



Perspectives on Cancer & Aging

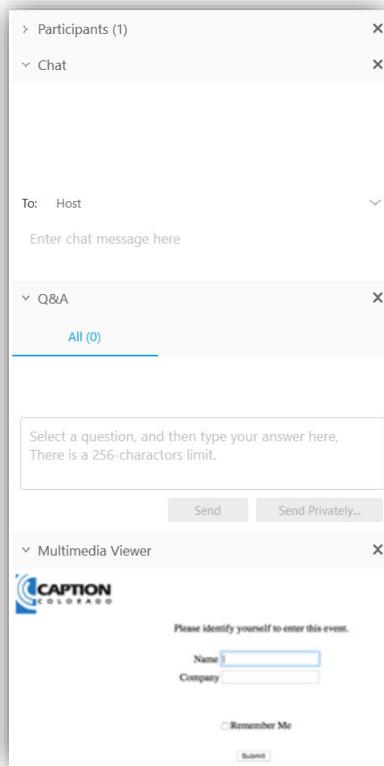
ARTI HURRIA MEMORIAL WEBINAR SERIES

Cancer and Aging: Biological and Phenotypic Measures of Aging

A WEBINAR TRIBUTE TO DR. ARTI HURRIA

Luigi Ferrucci, M.D., Ph.D.
Morgan Levine, Ph.D.
Cancer & Aging Research Group

Using WebEx and webinar logistics

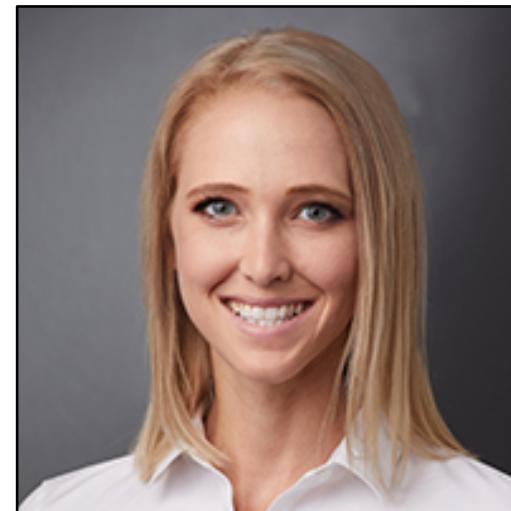


- All lines will be in listen-only mode
- Make sure icons are selected for them to appear as a drop-down option
- Submit questions at any time during the presentation by typing into the Q&A feature on the right-hand side of the WebEx interface.
 - Select Host and a moderator will ask the questions on your behalf
- Closed captioning available by selecting the Media Viewer Panel on the right-hand side of the screen
- This webinar is being recorded

Today's speakers:



Luigi Ferrucci, M.D., Ph.D.
Geriatrician and Epidemiologist
Scientific Director
National Institute on Aging



Morgan Levine, Ph.D.
Assistant Professor
Department of Pathology
Yale School of Medicine

April 9th 2020
Perspectives on Cancer and Aging Webinar

“Connecting the biological and phenotypic manifestations of aging: the case of muscle aging”.

Luigi Ferrucci - National Institute on Aging



The Metrics of Aging

Functional Aging (*impact on daily life*)

- Cognitive Function
- Physical Function
- Mood
- Mental Health



Phenotypic Aging (*phenotypes that change*)

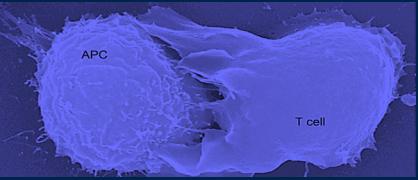
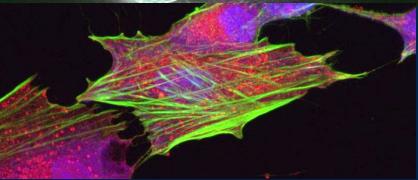
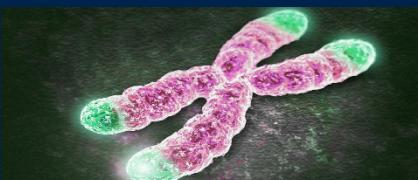
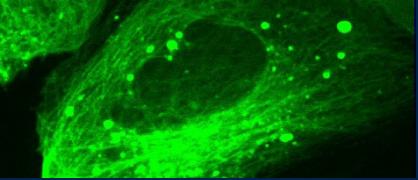
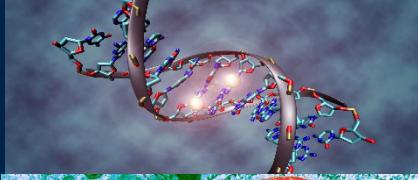
- Body Composition
- Energetics
- Homeostatic Mechanisms
- Brain health



Biological Aging (*root mechanisms*)

- Molecular damage
- Defective repair
- Energy exhaustion
- Signal/noise reduction





Genomic Instability

The Accumulation of Somatic Mutations with Aging

Cellular Senescence

Trade-off Between Cancer and Aging

Epigenetics (methylation)

The “Epigenetic Clock”

Mitochondrial Dysfunction

The Power Plant

Proteostasis (autophagy)

Repair, Recycle or Trash?

Telomere Length

Protecting the DNA During Replication

Stem Cell Exhaustion

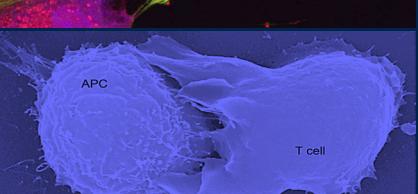
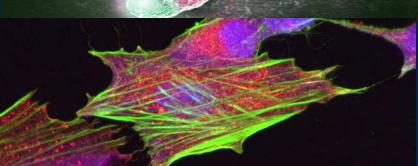
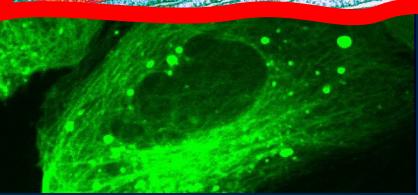
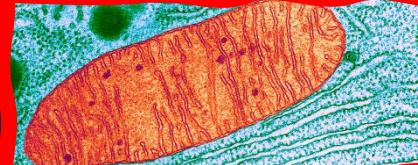
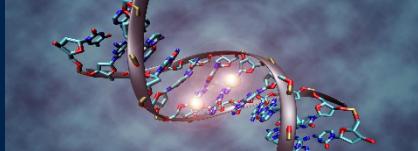
Templates for Cells Restoration

Cell to Cell Communication

Accuracy and Context in the Flow of Information



The Hallmarks of Aging
Carlos Lopez-Otin et al.



Genomic Instability

The Accumulation of Somatic Mutations with Aging

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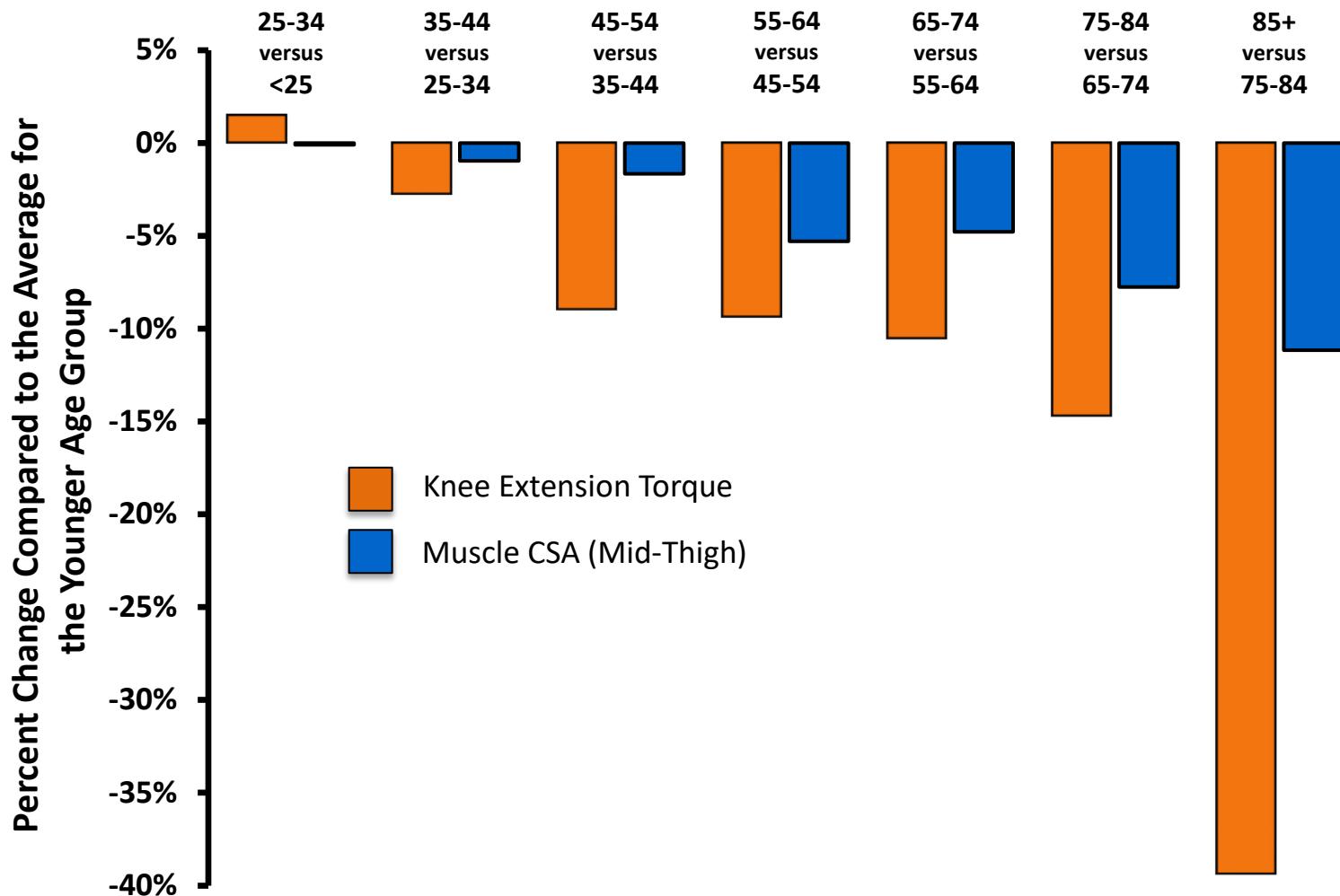
Accuracy and Context in the Flow of Information



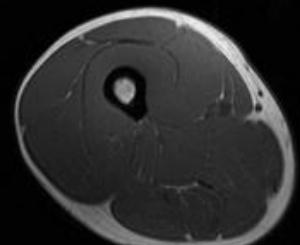
The Hallmarks of Aging
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Muscle

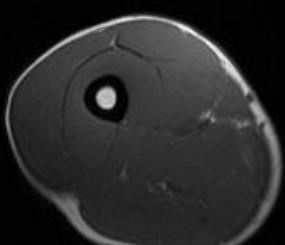
Strength/ Mass Ratio in BLSA Participants 60-70 yrs Old



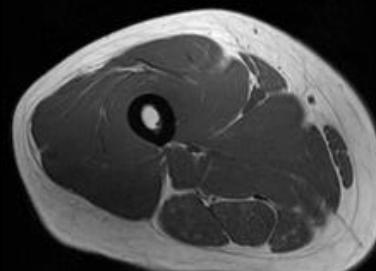
Mid-thigh T1w MRI Images (Men; GESTALT)



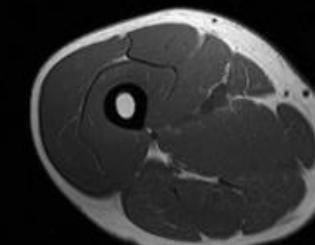
Age 23 Years



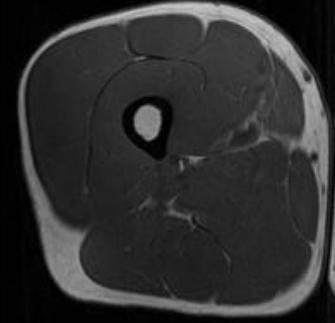
Age 26 Years



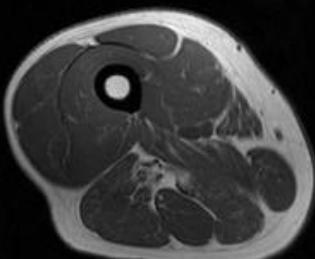
Age 28 Years



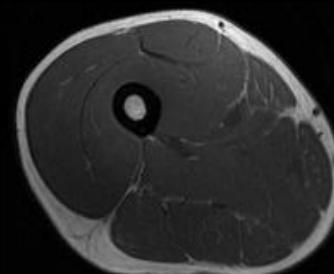
Age 31 Years



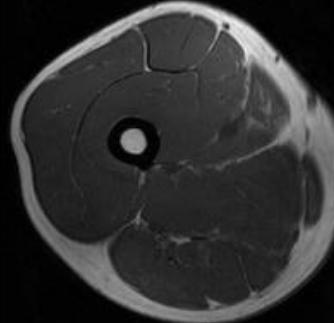
Age 31 Years



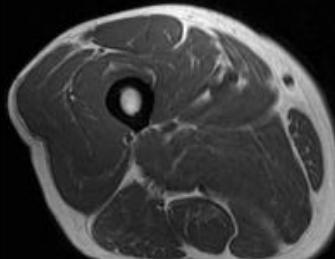
Age 42 Years



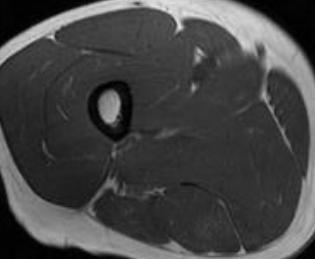
Age 45 Years



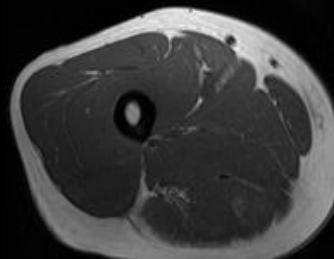
Age 45 Years



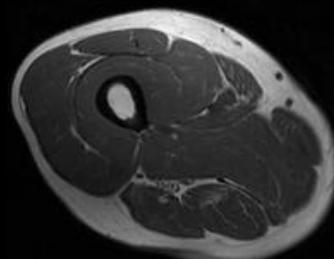
Age 51 Years



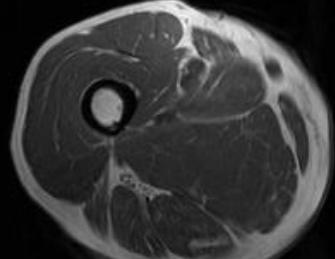
Age 52 Years



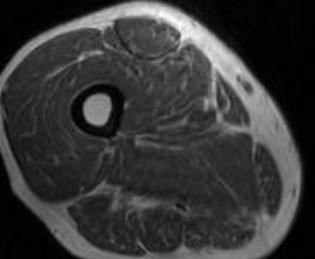
Age 57 Years



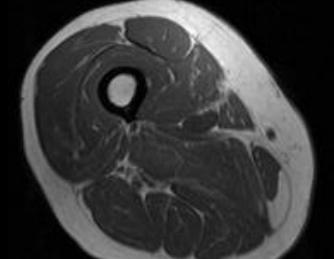
Age 60 Years



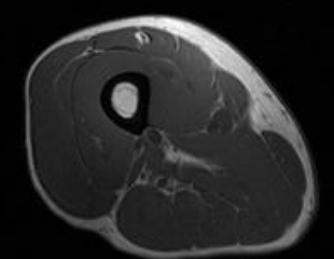
Age 67 Years



Age 72 Years



Age 81 Years



Age 83 Years



David Reiter

NESTED CASE-CONTROL STUDY in BLSA

Selection of 79 pairs of cases (low muscle quality) and controls (high muscle quality), matched by age (± 2.5 years), sex, and height (± 1.5 cm). Muscle quality defined as knee extension torque/mid-thigh muscle cross-sectional area.

