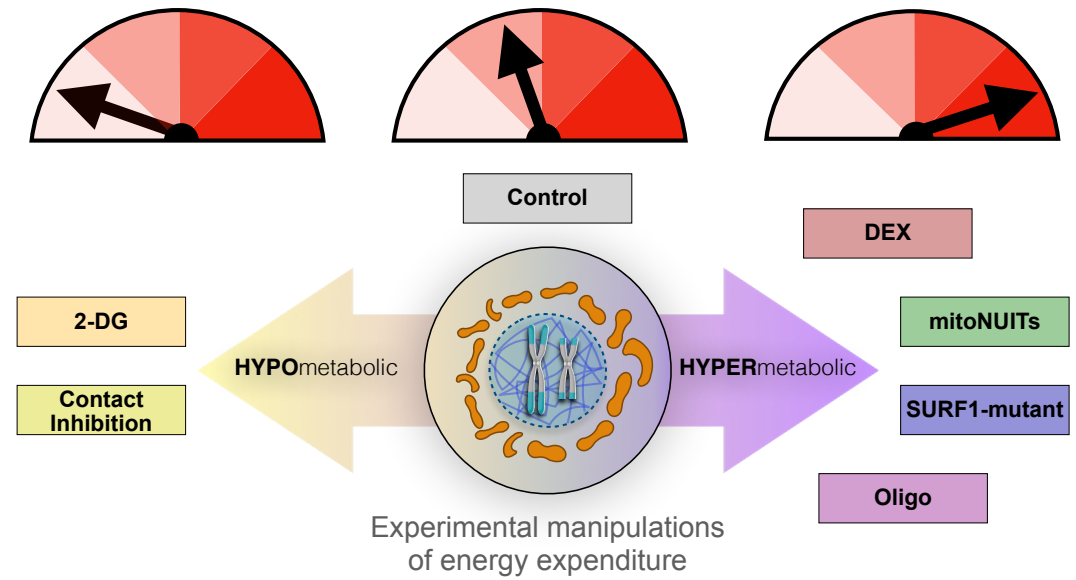
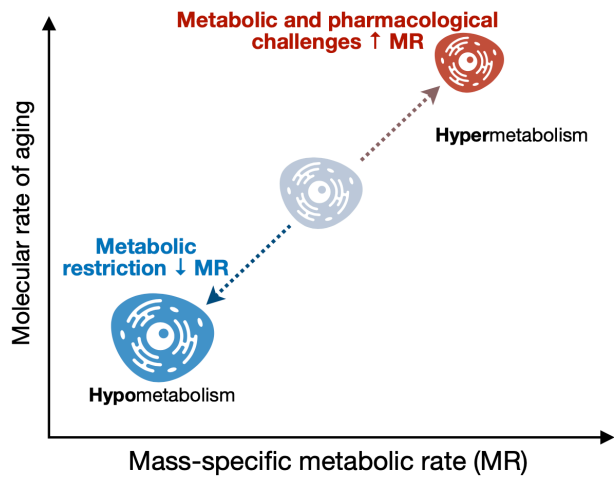


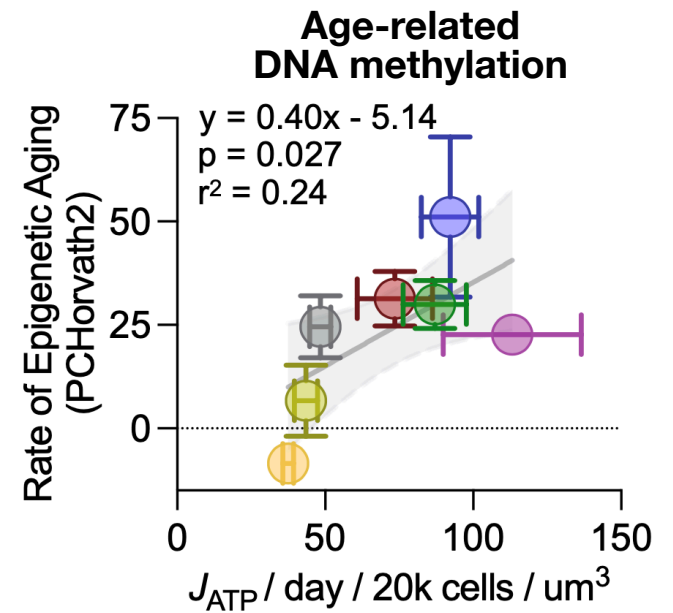
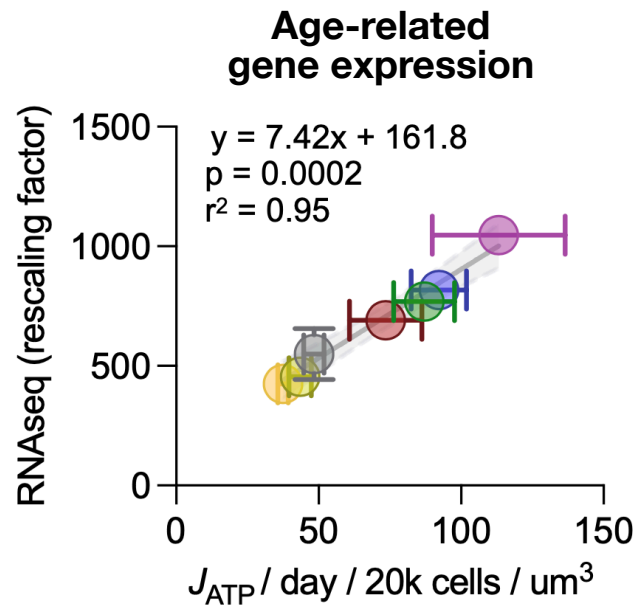
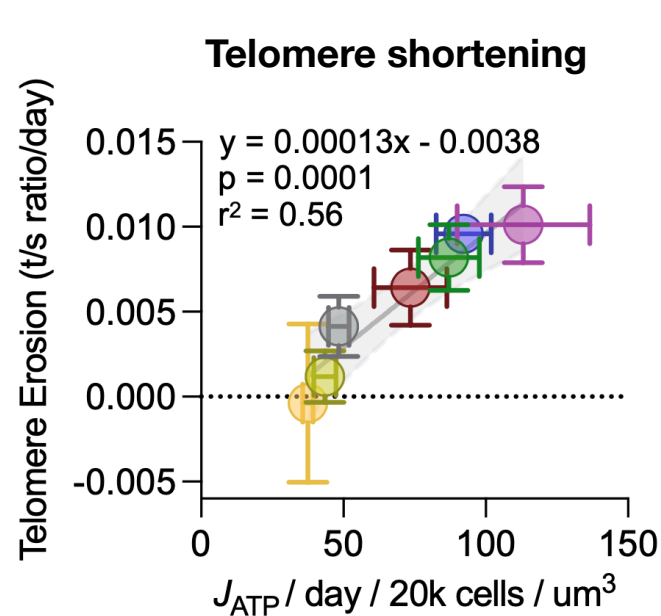
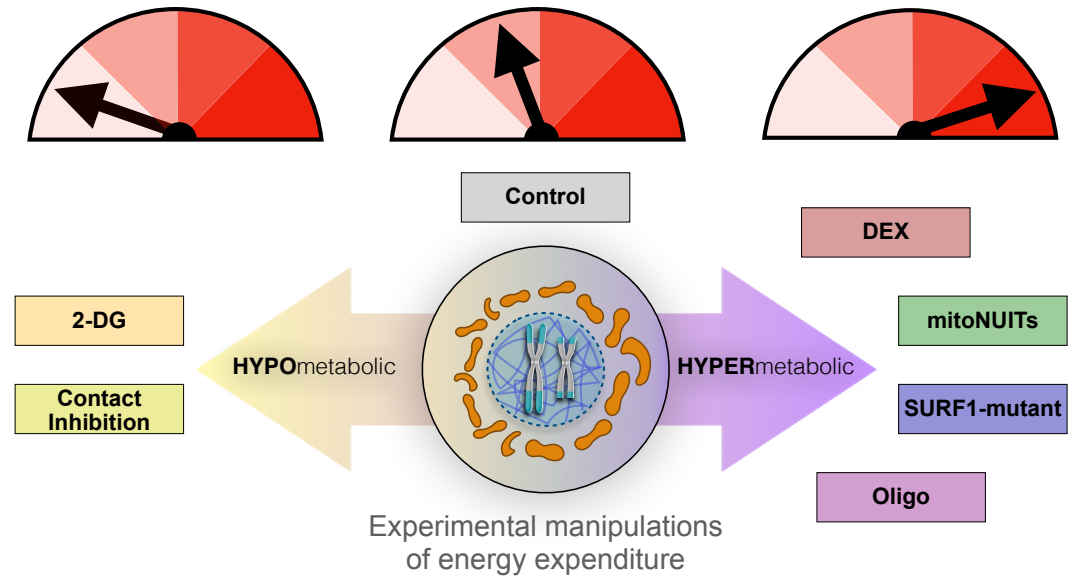
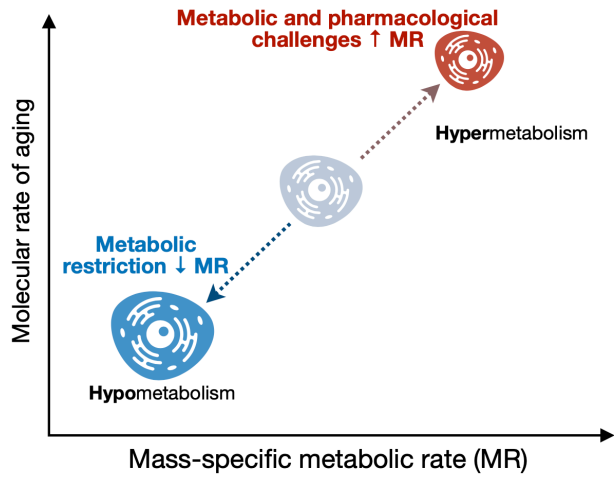
The **rate** of development can be altered by manipulating **mitochondrial energy metabolism**

(Presomitic mesoderm) Diaz-Cuadros et al. *Nature* 2023
(Neuronal development) Iwata et al. *Science* 2023

Scaling of metabolic rate (MR) & rate of aging







Rate of living hypothesis?

2nd law of thermodynamics

HYPERMETABOLISM



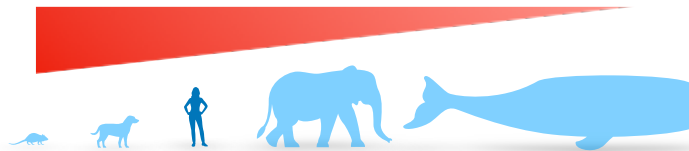
Seconds



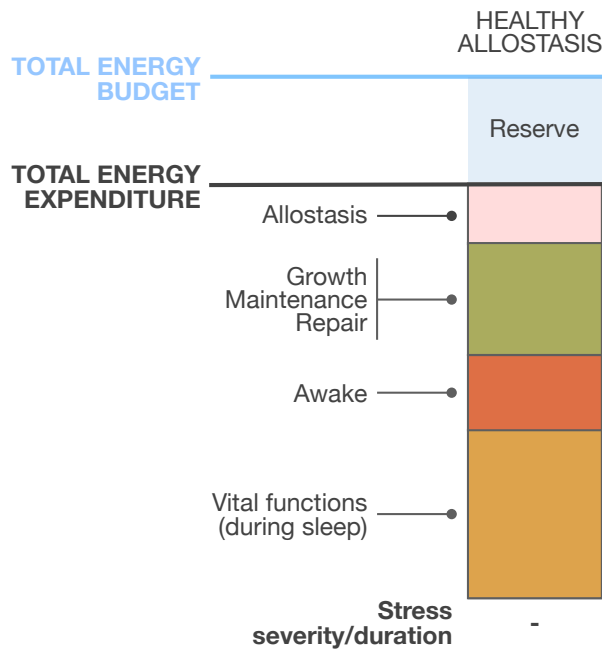
Years / Decades



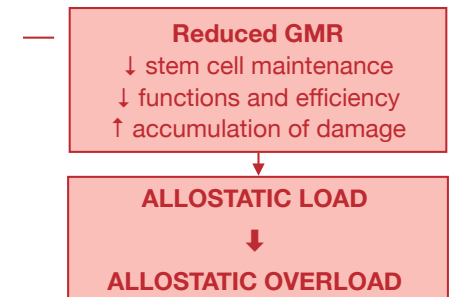
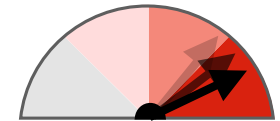
Centuries



Human aging and lifespan determined by partitioning of limited energetic resources ?



$$\text{TOTAL ENERGY EXPENDITURE} = \text{TOTAL ENERGY BUDGET}$$



PART 1

Mitochondrial Signal Transduction, Hair greying

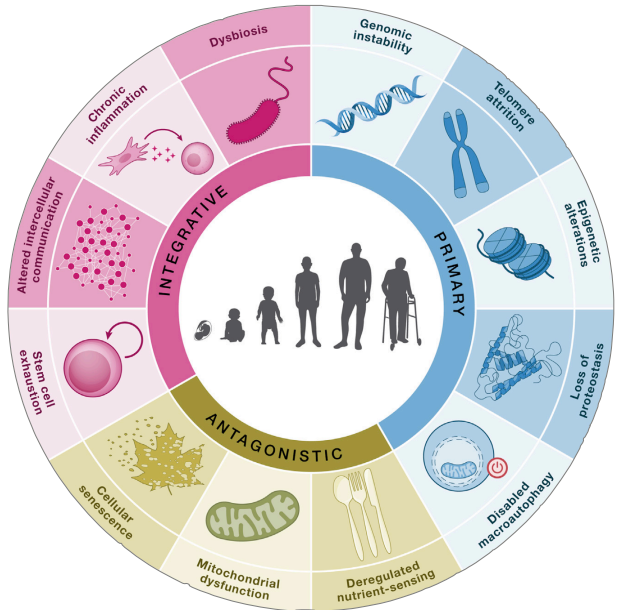
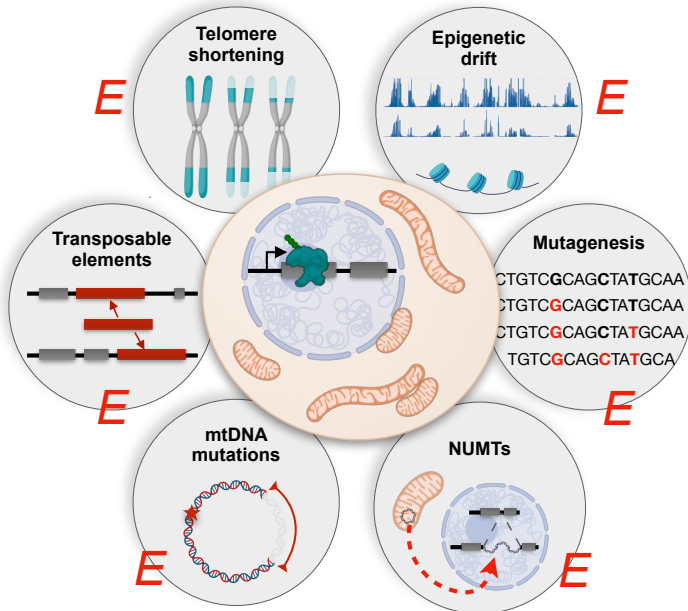
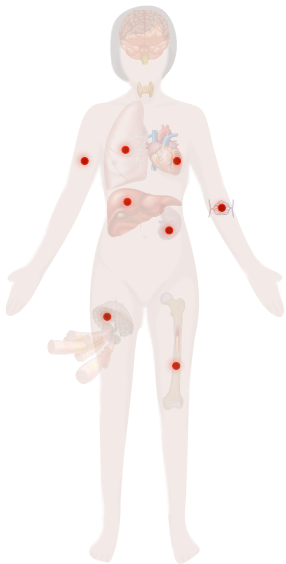
PART 2

Energy, OxPhos defects, Hypermetabolism, Rate of aging

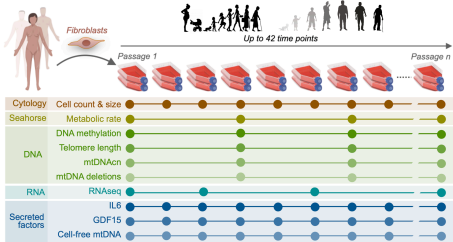
PART 3

Somato-cognitive Energy Conservation — SEC

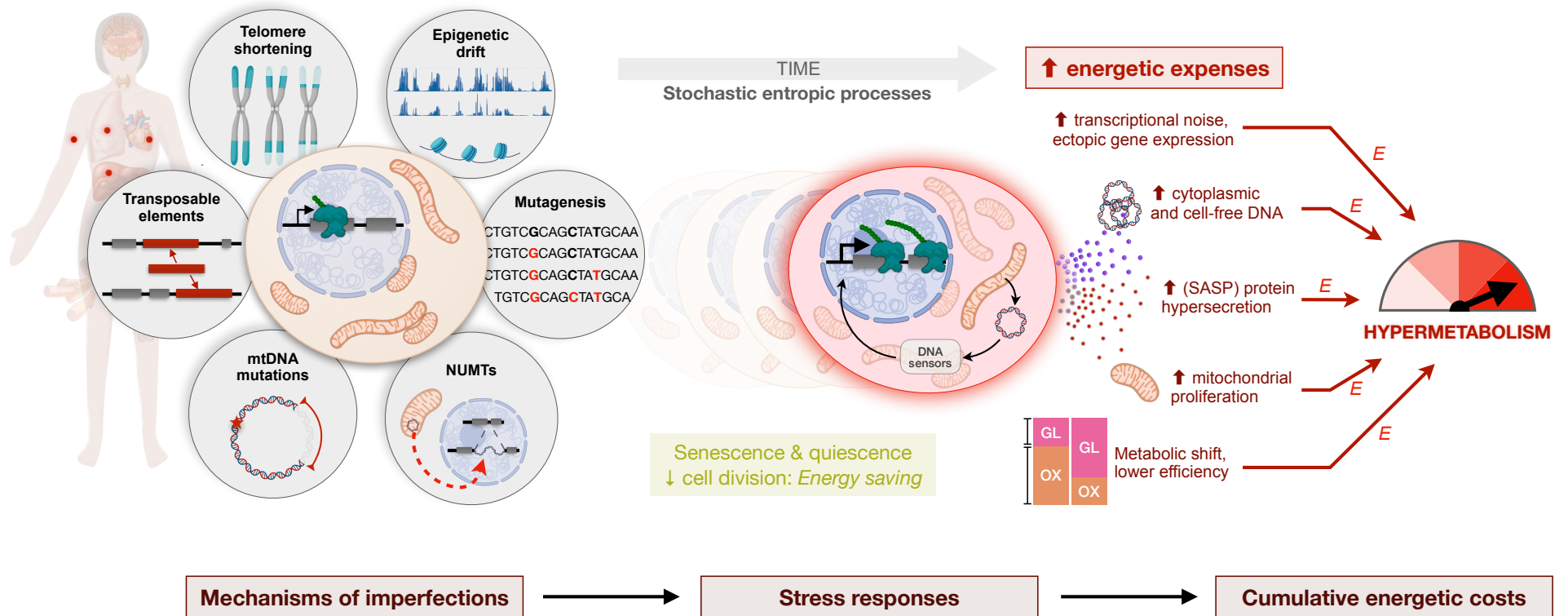
The imperfection of life as (costly) driver of cellular aging