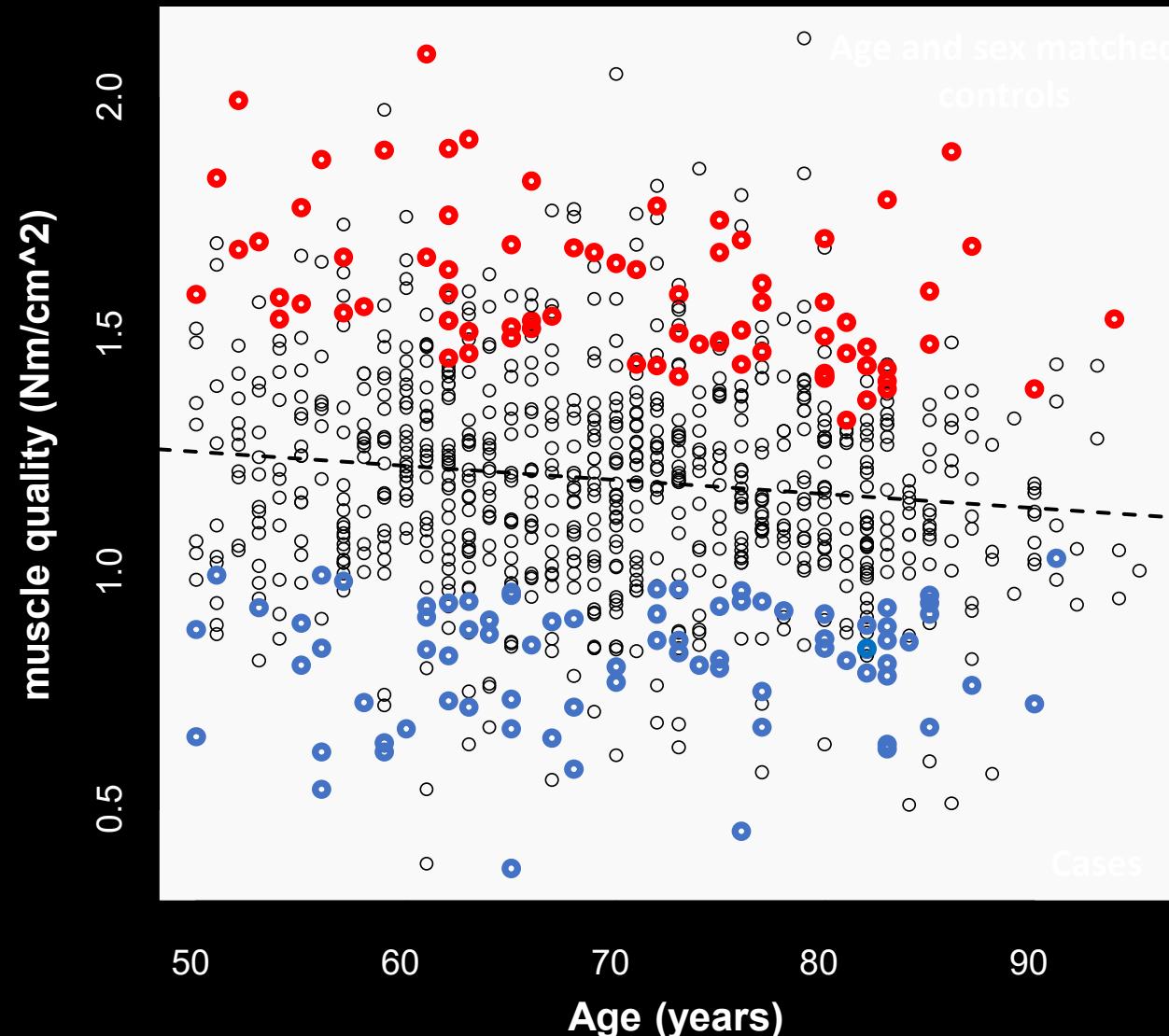
Moaddel R et al. J Gerontol A Biol Sci Med Sci. March 2016. doi:10.1093/gerona/glw046



Ruin Moaddel



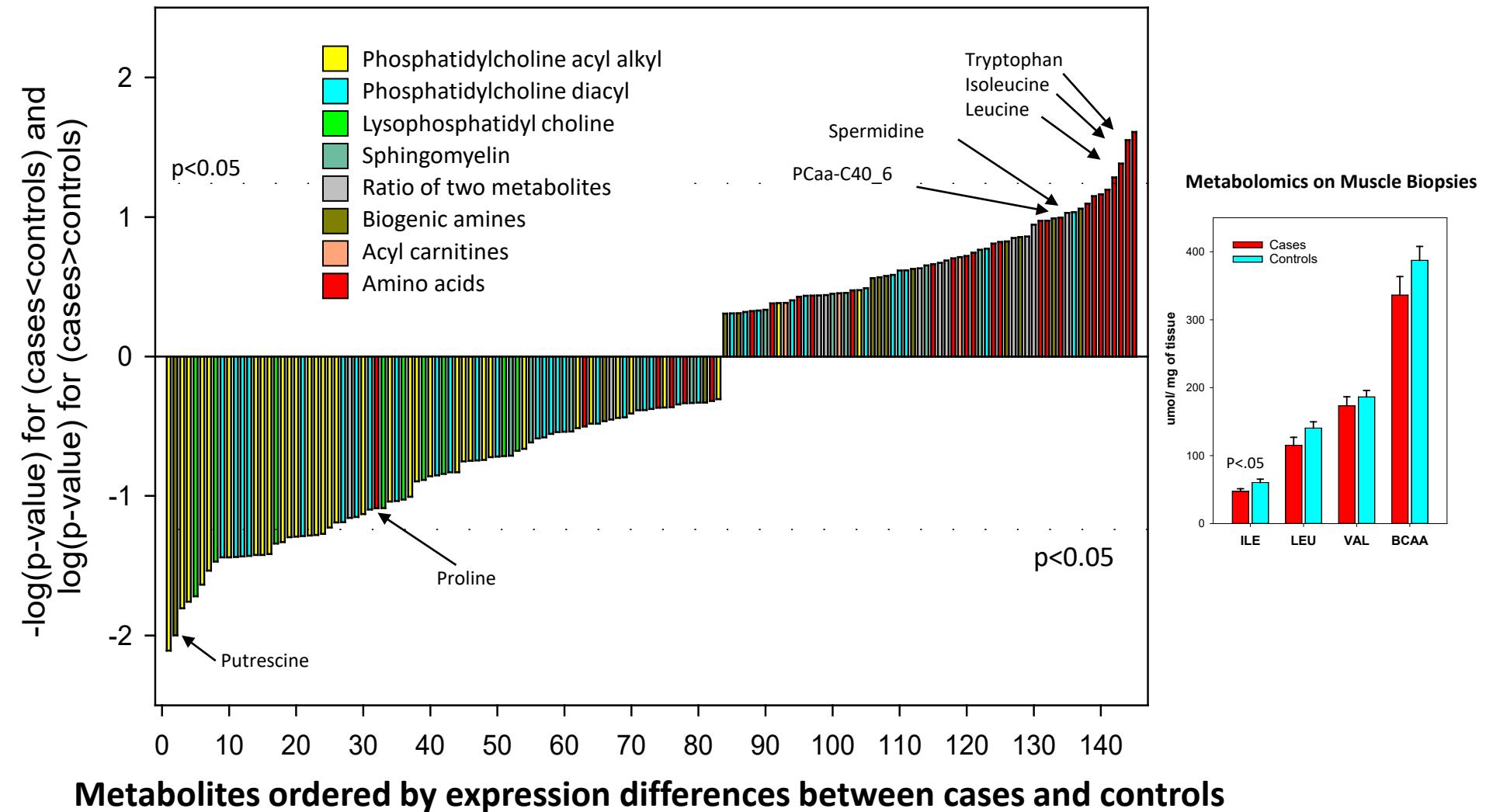
Elisa Fabbri



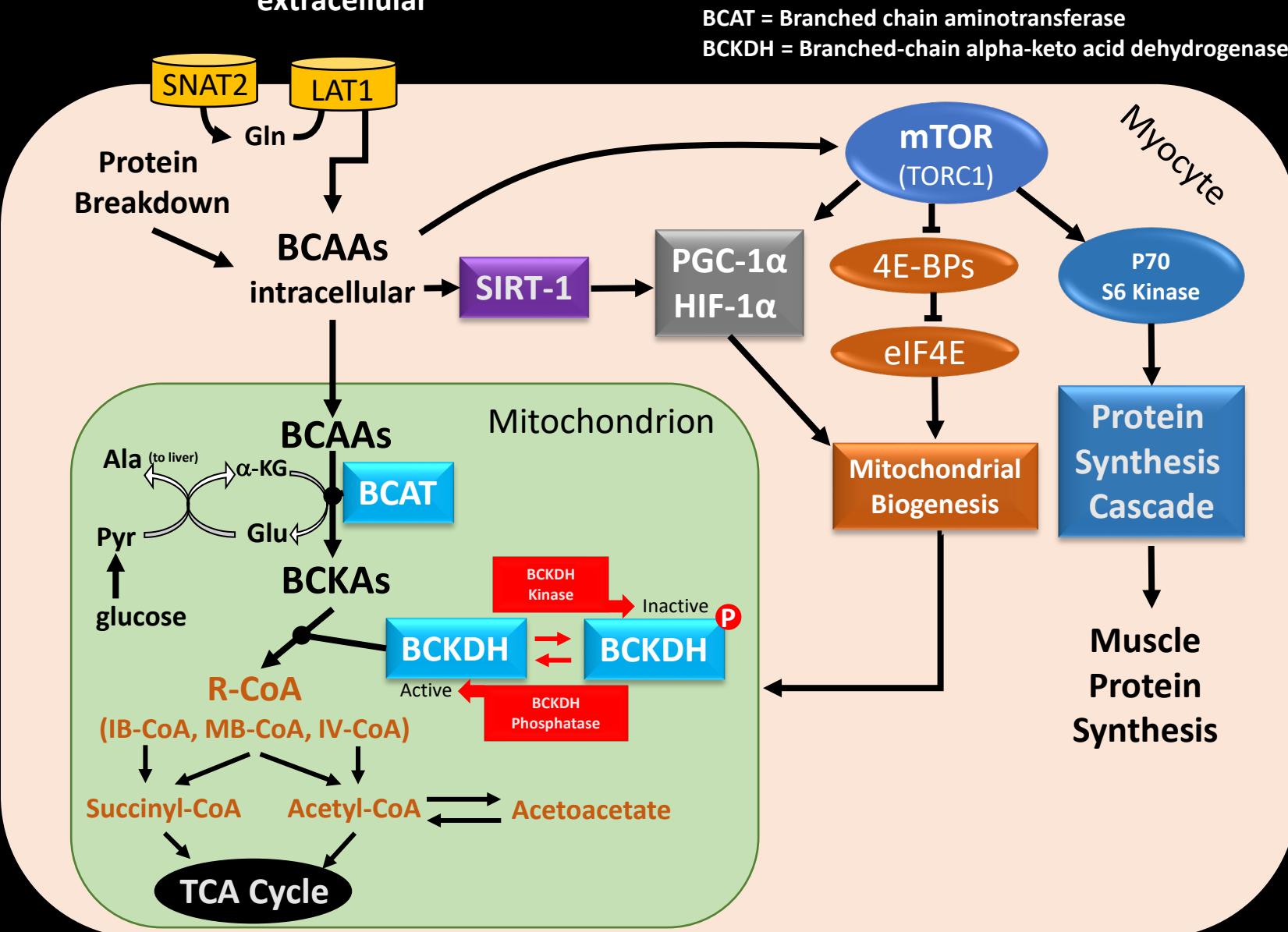
NESTED CASE-CONTROL STUDY in BLSA

126 Metabolites according to down-regulation or up-regulation in cases (low muscle quality) compared to controls (high muscle quality)

Moaddel R et al. J Gerontol A Biol Sci Med Sci. March 2016. doi:10.1093/gerona/glw046