Mechanisms of imperfections



Lopez-Otin et al. *Cell* 2023

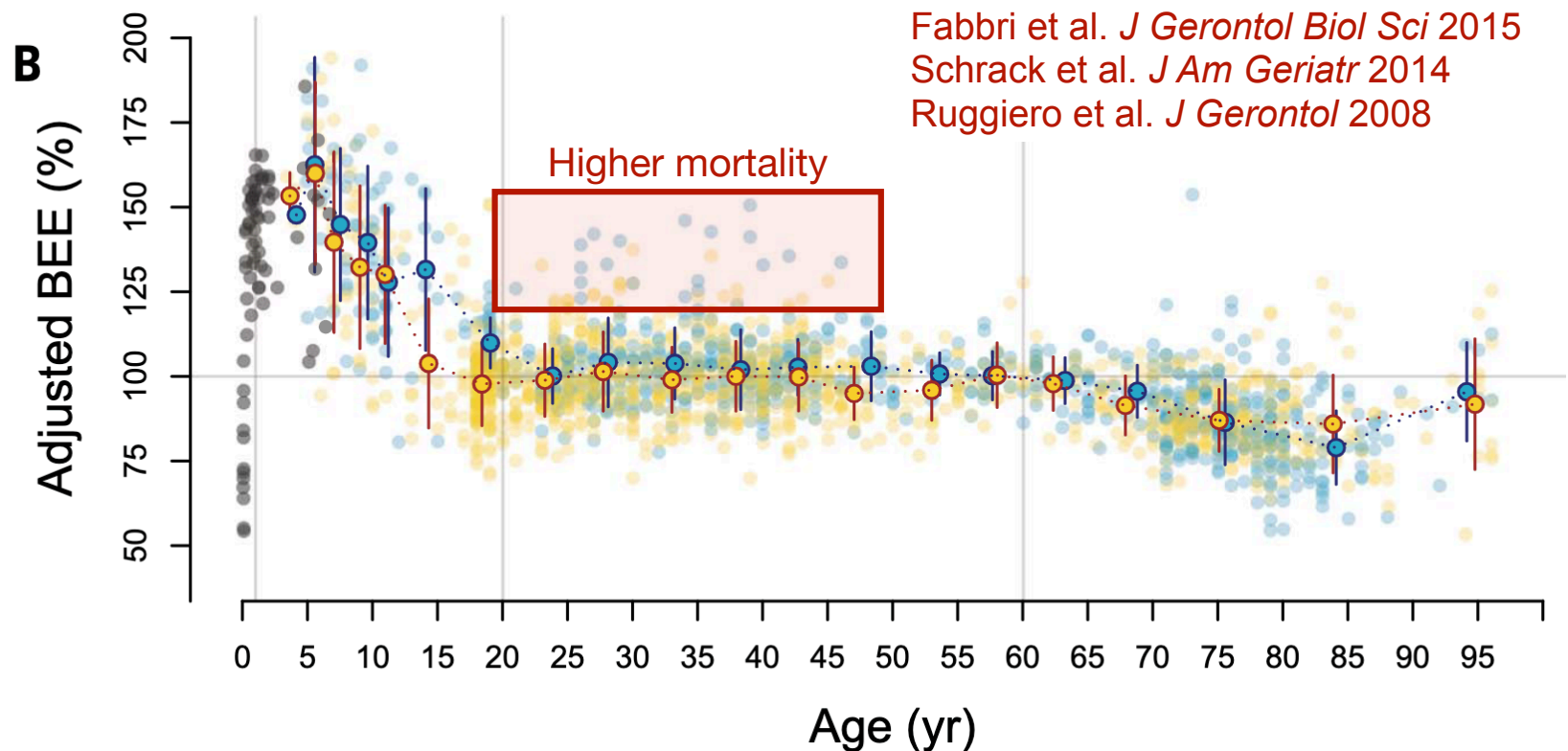


Aging (senescent) cells become **hypermetabolic**

Aging (senescent) cells become **hypermetabolic**

But the whole body does not

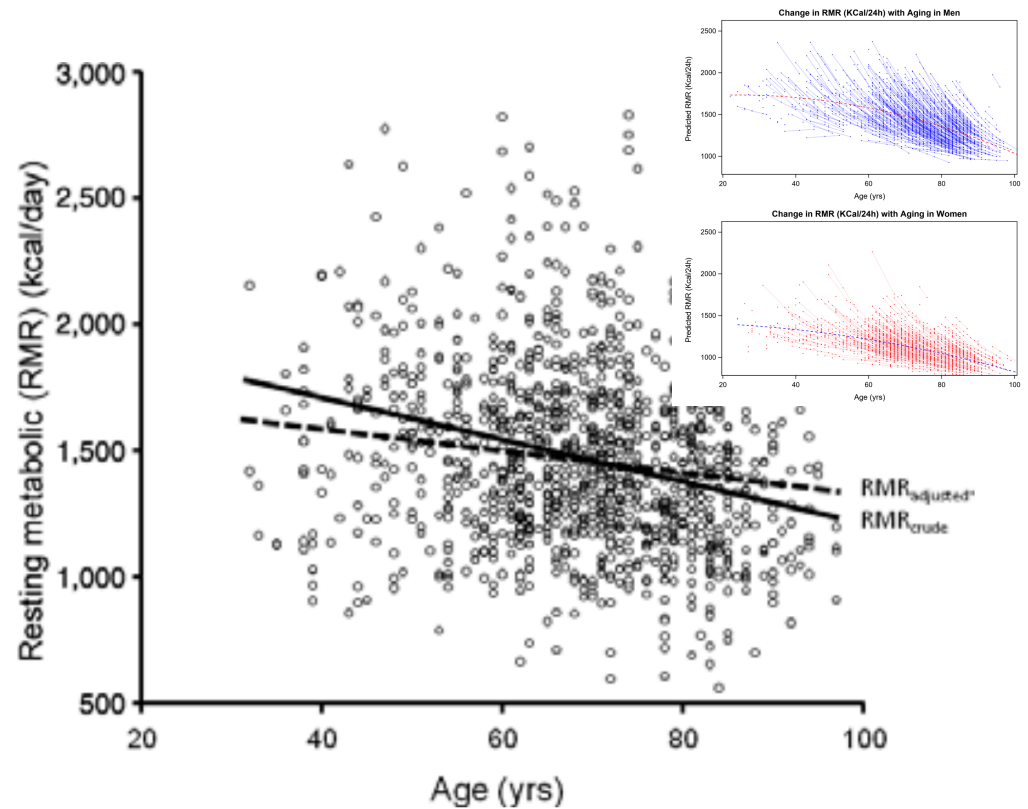
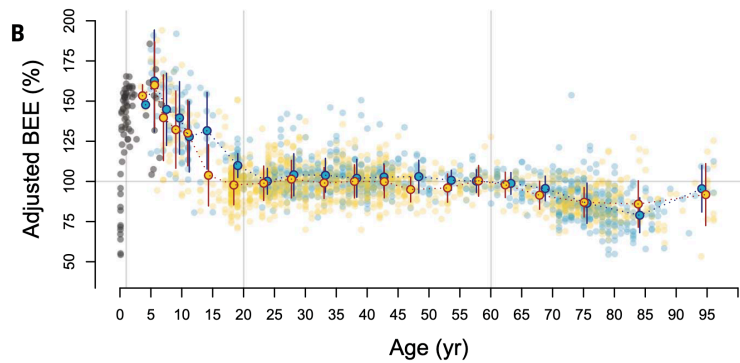
In fact, whole body energy expenditure **declines with age**



Aging (senescent) cells become **hypermetabolic**

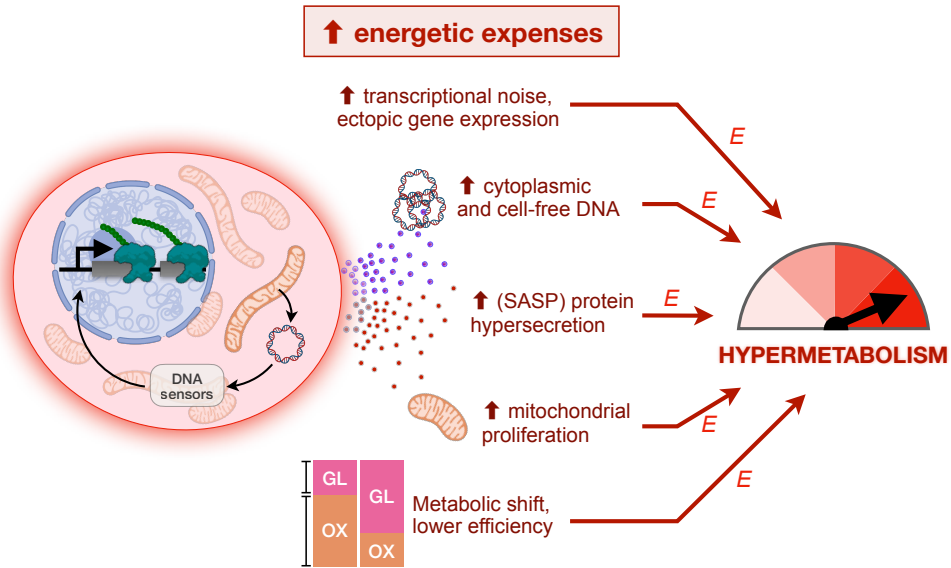
But the whole body does not

In fact, whole body energy expenditure **declines with age**

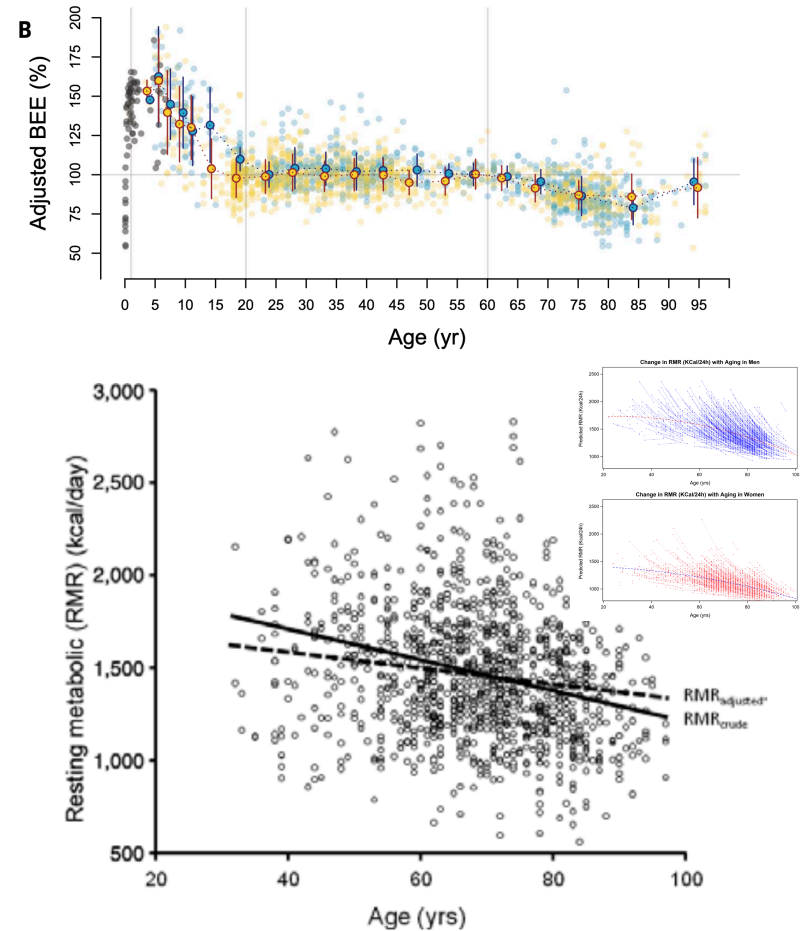


Energetic paradox ?