BCAAs stimulates energy production and protein synthesis



KPCr by Age in Men and Women

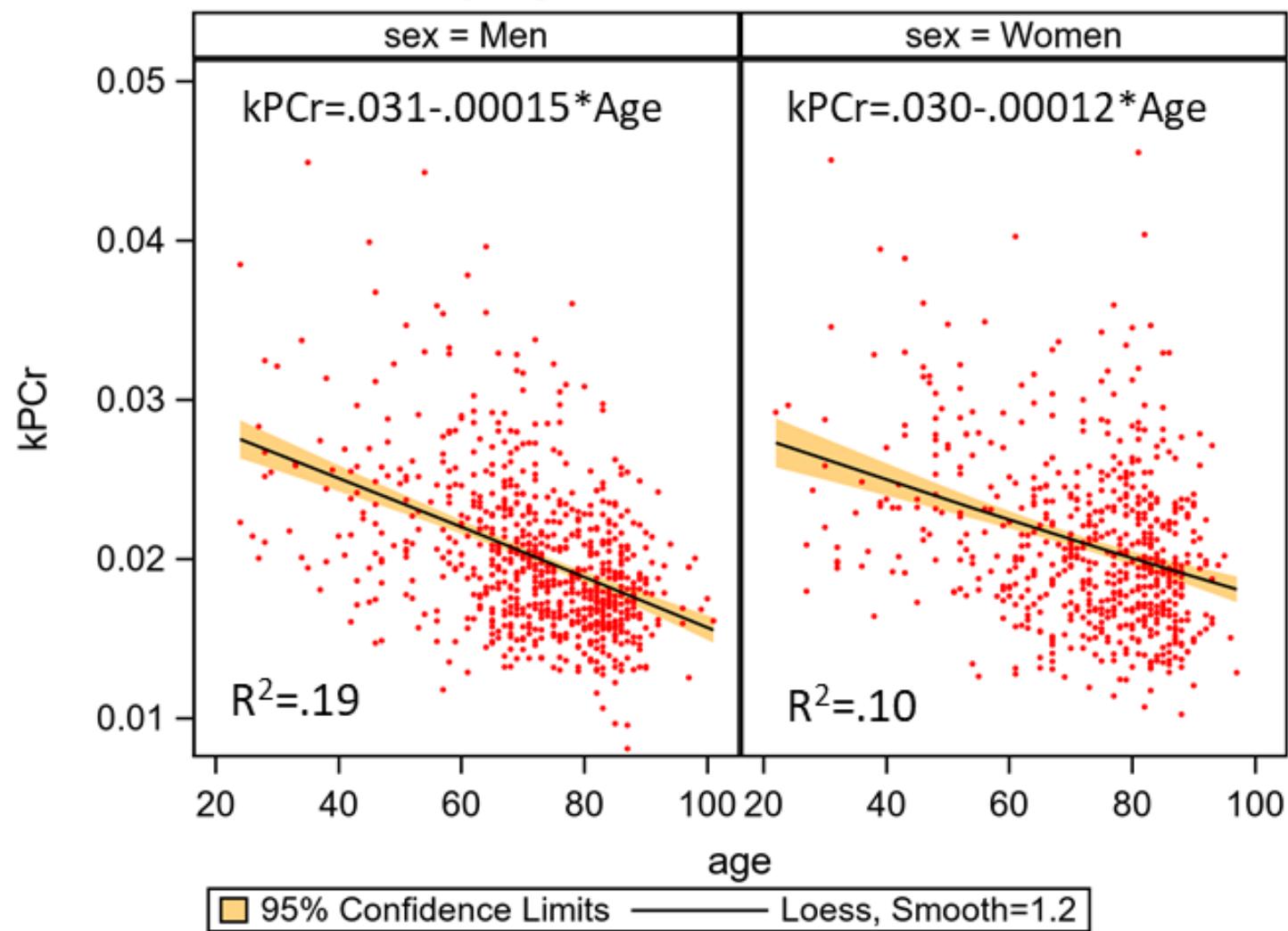
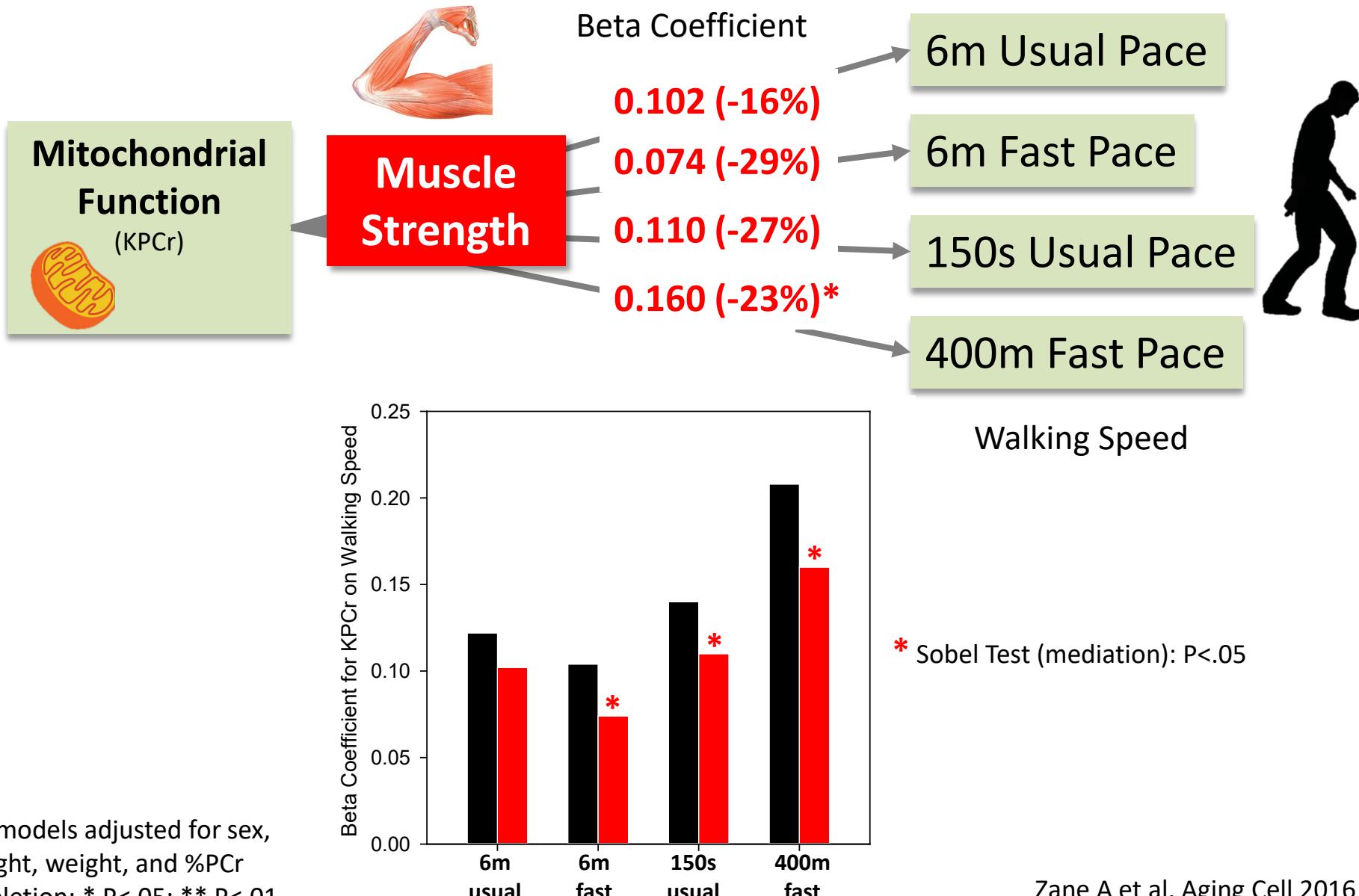


Figure 1. Skeletal muscle oxidative capacity, a proxy marker of mitochondrial function, declines with aging both in men (n=400) and women (n=331). BLSA 2020 limited to first measures.

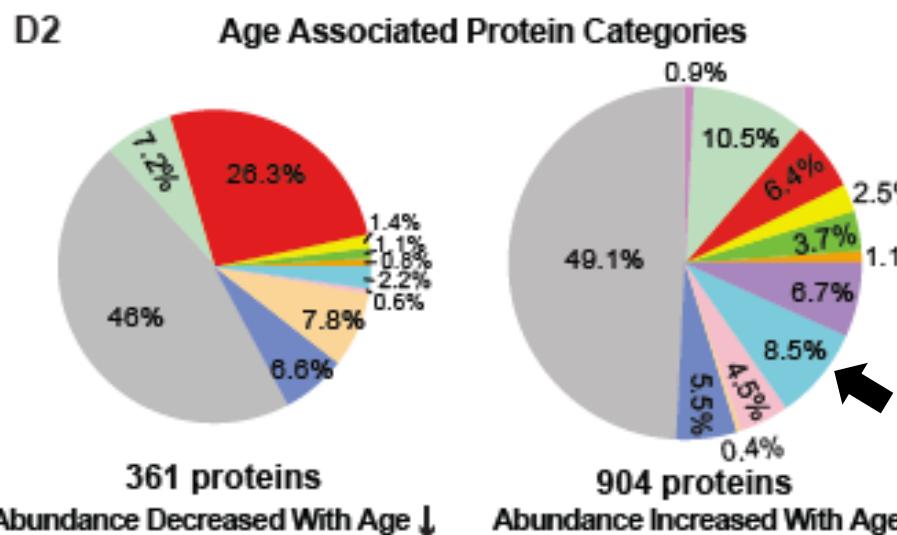
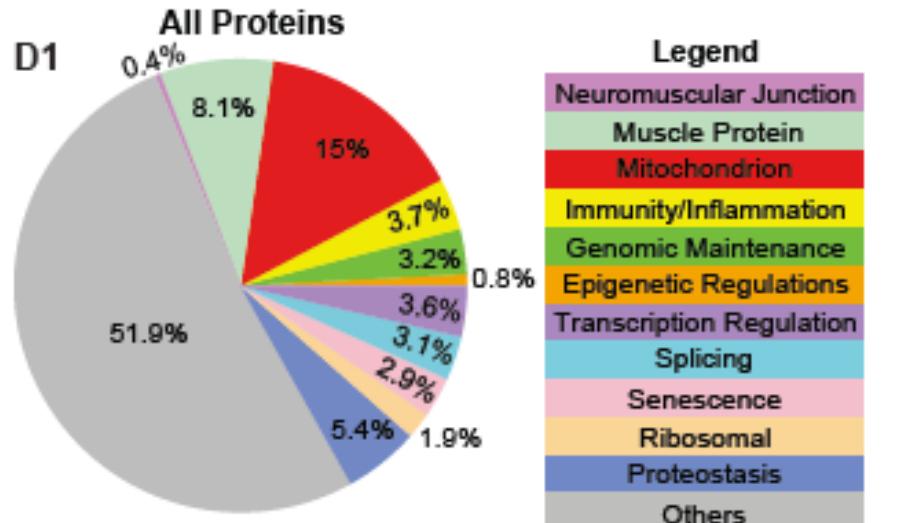
Muscle Strength Mediates the Effect of Mitochondrial Function on Walking Performance.





Classification of Age-associated Proteins In Skeletal Muscle

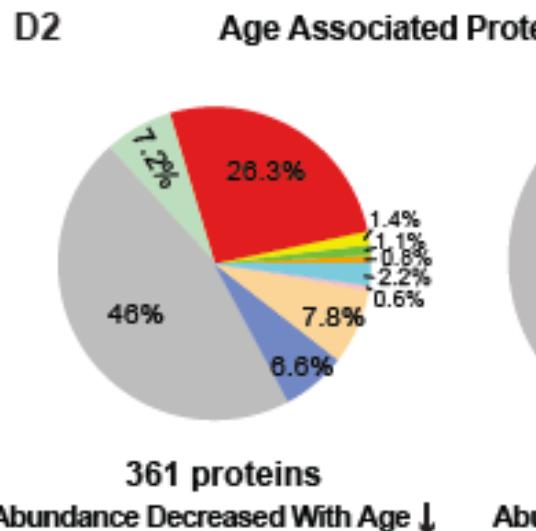
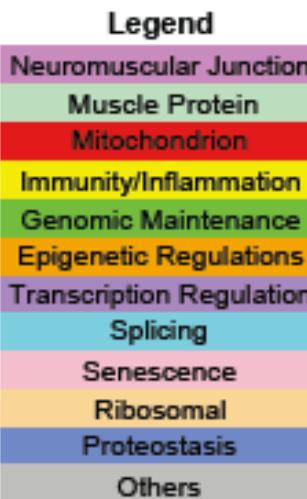
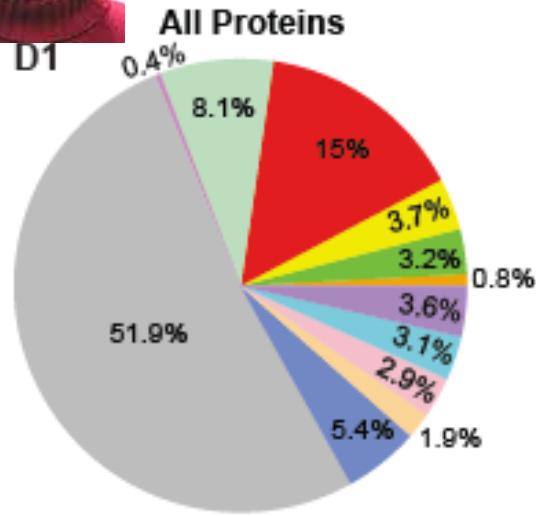
Proteins associated with Aging