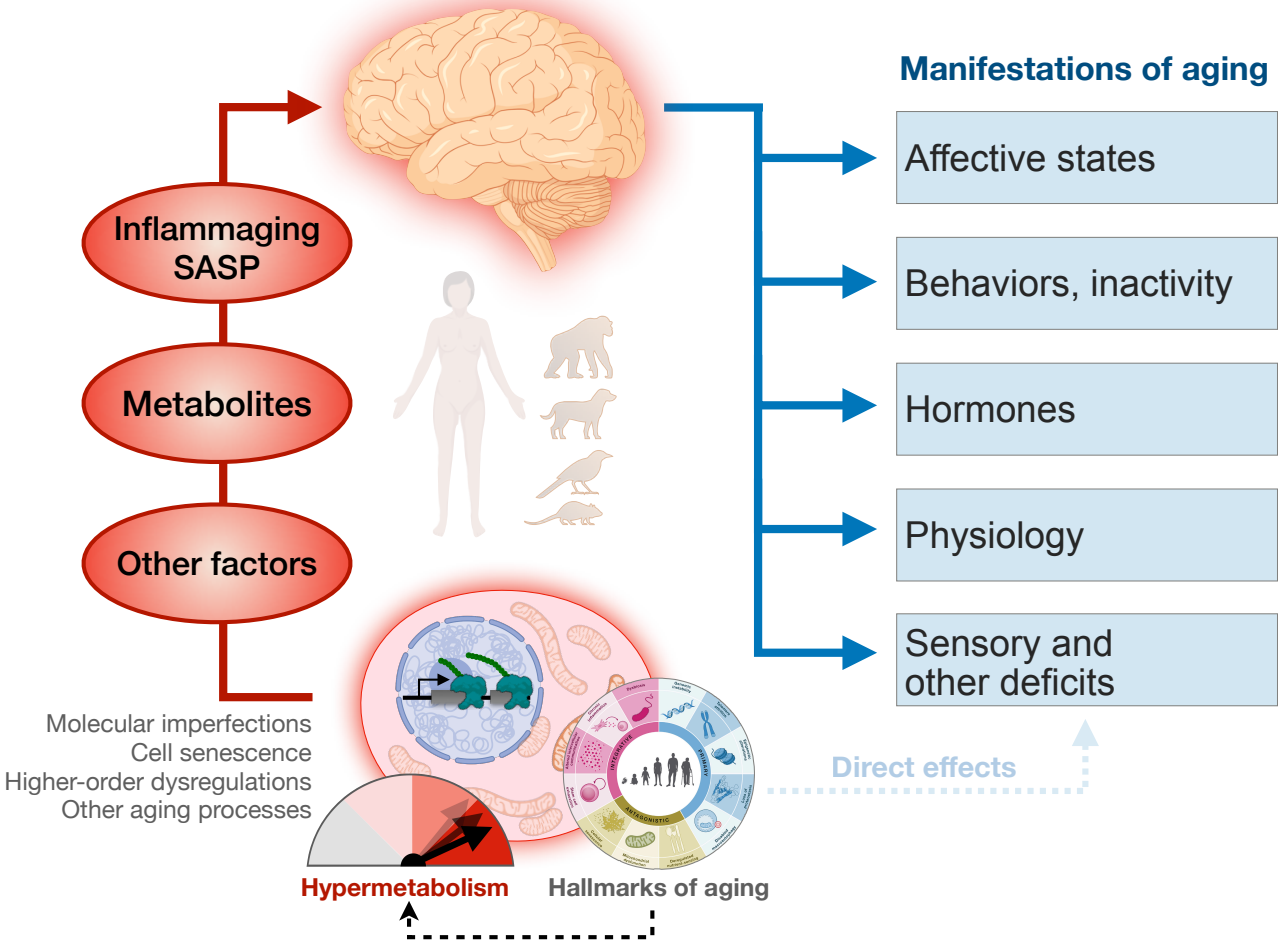
CELLULAR HYPERMETABOLISM



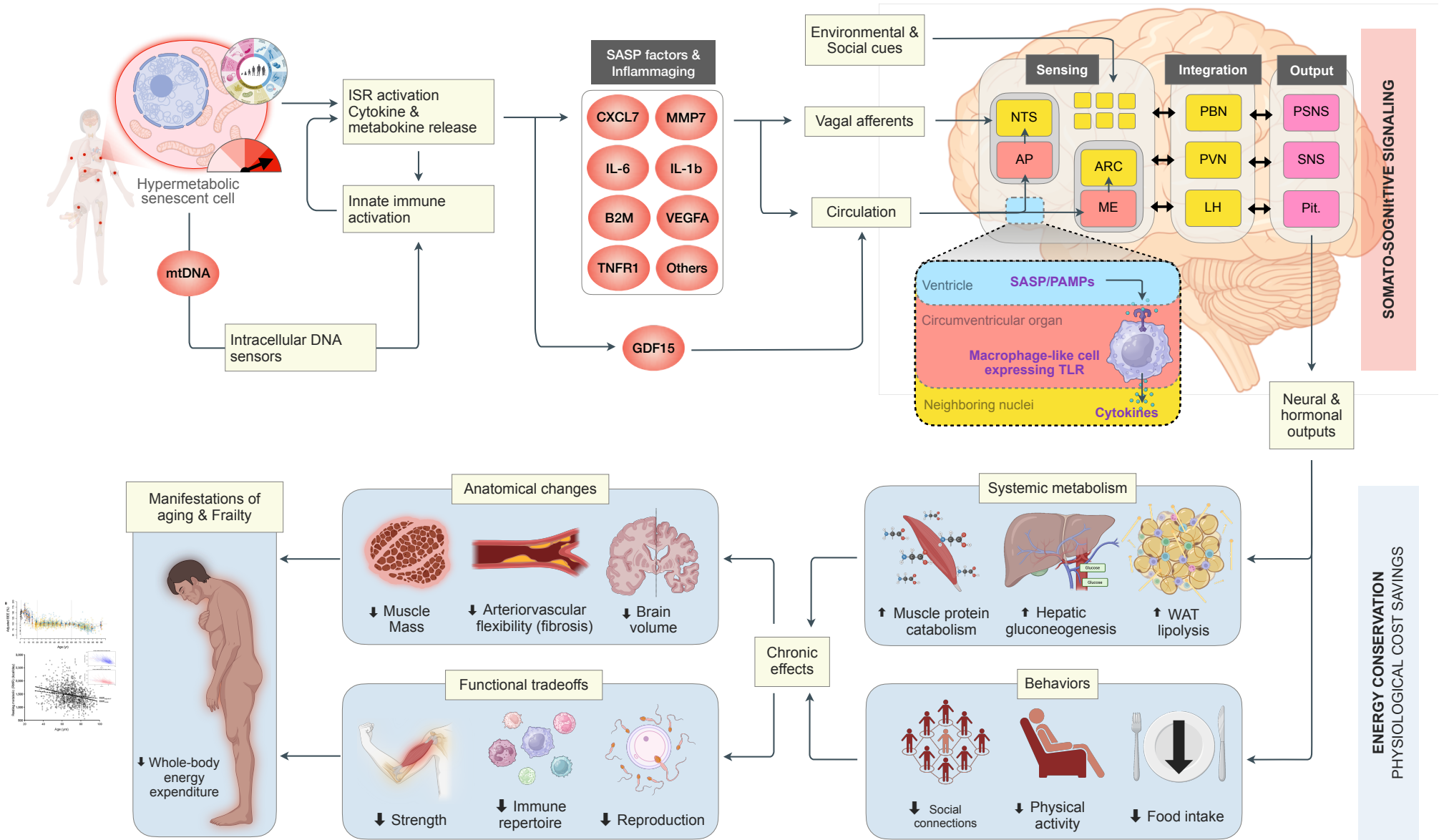
WHOLE-BODY HYPOMETABOLISM



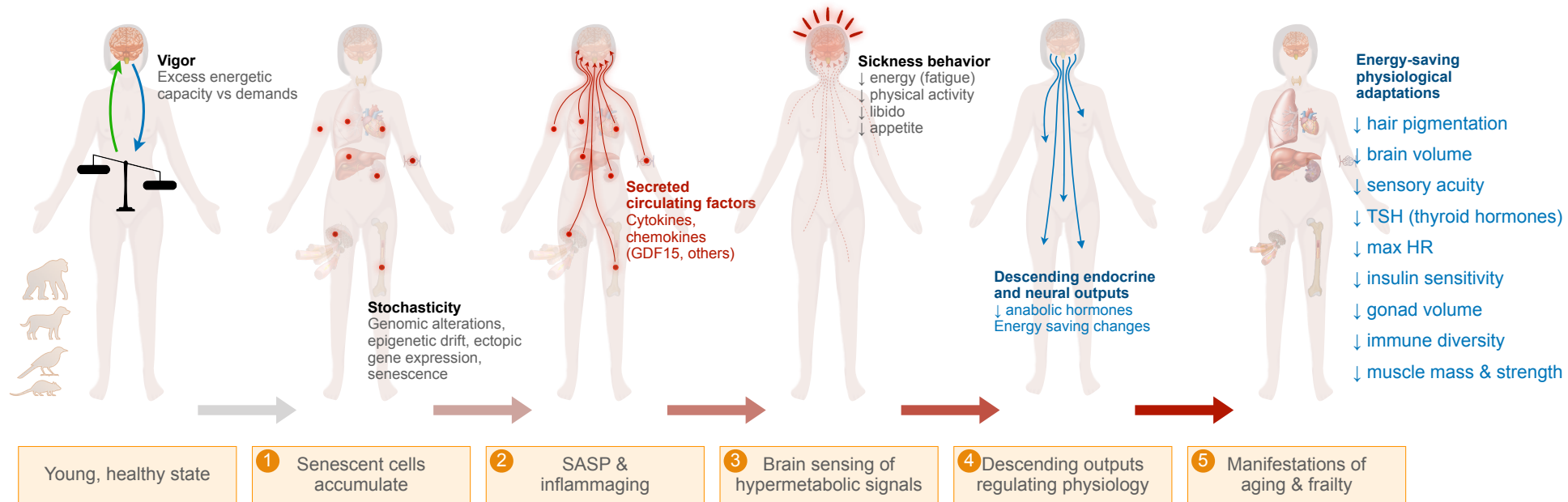
Somato-cognitive Energy Conservation (SEC)



Somato-cognitive energy conservation (SEC)

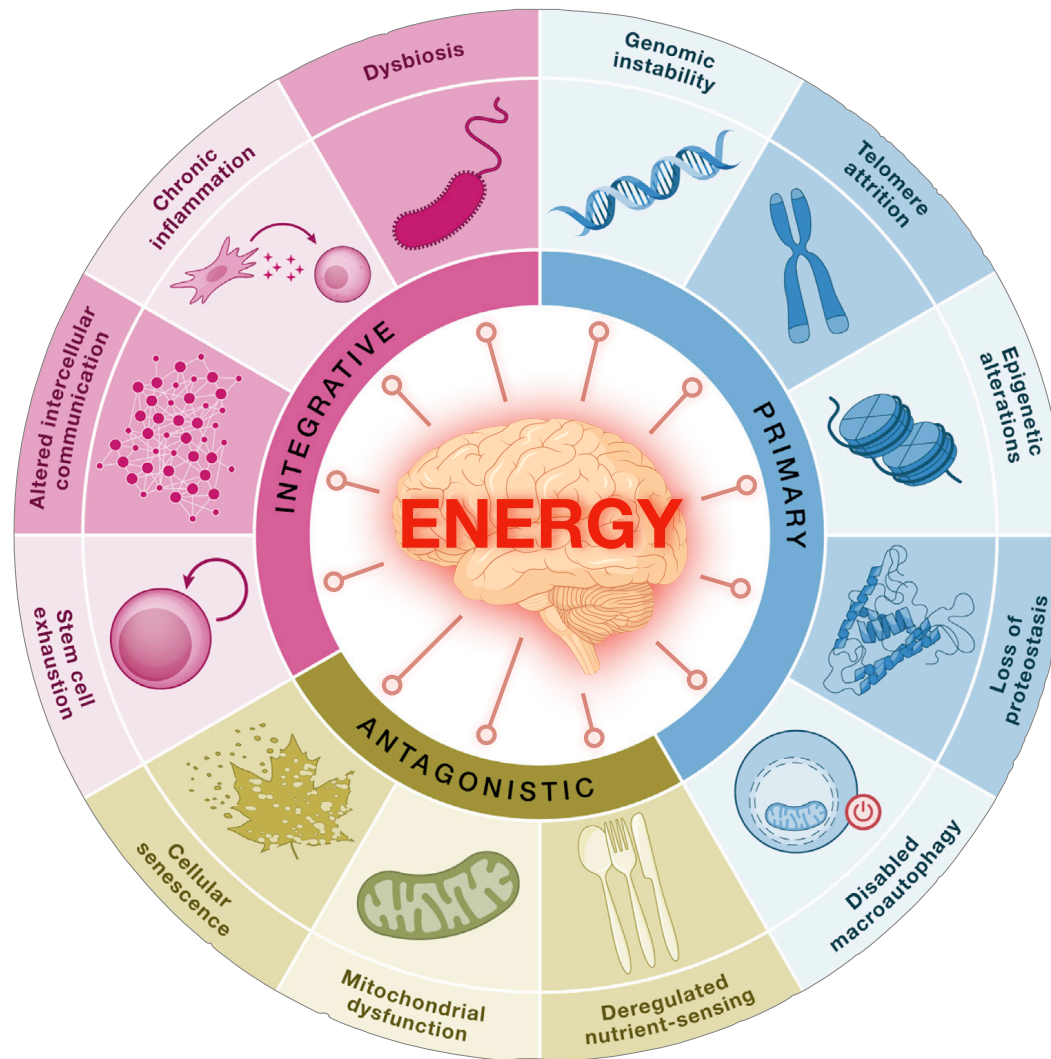


Somato-cognitive energy conservation (SEC)