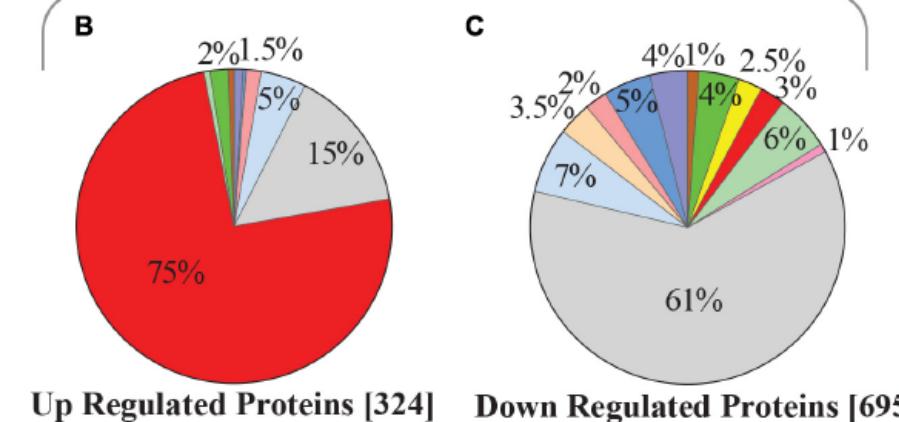
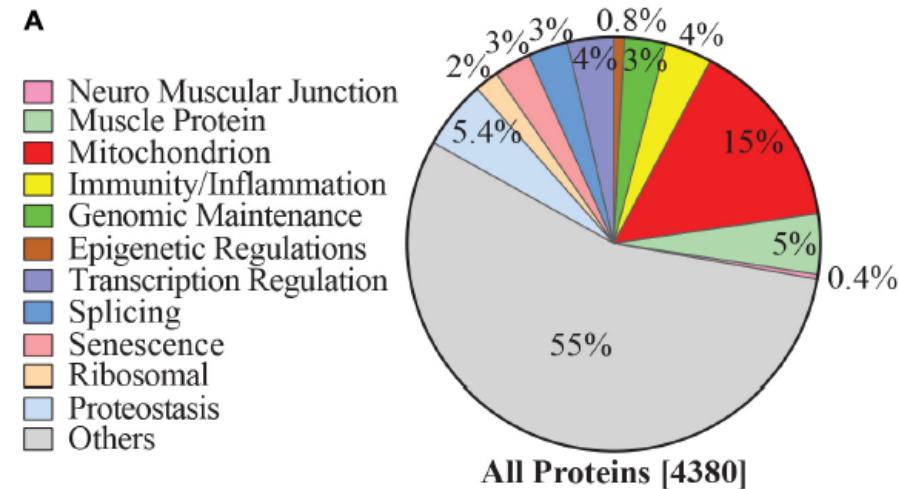


Classification of Age-associated Proteins In Skeletal Muscle

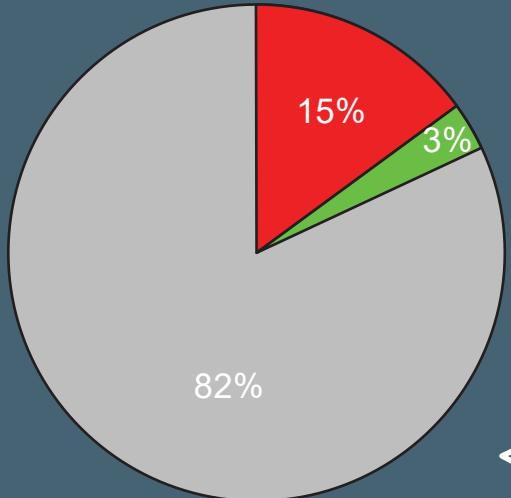
Proteins associated with Aging



Proteins associated with Physical Activity



Hallmarks of Aging



Mitochondrion

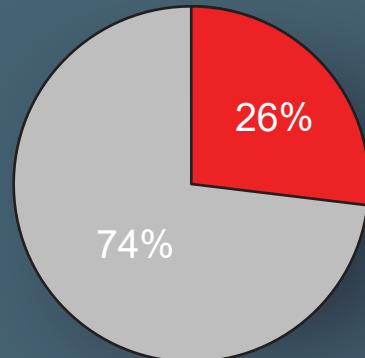
Splicing

Others

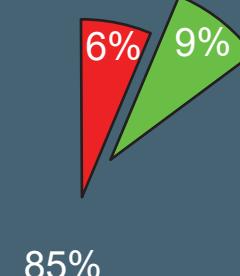
(Immunity/Inflammation,
NMJ, Proteostasis,
Senescence, Transcription
Regulation, Epigenetic
Regulations, Genomic
Maintenance, Ribosomal,
Unknown)

Age Associated Protein Categories

Adjusted for Gender, PA, Race, Fiber ratio, BMI and Batch effects



Abundance Decreased
With Age (361)

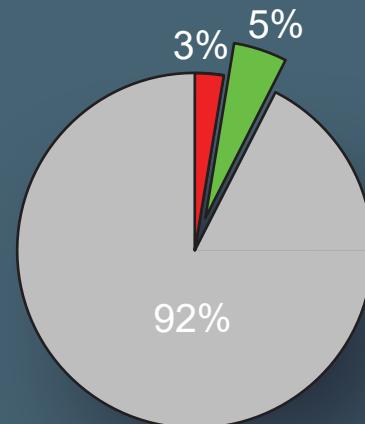


85%

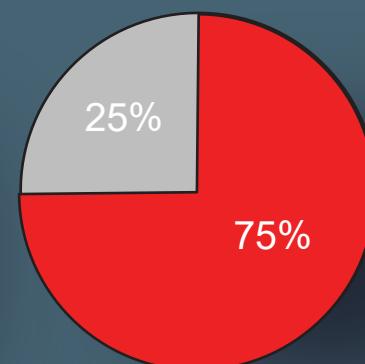
Abundance Increased
With Age (904)

Physical Activity Associated Protein Categories

Adjusted for Gender, Age, Race, Fiber ratio, BMI and Batch effects



Abundance Decreased
With PA(695)



Abundance Increased
With PA(324)

Figure 3 Functional Decline of Mitochondrial Proteins with Age

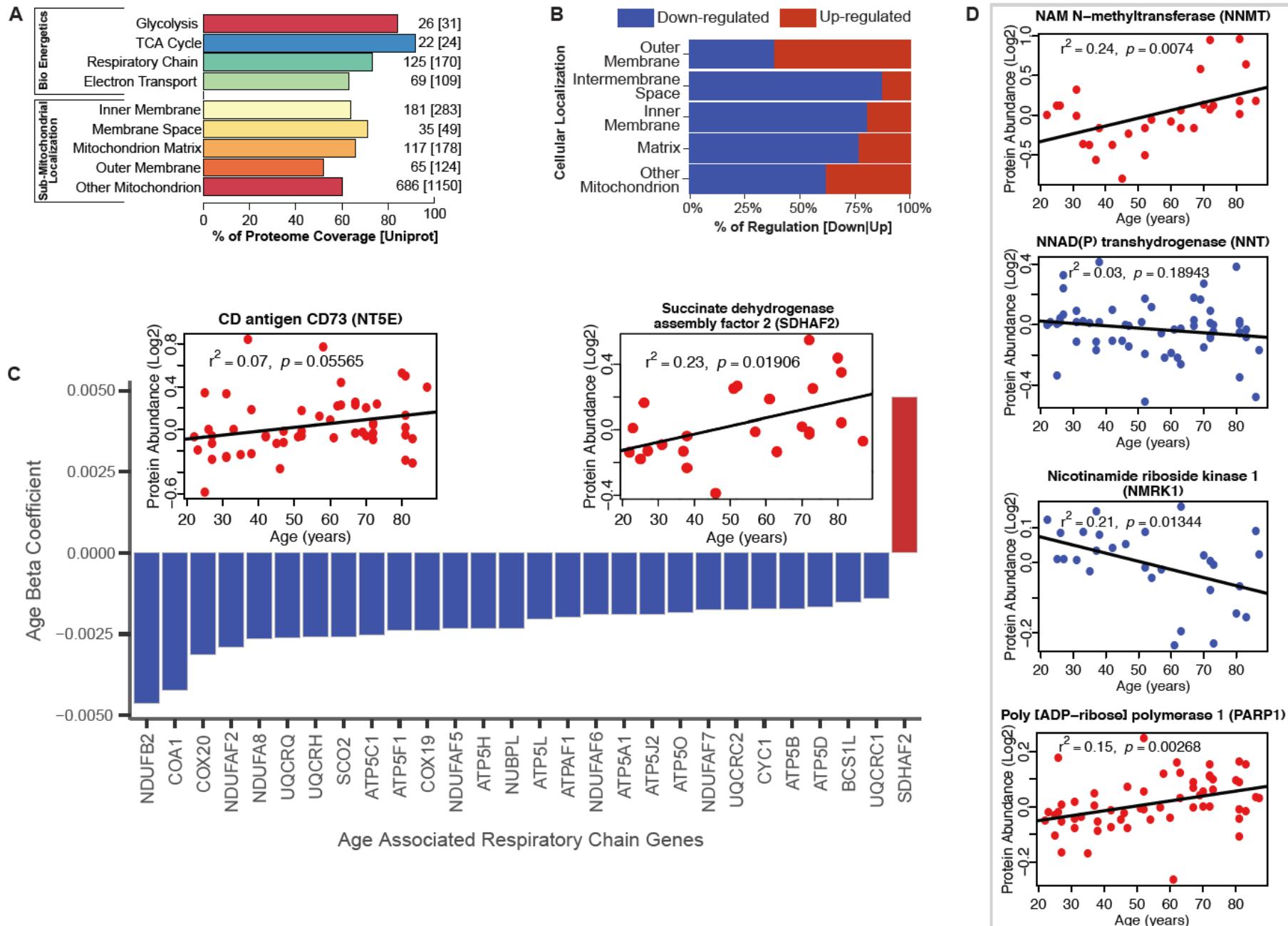
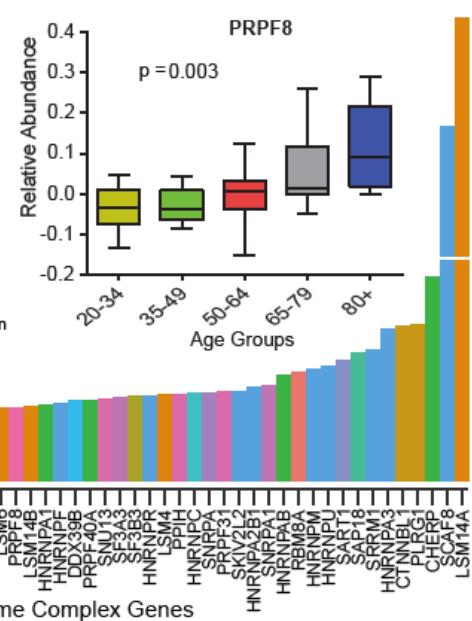
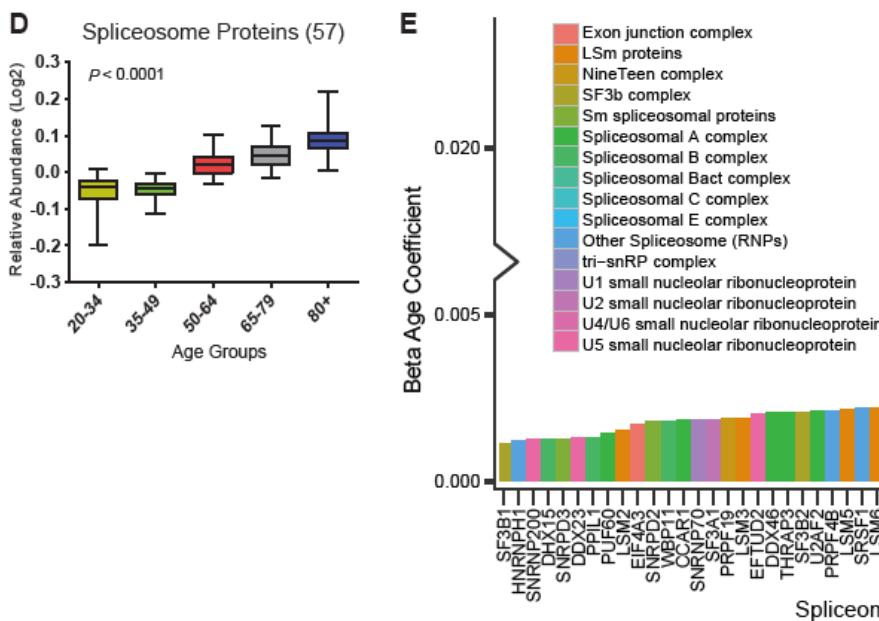
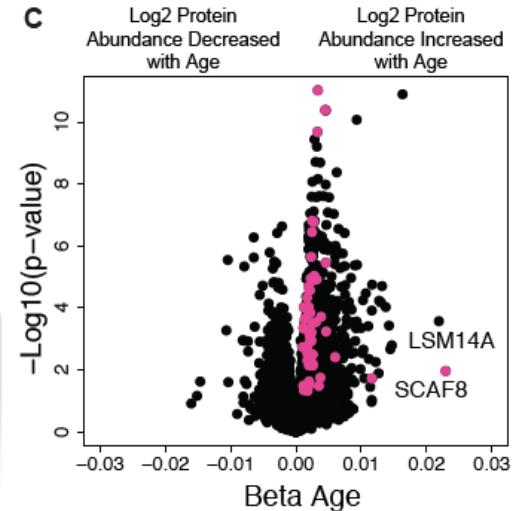
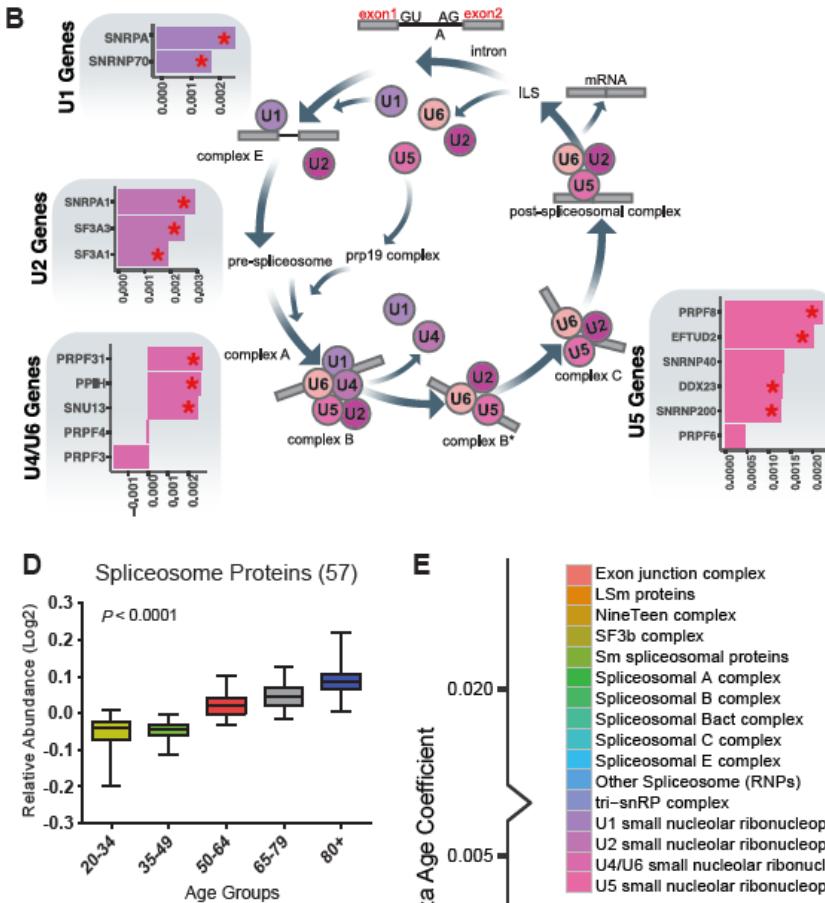
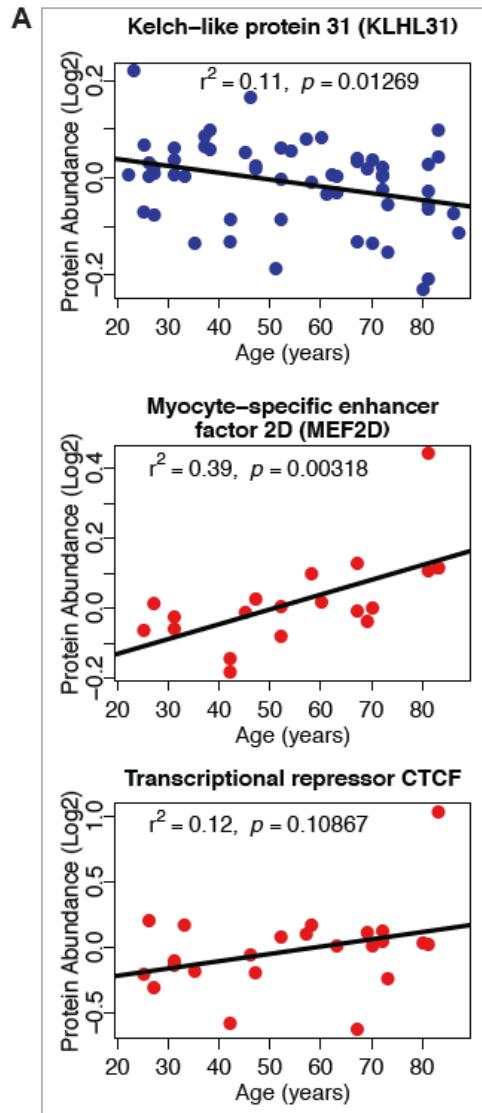
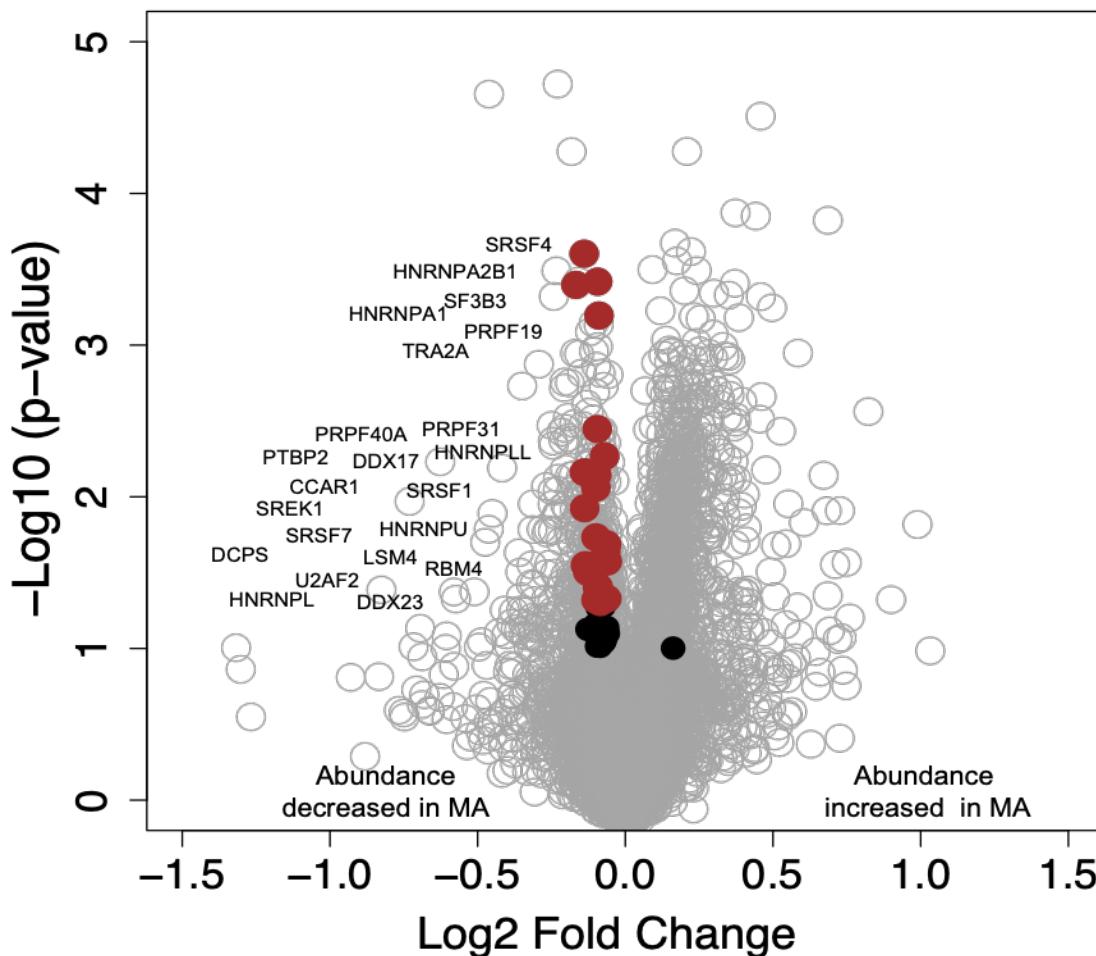


Figure 4 Implications of Proteins that Modulate Transcription and Splicing



Relative Abundance of Spliceosome Proteins in Master Athletes Compared to Age-Matched Controls

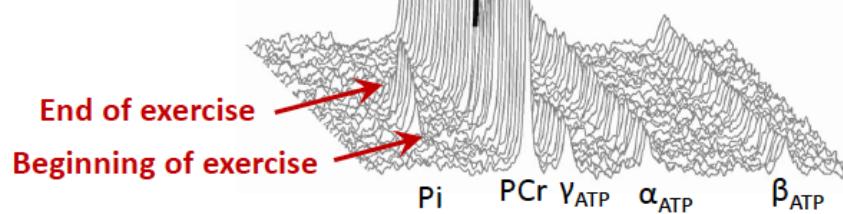
Collaboration with Russel Hepple, PhD (University of Florida)



Discovery proteomics on muscle biopsies quantified using TMT and LC-MS methods. Overall, 132 spliceosome pathway proteins were quantified. Of these, 122 were underrepresented in master athletes compared to controls, and for 22 of them the difference was significant.

Adjusting for Age and Physical Activity, Up-Regulation of the Splicing Machinery is Associated with Better Mitochondrial Function

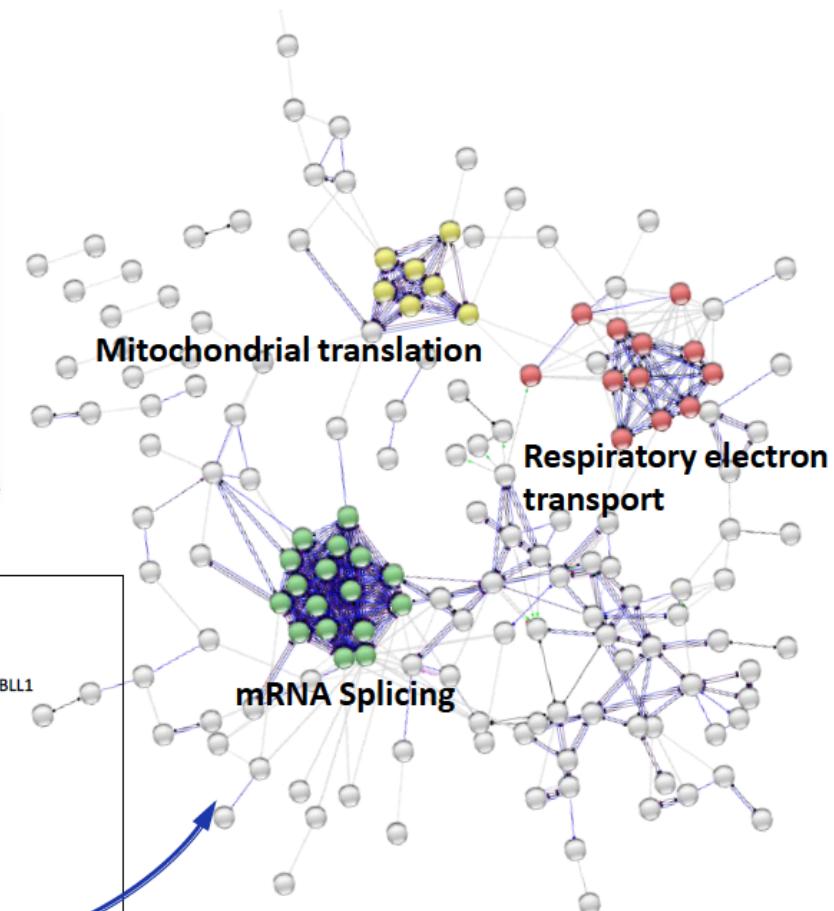
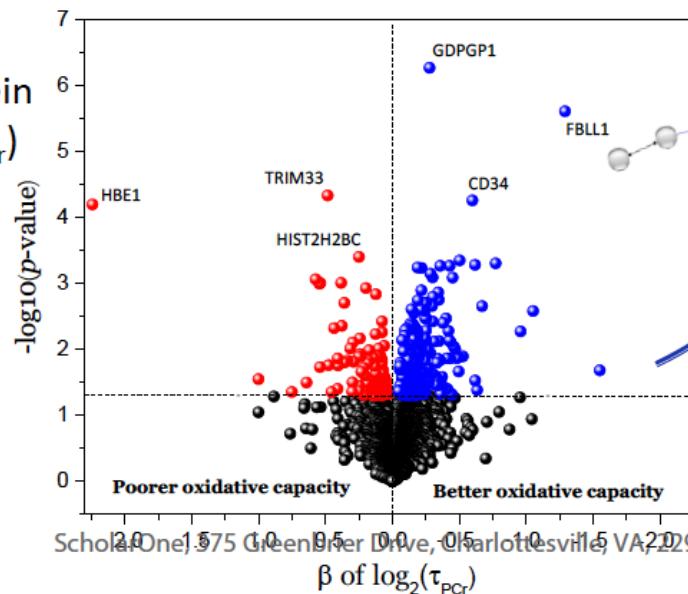
Post-exercise phosphocreatine (PCr) recovery time (τ_{PCr}) measurement by ^{31}P MRS; reflects *in vivo* oxidative capacity of skeletal muscle.



Skeletal muscle protein association with (τ_{PCr})



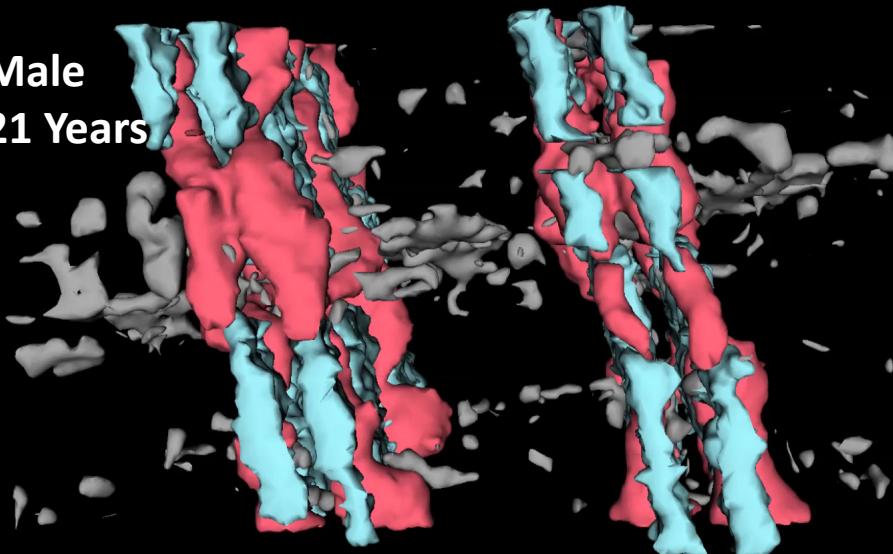
Fatemeh Adelnia



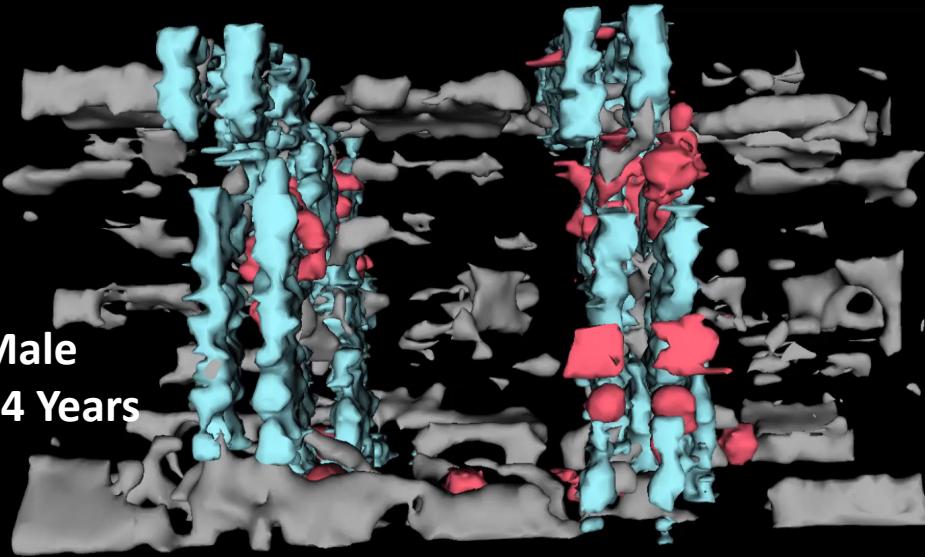
Network representation of the 253 proteins associated with better muscle oxidative capacity

3D Reconstructions of FIB-SEM images in 3 different age groups. Mitochondria are pink, Z-bands are cyan, and voided areas are gray.