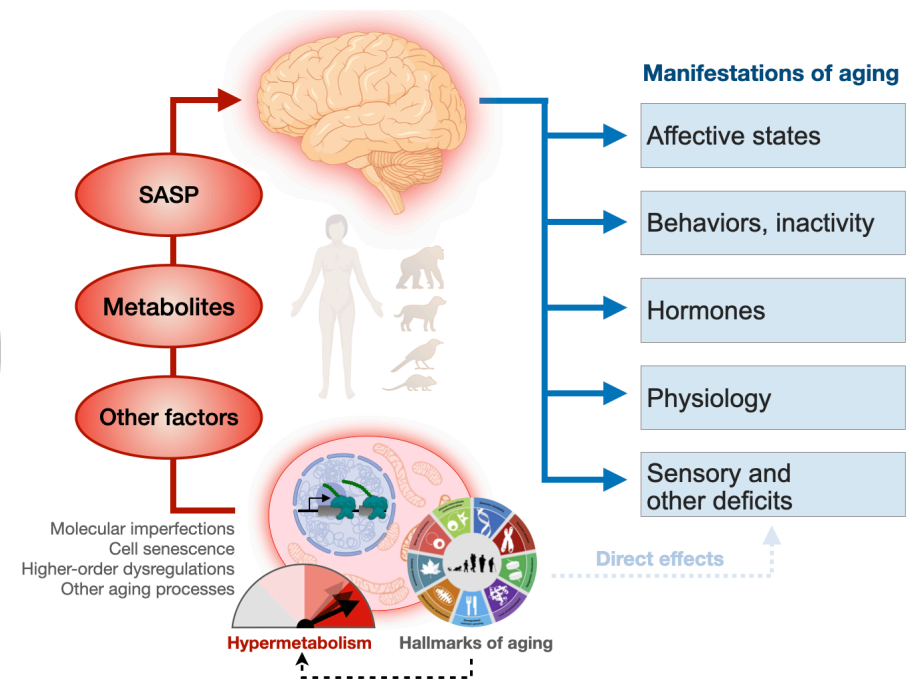
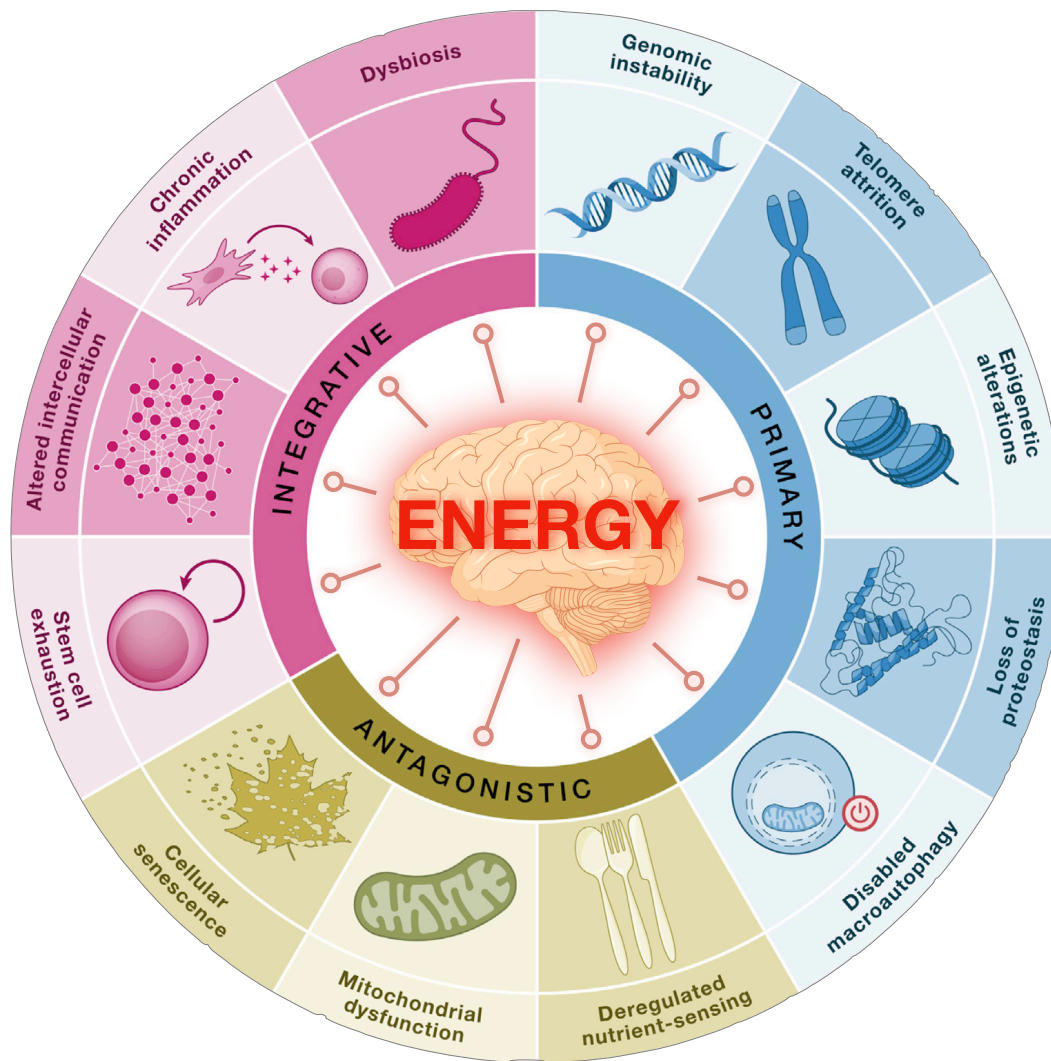


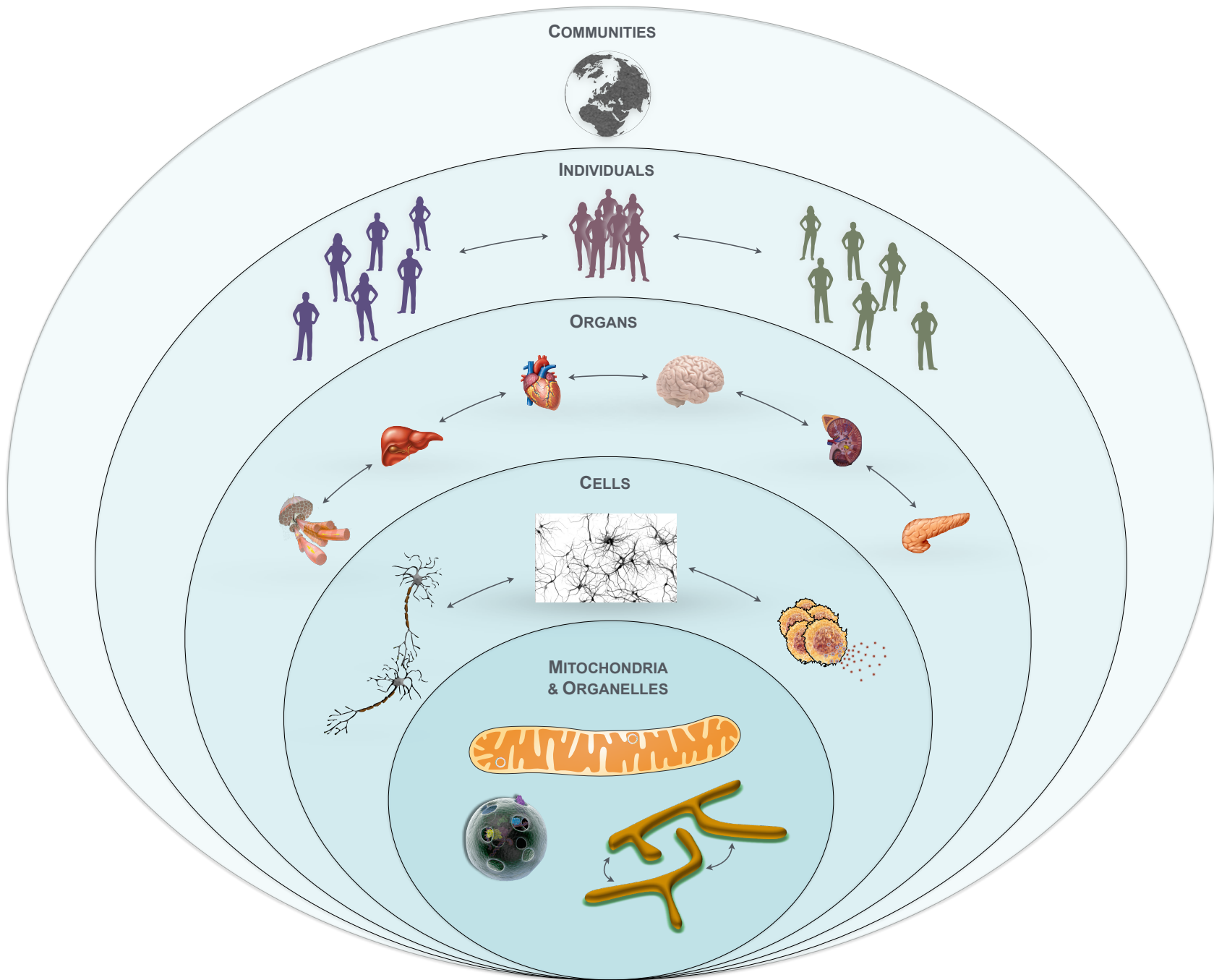
Is the shared factor among hallmarks of aging **ENERGY** ?

Do the hallmarks physically intersect in the **BRAIN** ?

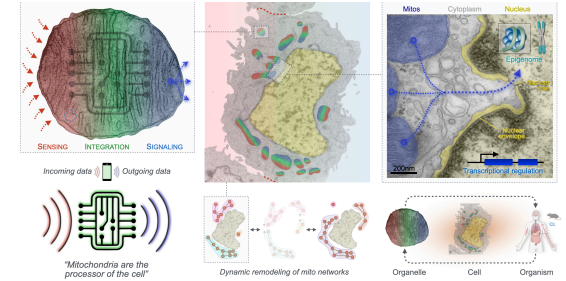
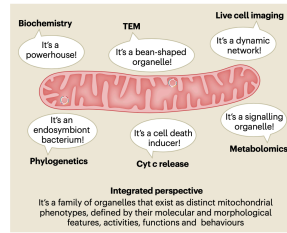
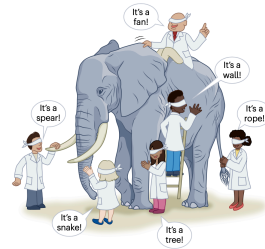
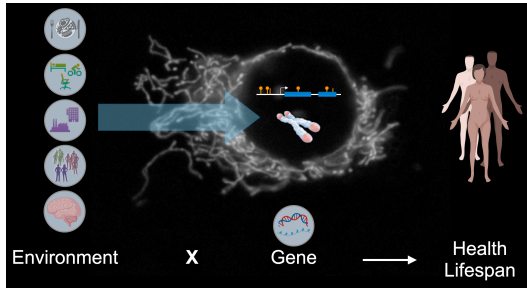


Somato-cognitive energy conservation (SEC)

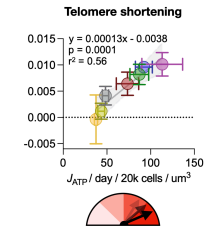
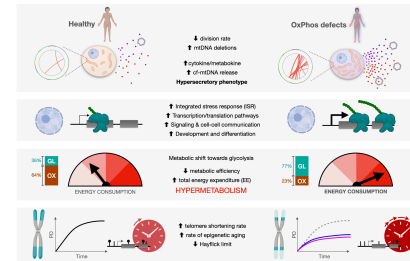
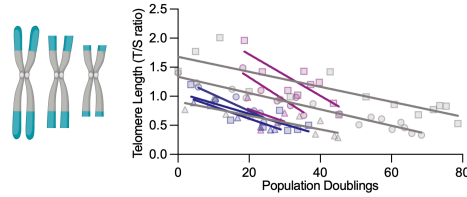
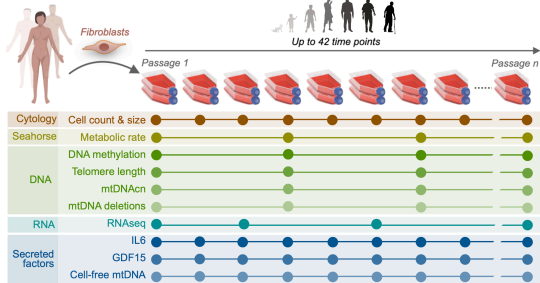