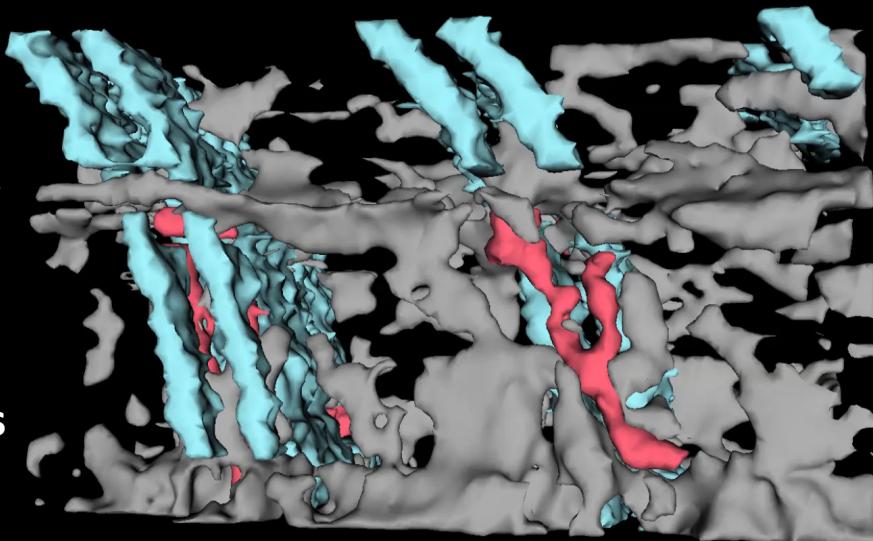
Male
21 Years



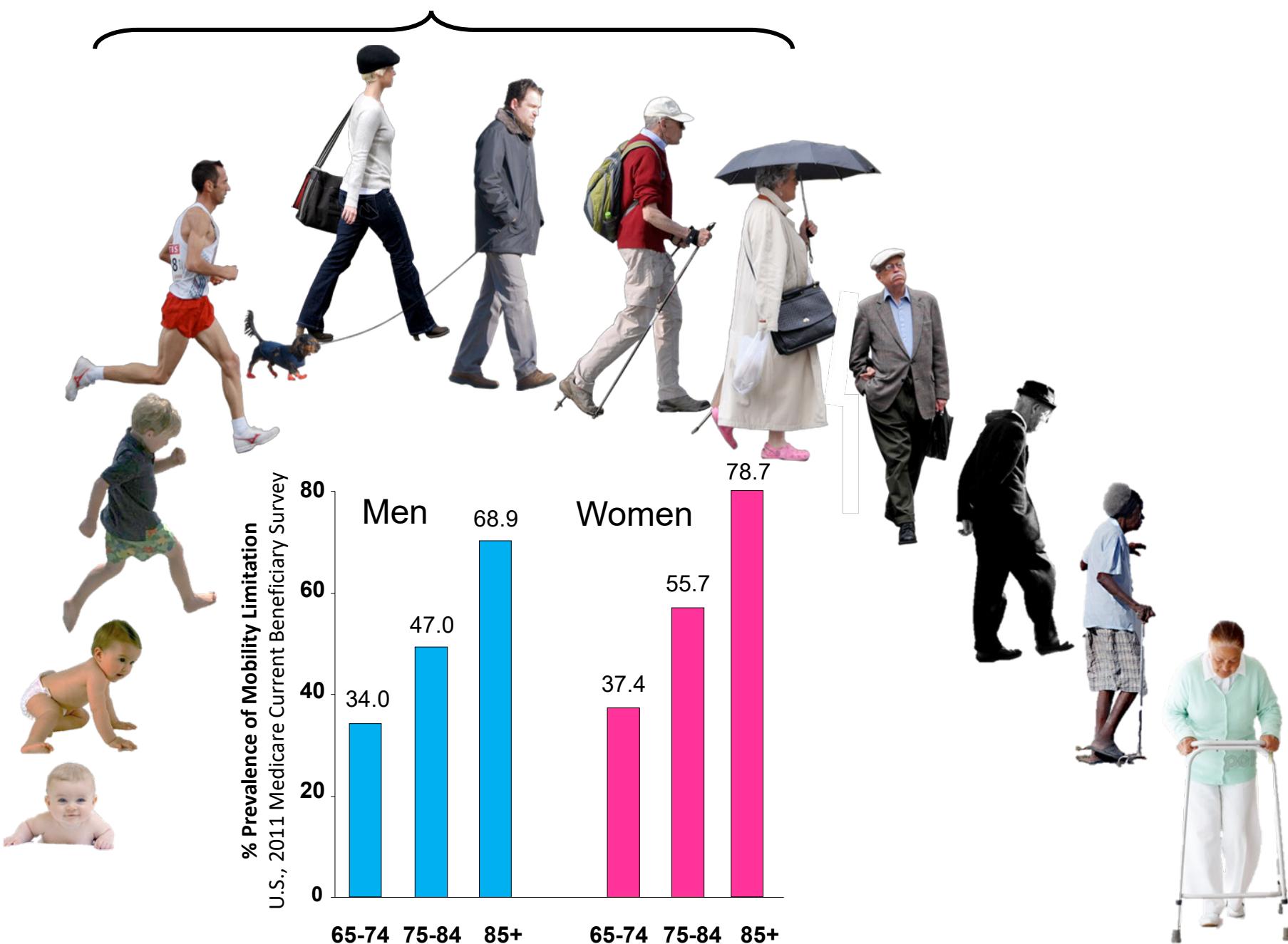
Male
54 Years



Male
82 Years

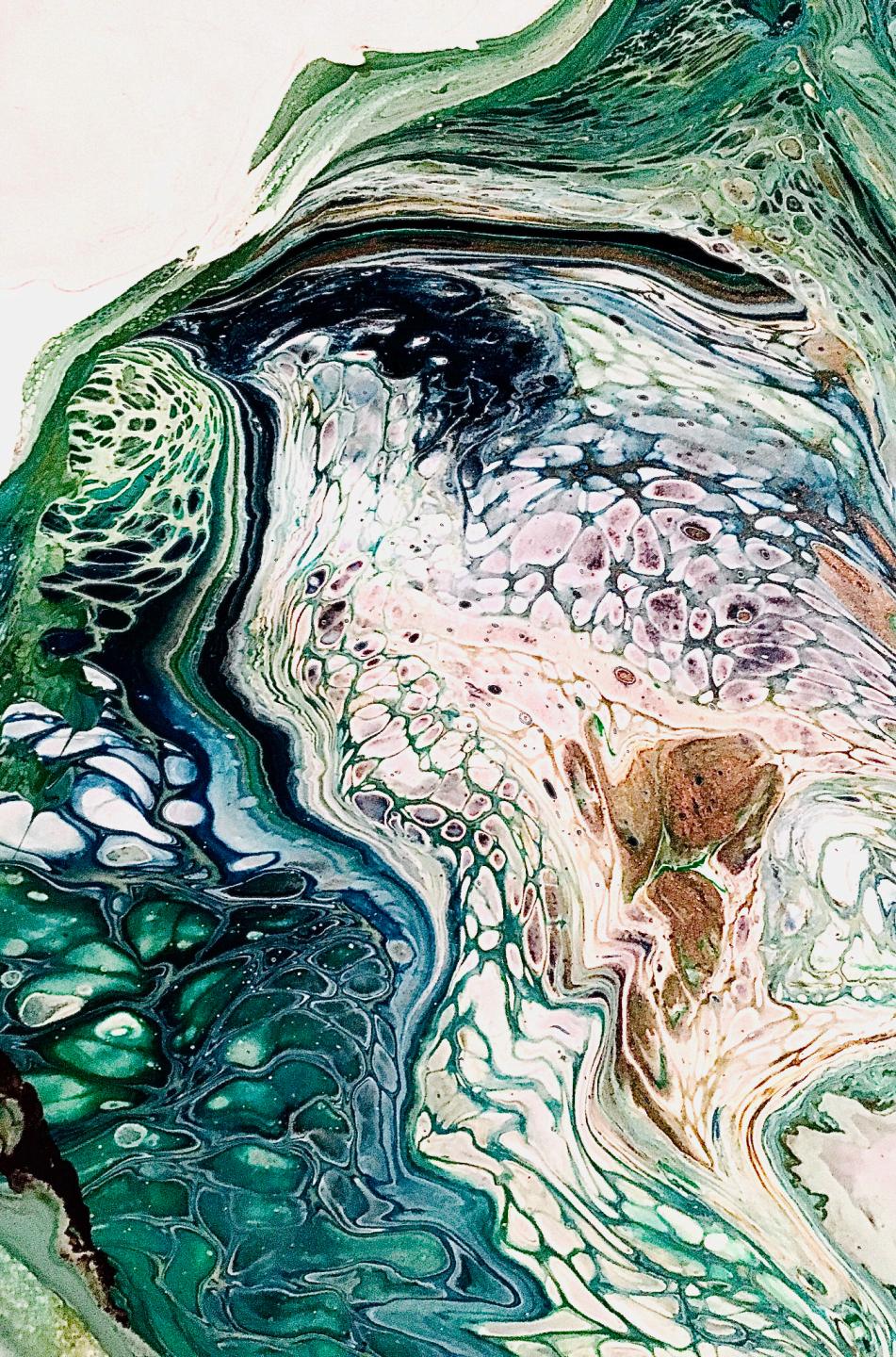


Geroscience anticipates secondary prevention



DNA Methylation Landscapes in Aging & Disease

Morgan Levine
Assistant Professor
Department of Pathology
Yale University School of Medicine



Cancer Risk

What is the biggest risk factor for lung cancer?

Smoking increases lung cancer incidence and death by 15 to 30 fold

1 in 200k chance for ages 25-29, nearly 400 in 100k chance ages 75-79

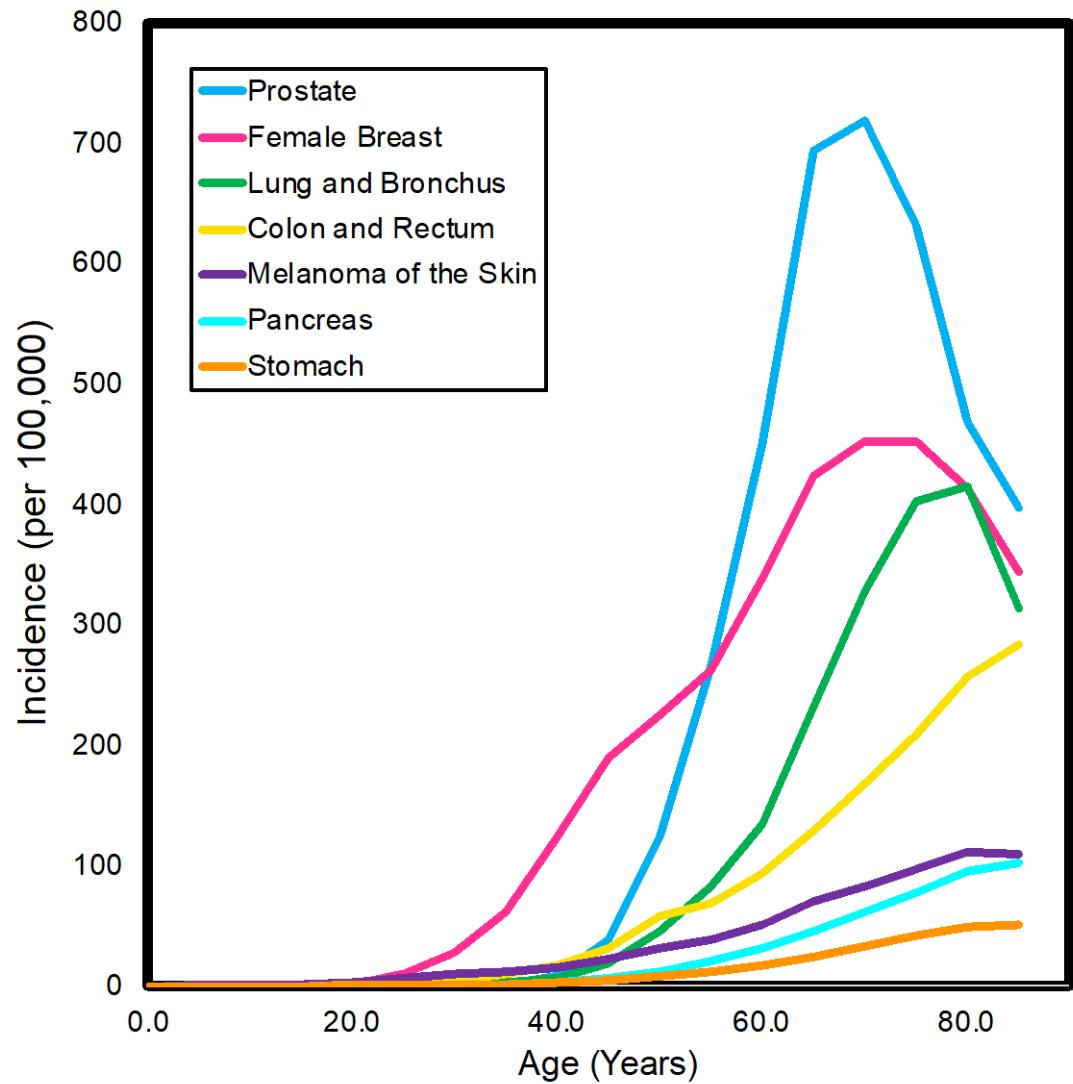


Geroscience

The aging process is thought to play a causal role in the etiology of most major chronic diseases.

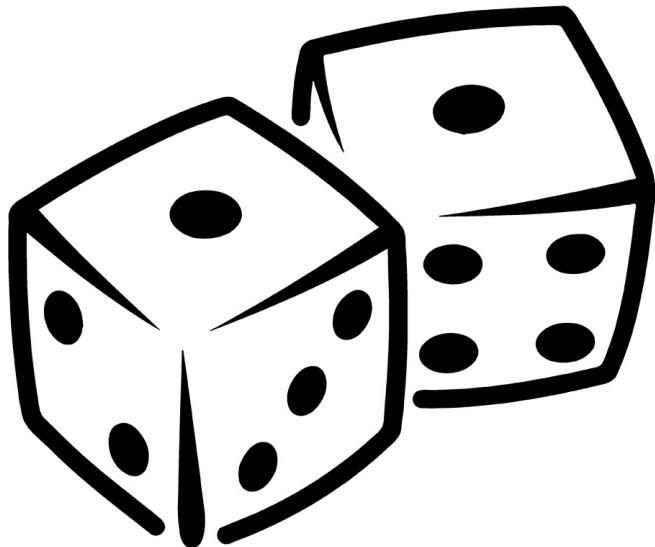


'YOU'RE DELIBERATELY PUTTING YOURSELF
AT RISK OF ILL HEALTH BY BEING OVER 65...'



Cancer Risk & Age

Is aging causal or consequential in cancer?

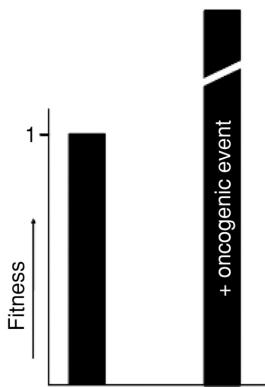


The more times you roll the dice
(function of chronological time)
the more likely your chances?

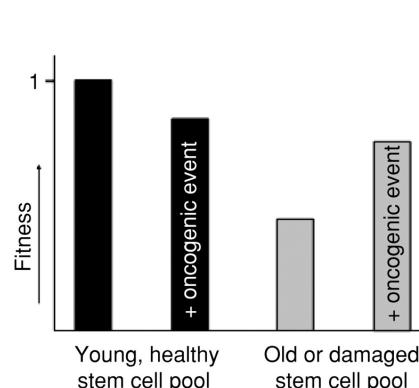
Cancer Risk & Age

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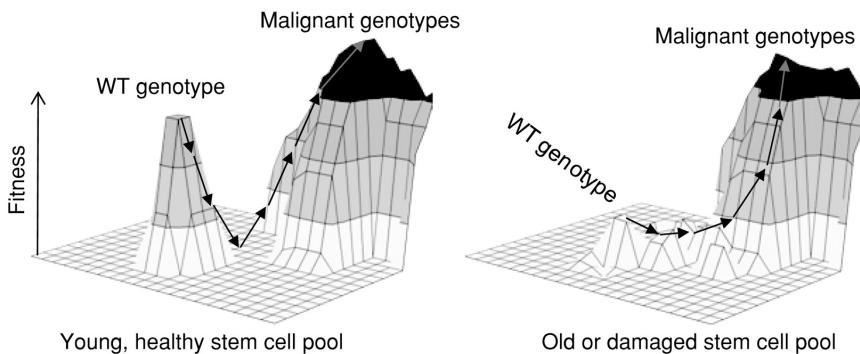
A Conventional model



B Adaptive Oncogenesis



C Adaptive landscapes



Adaptive Oncogenesis Model

Dr. James DeGregori

Roughly half of all mutations occur before full body maturation

Context matters!

system-level dynamics that change with age alter the fitness landscape

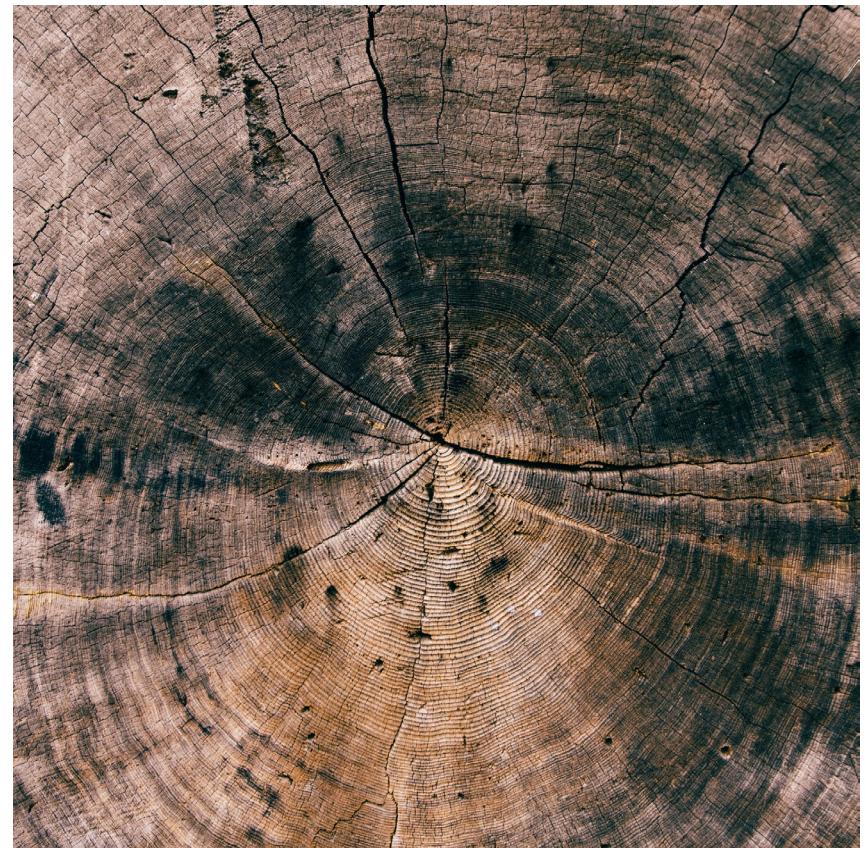
Biomarkers of Aging

Useful proxies that estimate aging
(or agedness) of a sample.

Should Answer:

Biologically, what differentiates
the average 20 year old from the
average 80 year old?

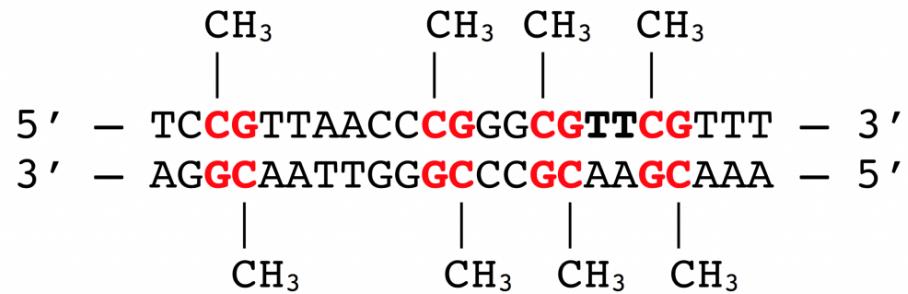
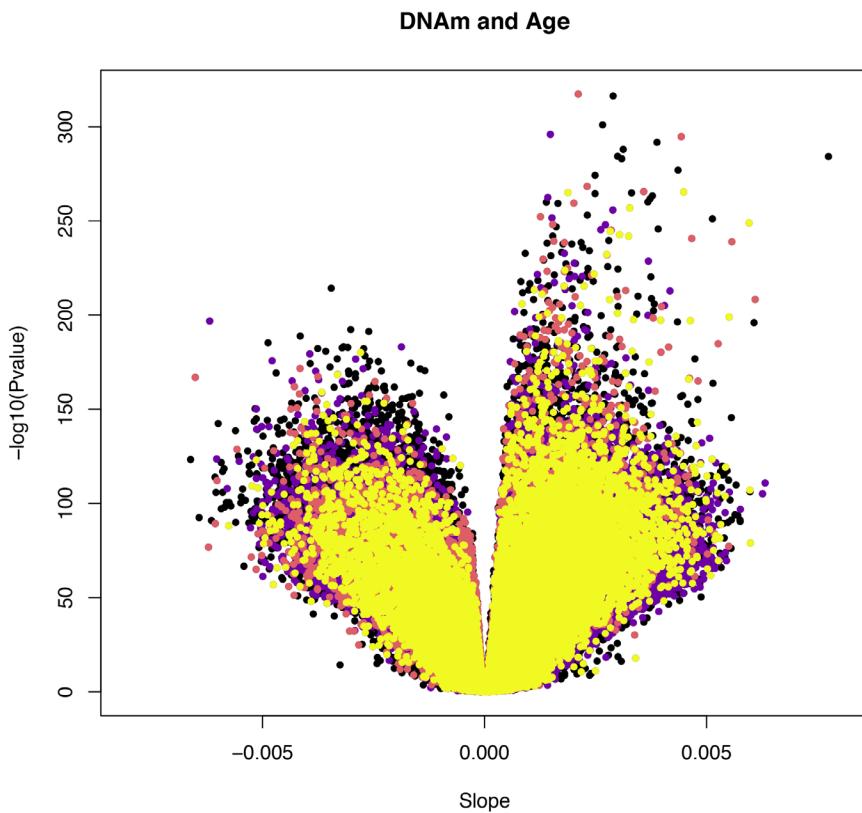
What differentiates a healthy 80
years old from an unhealthy 80
year old?



Epigenetics: Molecular OS

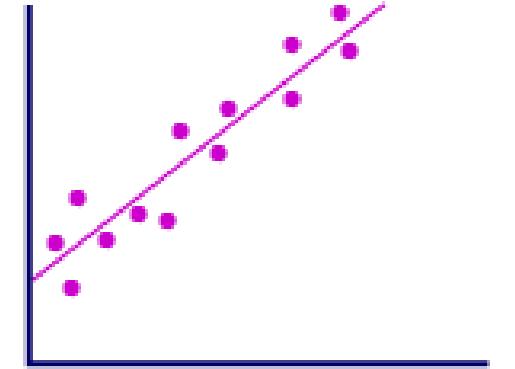
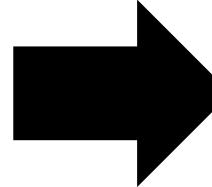
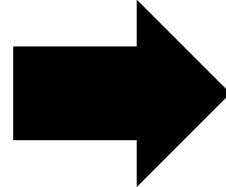
DNA Methylation (DNAm)

Involved in cell proliferation/differentiation, transcriptional repression, genomic imprinting, organization of chromatin.



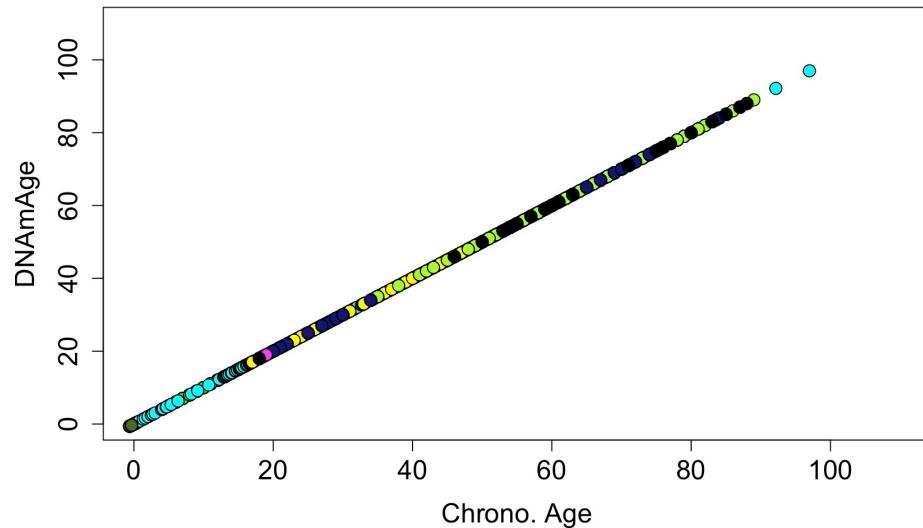
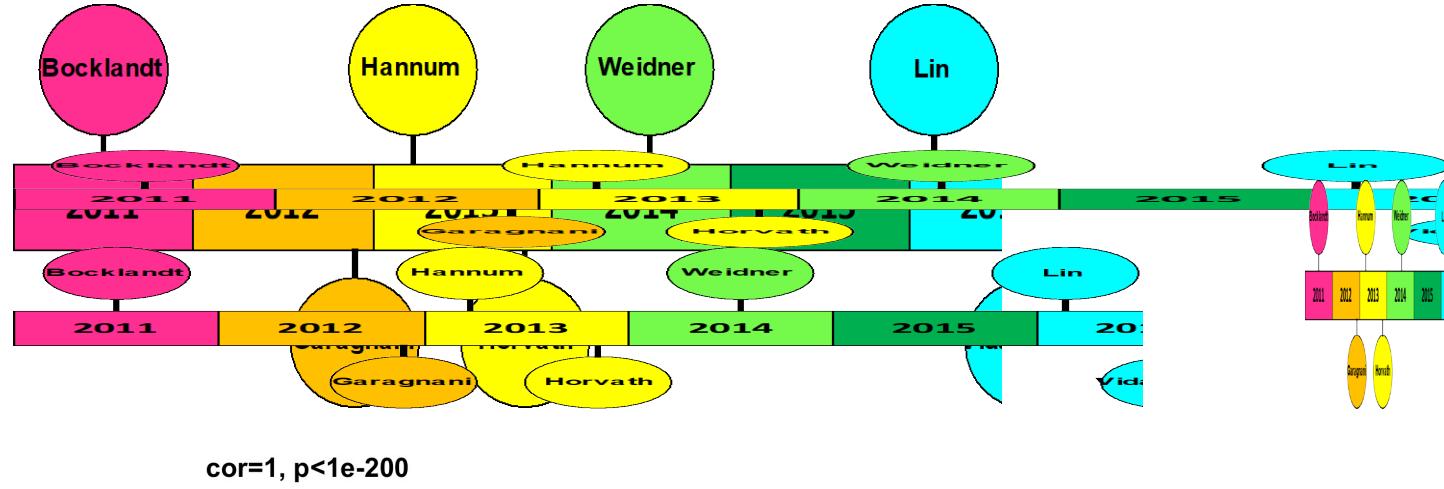
Epigenetic Clocks

Because of the precise age changes,
we can use machine learning to predict “the age”
of a sample based on its DNAm levels.