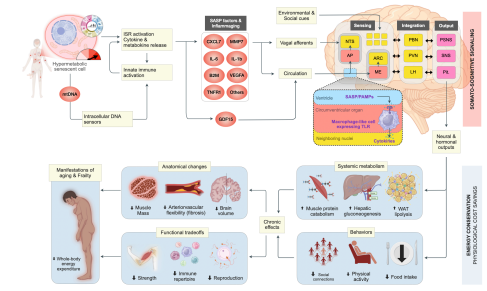
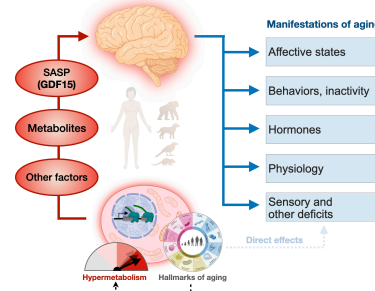
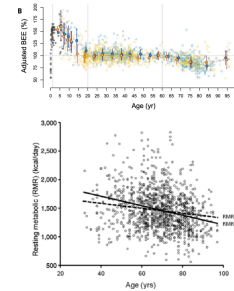
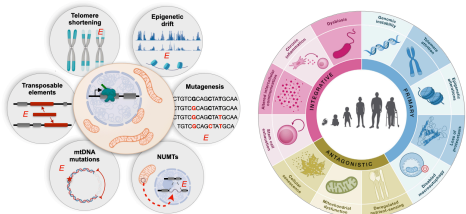
PART 1



PART 2



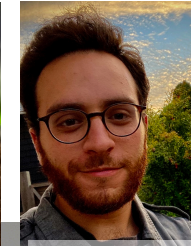
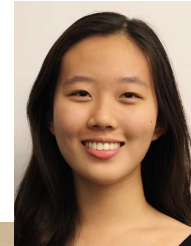
PART 3



Mitochondrial PsychoBiology Lab

Linking molecular processes within mitochondria with the human experience

OUR RESEARCH



Evan

Gabriel



Caroline

Hannah

Natalia

Anna

Collaborators

Mitochondrial Biology & Medicine

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Catarina Quinzii
CUIMC Neurology
- Brett Kaufman
Pittsburgh University
- Gyuri Hajnóczy
Erin Seifert
Thomas Jefferson University
- Orian Shirihai
Mike Irwin
UCLA
- Vamsi Mootha
Rohit Sharma
Harvard & MGH
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University of Michigan
- Gilles Gousspillou
UQAM
- Tonio Enriques
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- Catherine Kelly
- Shufang Li
Anna Monzel
Mangesh Kurade

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- Jue Lin
Aric Prather
Ashley Mason
UCSF
- Eli Puterman
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- Clemens Kirshbaum
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Vincenzo Lauriola
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Manish Saggarr
Stanford

Anne Grunewald
University of Luxembourg

Carmen Sandi
EPFL

Efrat Levy
Pasquale D'acunzo
NYU

Biological Aging

- Steve Horvath
- Morgan Levine
Altos
- Albert Higgins-Chen
Yale
- Marie-Abèle Bind
Harvard
- Luigi Ferrucci
NIA Intramural
- Alan Cohen
- Dan Belsky
Linda Fried
CUIMC Mailman & Aging Center

BASZUCKI
BRAIN RESEARCH FUND

The Nathaniel Wharton Fund 

 National Institute of Mental Health

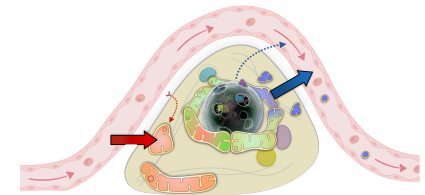
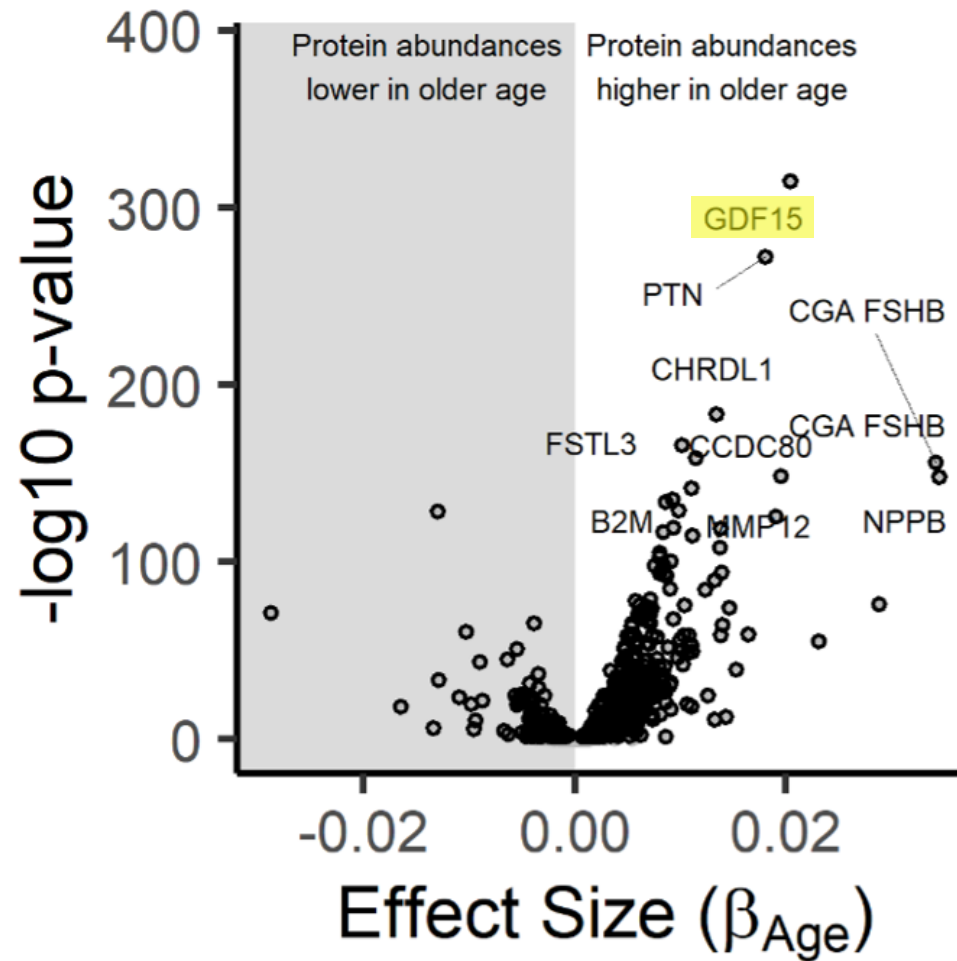
 National Institute of General Medical Sciences

 National Institute on Aging

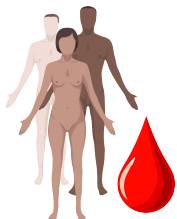
Downloadable presentation slides

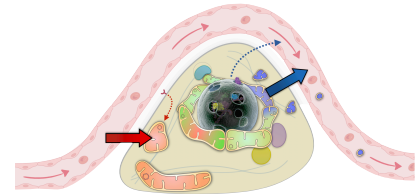
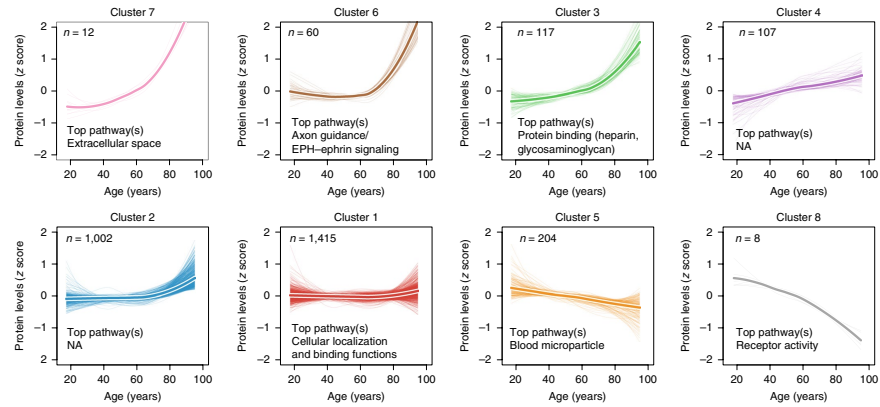
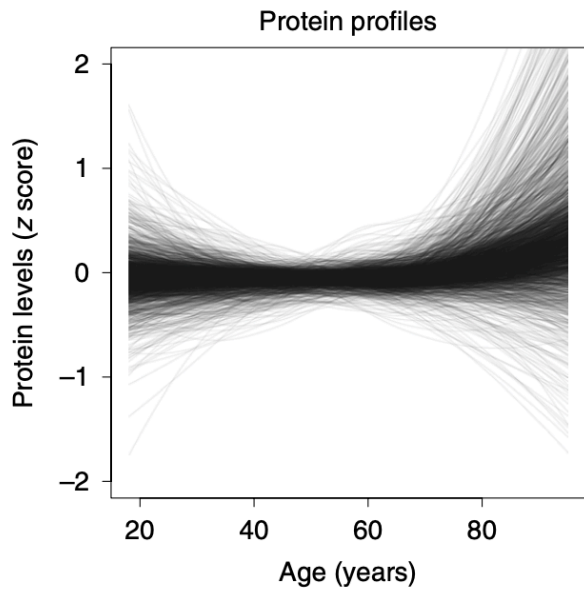


UPregulation of most signaling proteins in human aging

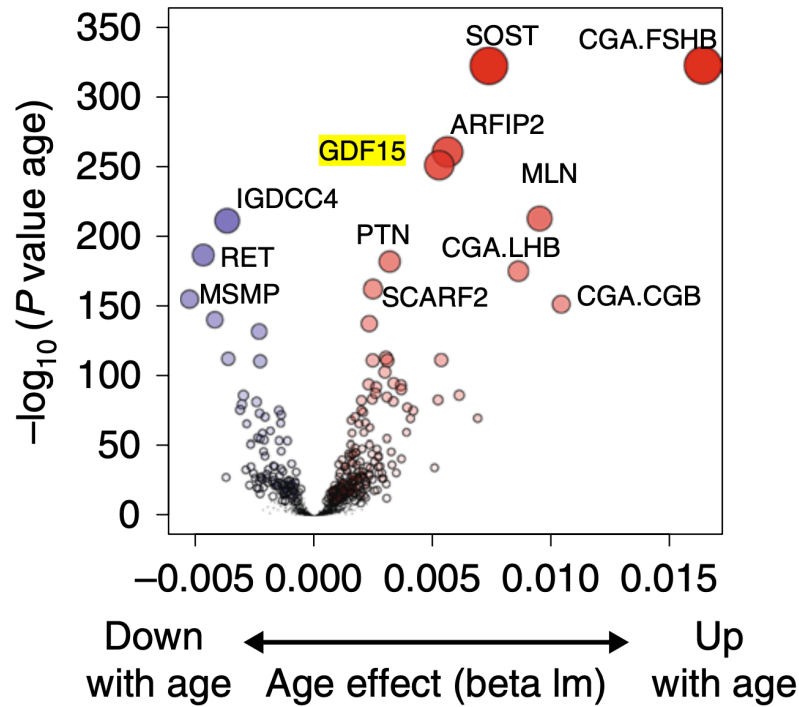


**Null hypothesis:
50:50**

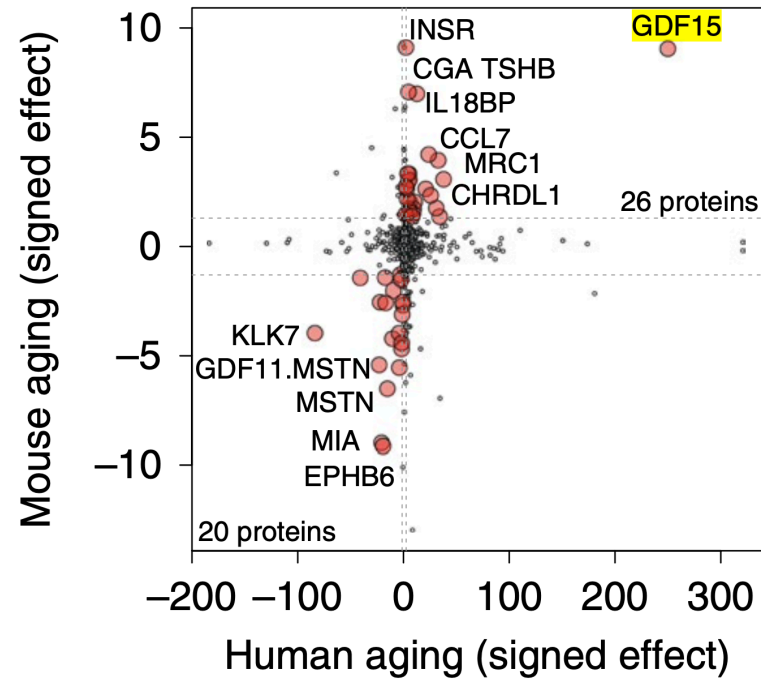




Aging proteome

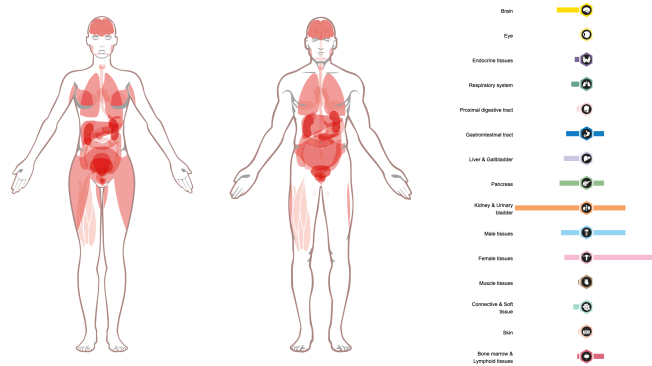


Conserved aging proteome

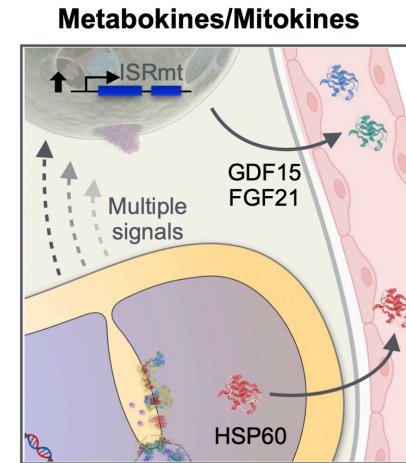


What does GDF15 mean to the organism?

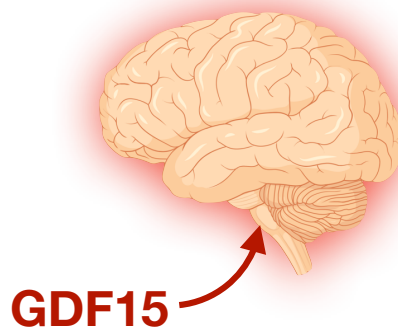
What does GDF15 mean to the organism?



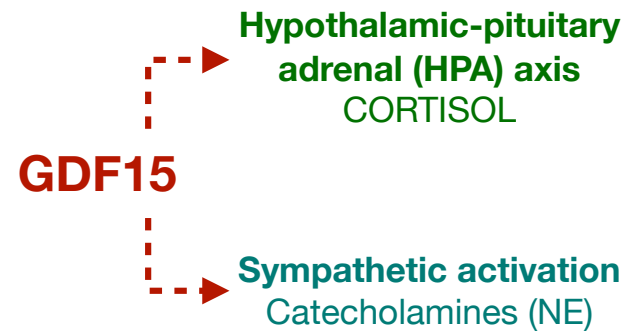
Expressed in >50% of somatic tissues



Triggered by cellular stressors (ISR)

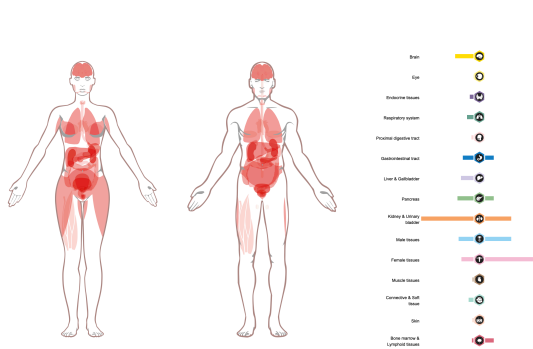


Signals on the brainstem

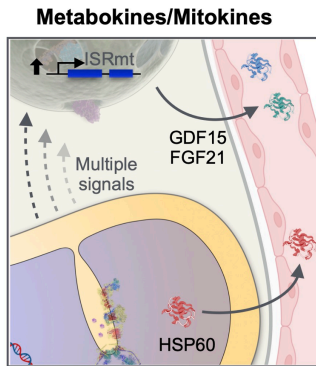


Activates canonical stress axes

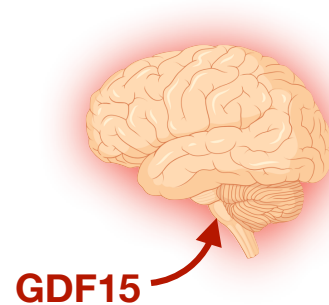
What does GDF15 mean to the organism?



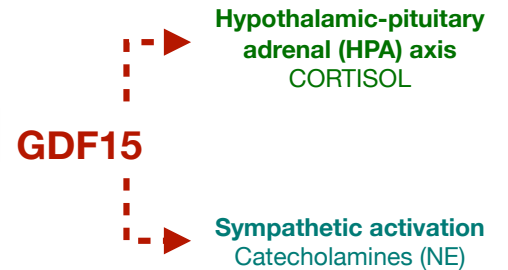
Expressed in >50% somatic tissues



Triggered by cellular stressors (ISR)

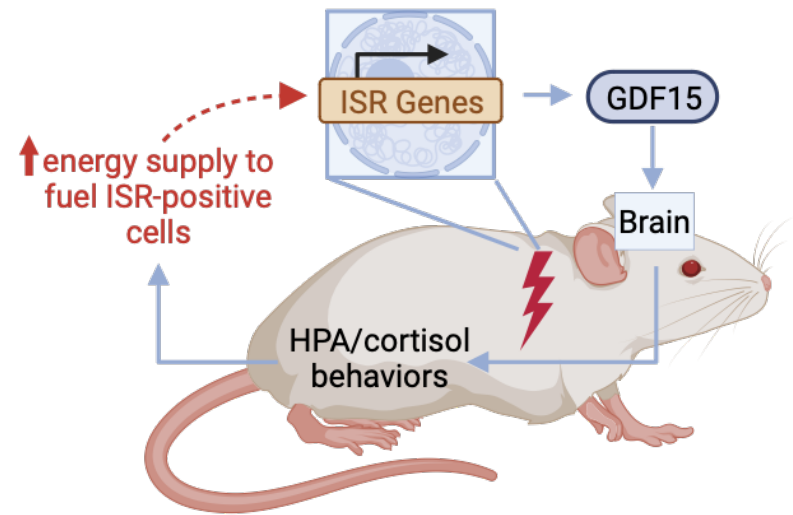


Signals on the brainstem

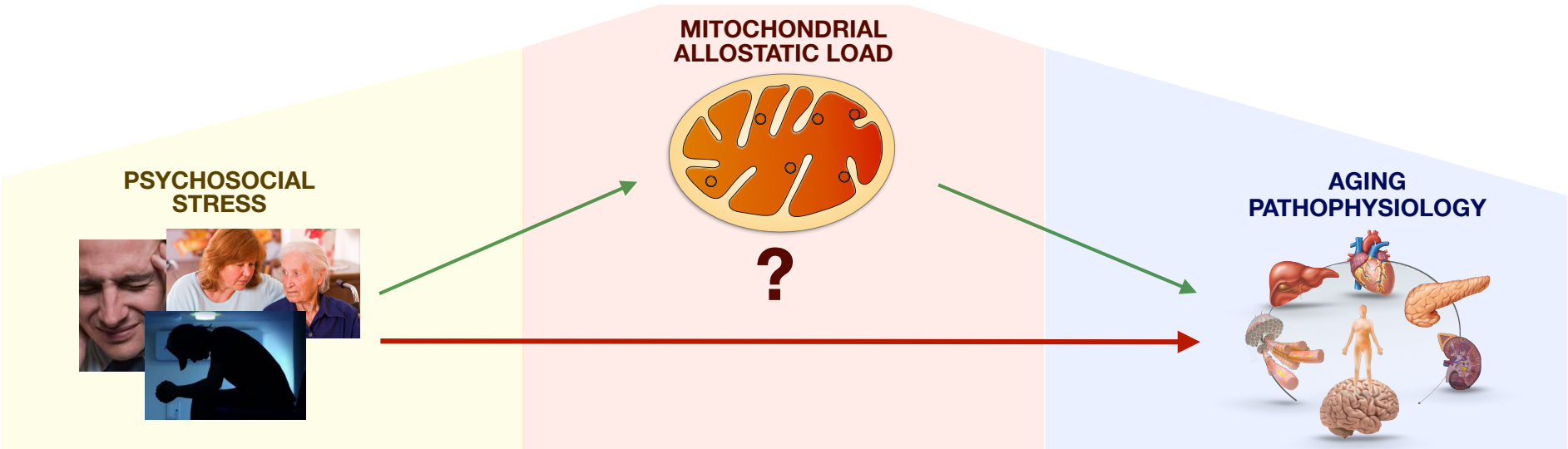
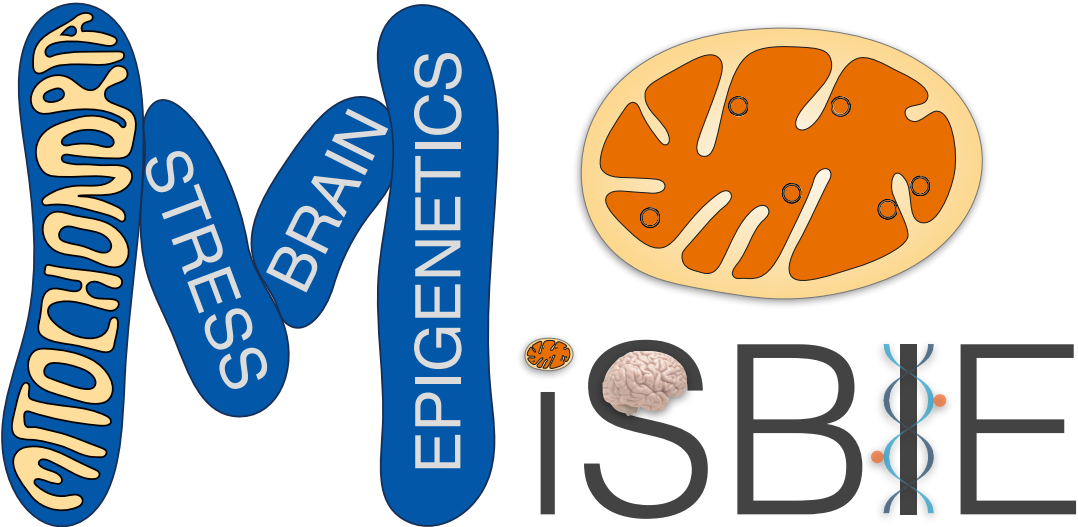


Activates canonical stress axes

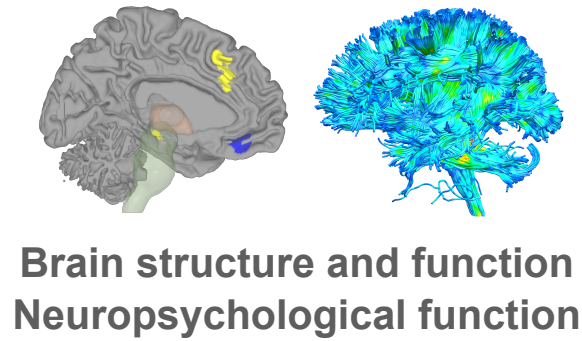
Organismal body-brain signaling



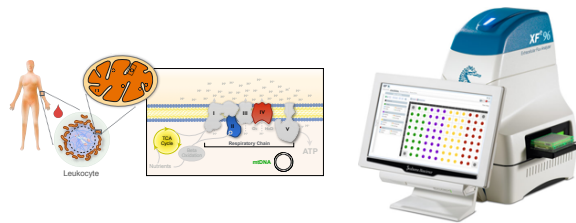
Mitochondrial Stress, Brain Imaging, and Epigenetics — MiSBIE



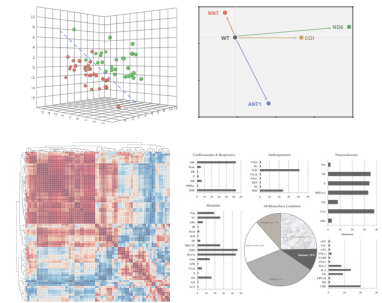
Mitochondrial Stress, Brain Imaging, and Epigenetics — MiSBIE



mtDNA heteroplasmy
Mitochondrial OxPhos
Lymphocytes, Monocytes,
Neutrophils, Platelets

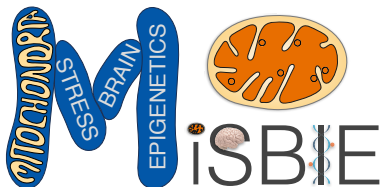


Disease biomarkers
Stress reactivity
Energy expenditure
e.g., GDF15, Lactate,
Pyruvate, Alanine, etc.



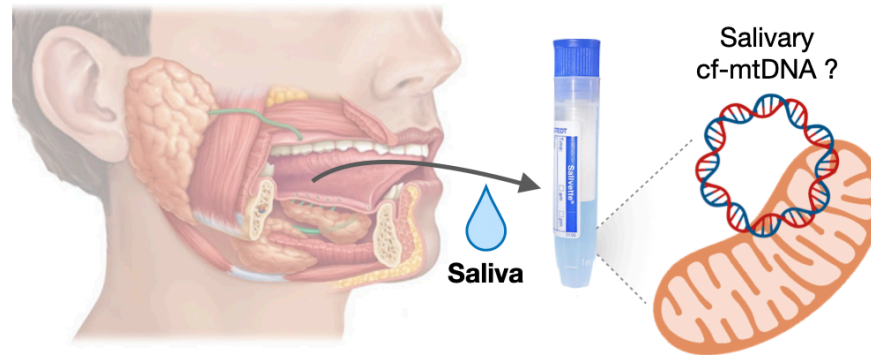
Total N = 110

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



Stress hormones & metabolites are detectable in saliva

Can we quantify cf-mtDNA and other “mitokines”
in human saliva?



If so: this would make possible epidemiological and
high-temporal resolution timecourse studies of GDF15

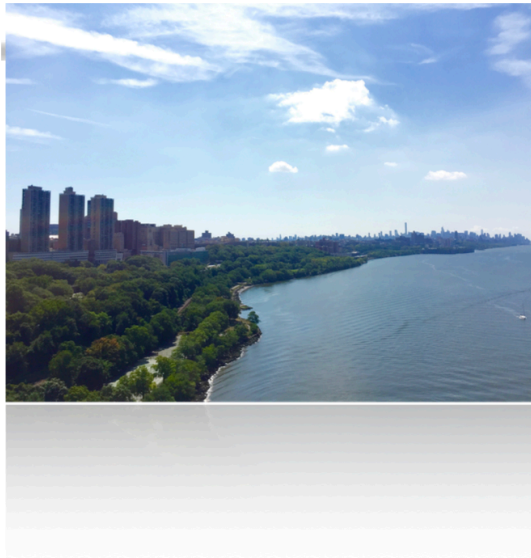


Shannon Rausser Caroline Trumpff

The MiSBIE Study

Mitochondrial Stress Brain Imaging and Epigenetics

Investigating the link between the mind and the body



 COLUMBIA UNIVERSITY
IN THE CITY OF NEW YORK



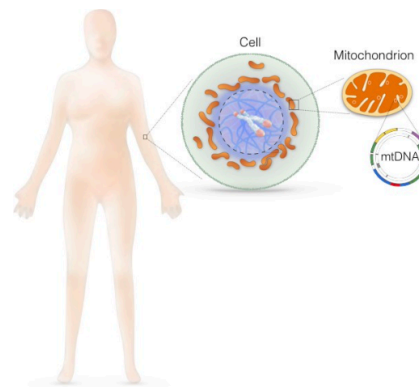
Understanding Mitochondrial Disease

Researchers are just starting to understand the factors that influence aging and the progression of various diseases. Life stress can change the function of the body and influence the development of certain age-related diseases, such as cardiovascular disease and neurodegeneration.

The goal of the MiSBIE study is to understand how an individual's life experience and emotions affect physical health, psychological functioning, and disease risk.

Each cell of the body contains hundreds of mitochondria, which have their own DNA: mitochondrial DNA (mtDNA). Mitochondria produce energy and signals enabling cells to function normally. The MiSBIE study investigates the link between mitochondria, brain function, and different organs to understand their interaction, and *the person as a whole*.

This study also aims to understand the behavior of genes, whether they are turned “on” or “off”. This is called “epigenetics” and is measured in DNA from different cells.



Columbia University Medical Center

The MiSBIE study is a research study taking place at the College of Physicians and Surgeons at the **Columbia University Medical Center (CUMC)**, a leading medical institution of care and research.

The partnering Department of Neurology and Department of Psychiatry have a long history of clinical care and research in studying the effects of stress on the body and in mitochondrial disease.

CUMC is located at 168th Street and Broadway in Upper Manhattan, by the Hudson River in New York City, NY.



 COLUMBIA UNIVERSITY
MEDICAL CENTER

The MiSBIE Study

A two-day visit

This research includes two visits of about 8 hours each. Participants stay overnight at a nearby hotel.

Breakfast and lunch are provided.

Transport

Would you need to travel to NYC? If so, the MiSBIE Team will arrange your travel and reimburse your expenses associated with the study.

Confidentiality

All results and biological samples are kept **strictly confidential**.

Compensation

Participants who complete the study receive a compensation of \$599.

Eligibility

You are eligible if you are a woman or man between the ages of 18 and 60, and willing to visit Columbia University Medical Center (CUMC) for a two-day visit.

We are recruiting individuals with the following mtDNA mutations:

- m.3243A>G (MELAS)
- Single, large scale deletion (CPEO)

The MiSBIE Team

The MiSBIE team is a group of caring clinicians and researchers from academic disciplines including mitochondrial medicine, physiology, neuroscience, epigenetics, and psychology.

For more information about the study, please contact the clinical coordinator (see contact information on the back).

Sponsors



Contact Information

Questions about the study? Interested to participate?

Catherine Kelly | Study Coordinator

MiSBIE@columbia.edu

646-774-8931

Kris Engelstad | Clinical Coordinator

ke4@cumc.columbia.edu

212-342-5767



Study tests

On Day 1, participants undergo a laboratory evaluation. Saliva and blood samples, and a small clip of hair are collected. Heart rate and blood pressure are monitored. Participants also complete a medical exam with a doctor.

On Day 2, magnetic resonance imaging (MRI, *picture above*) is used to safely measure brain activity, participants complete questionnaires on an iPad, and meet with a neuropsychologist.

Participants also collect saliva at home.

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clinical and translational research

The MiSBIE Team

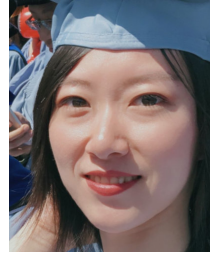
Laboratory and Clinical



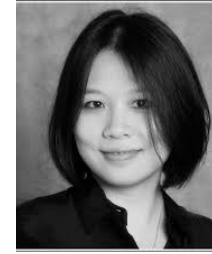
Catherine Kelly



Kris Engelstad



Shufang Li



Grace Liu



Lea Gregorio



Mangesh Kurade



Anna Monzel



Jeremy Michelson



Natalia Bobba-Alves



Janell Smith



Cynthia Liu



Alex Junker



Jack Baker



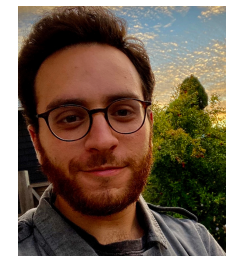
Vincenzo Lauriola



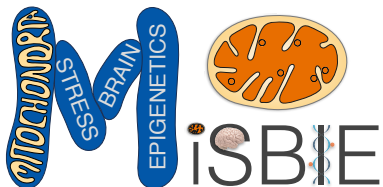
Sophia Tepler



Sophie Basarrate



Gabriel Sturm



The MiSBIE Team

Core Investigators



Frances Champagne
UT Austin



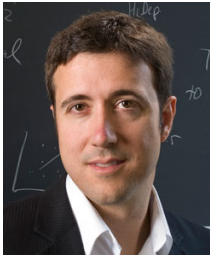
Stephanie Assuras
Columbia



Caroline Trumpff
Columbia



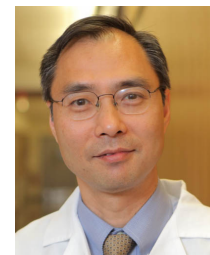
Richard Sloan
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Tor Wager
Dartmouth



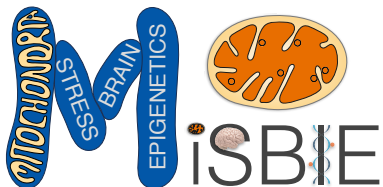
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