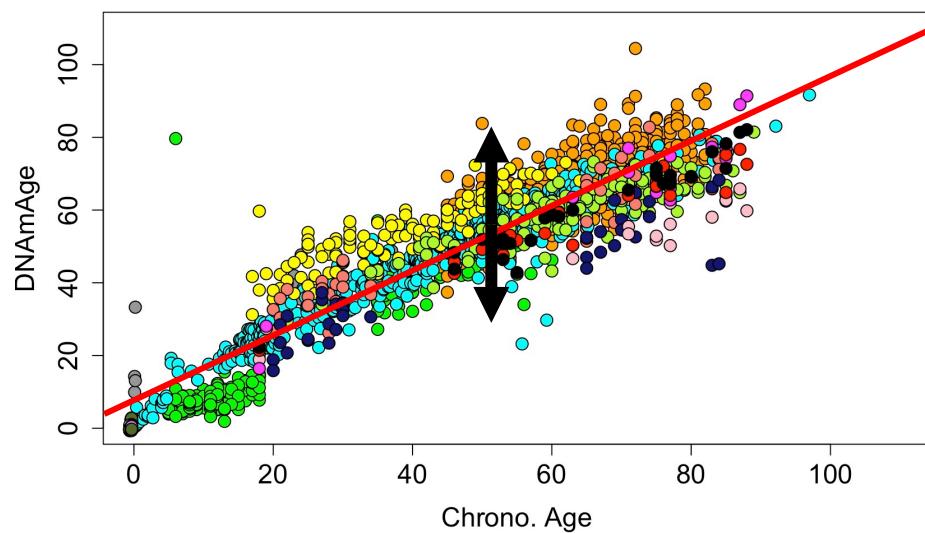
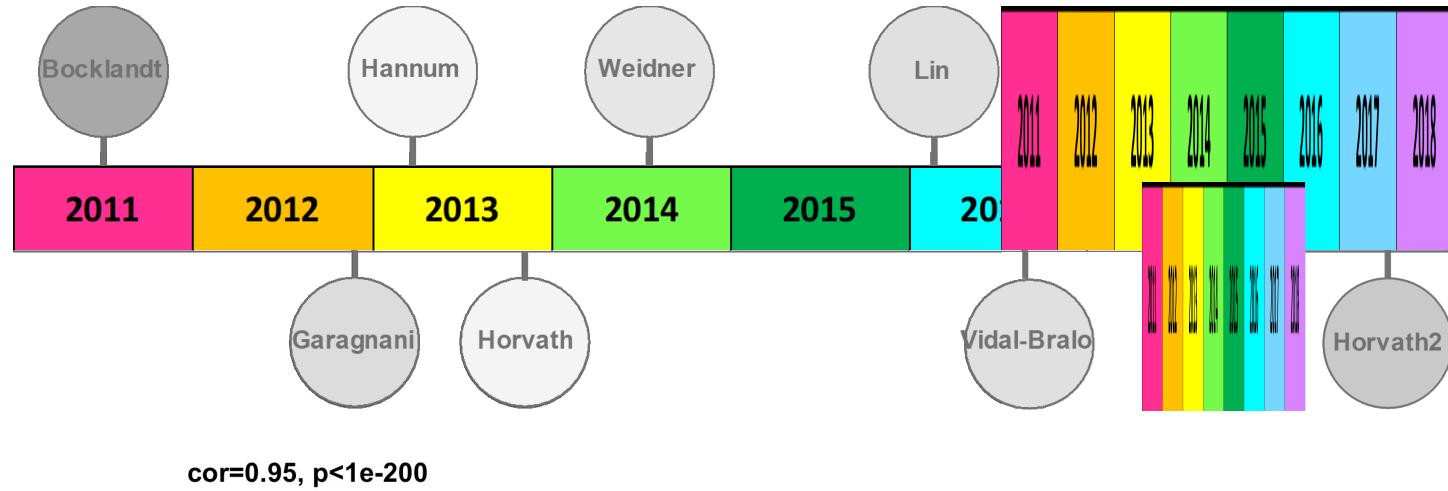
Independent variable

Epigenetic Clocks



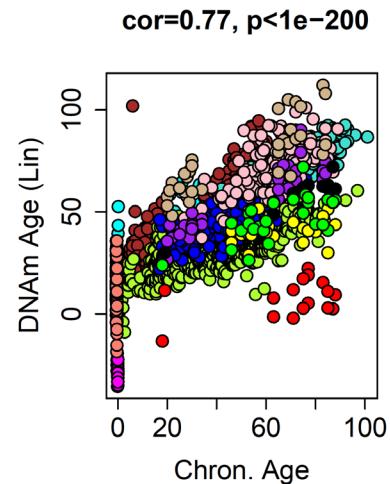
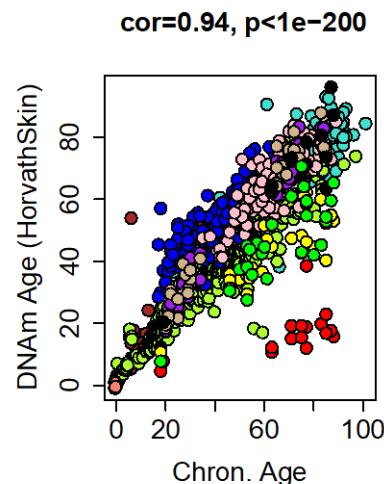
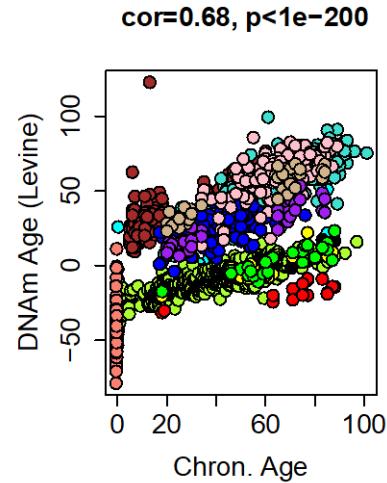
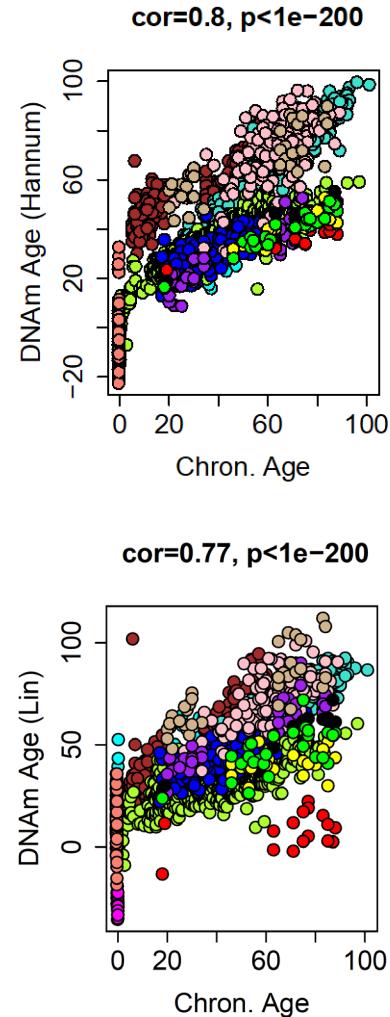
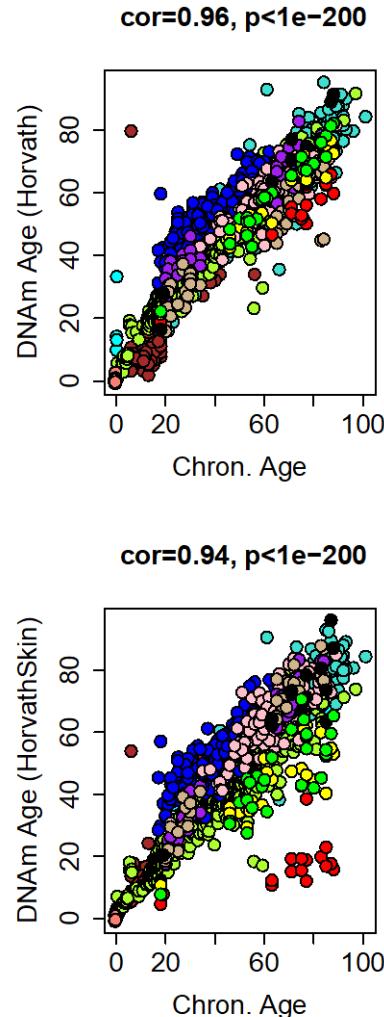
1st Generation:
Age Predictors
Supervised machine learning
to predict age from
hundreds of CpGs

Epigenetic Clocks



2nd Generation:
Aging Outcome Predictors
Supervised machine learning
to predict age-related outcomes
from hundreds of CpGs

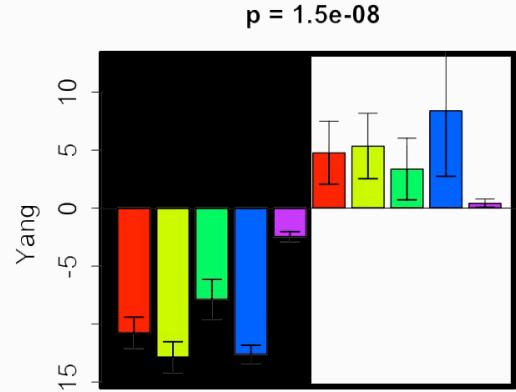
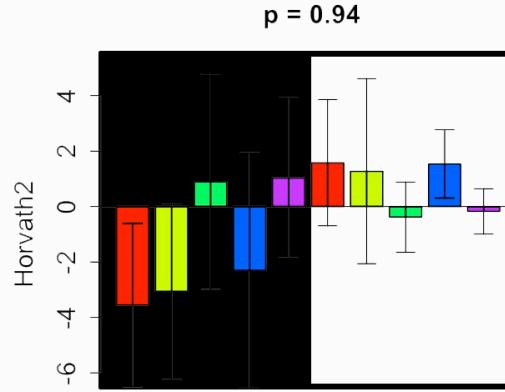
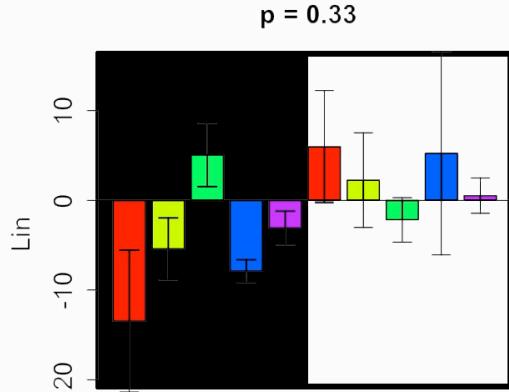
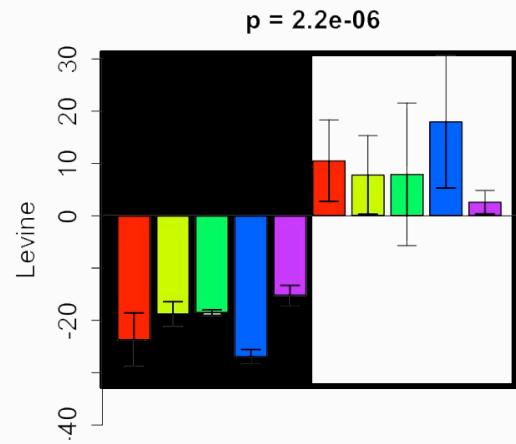
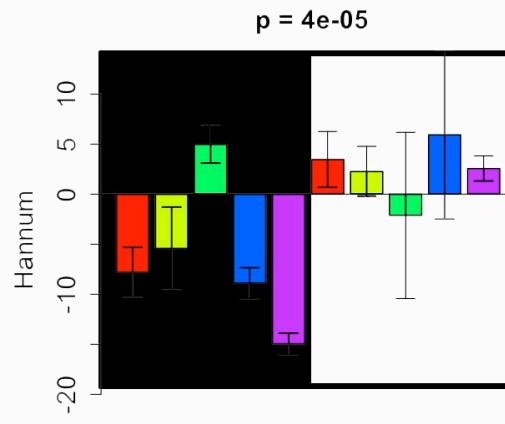
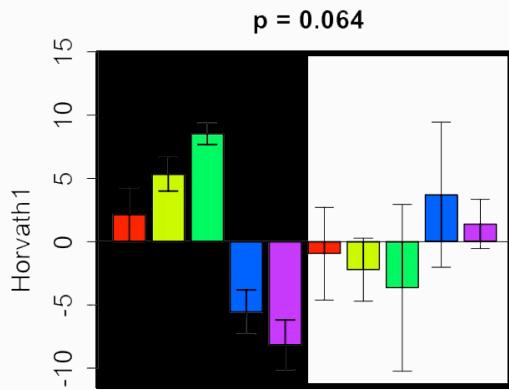
Epigenetic Clocks



Epigenetic Clocks

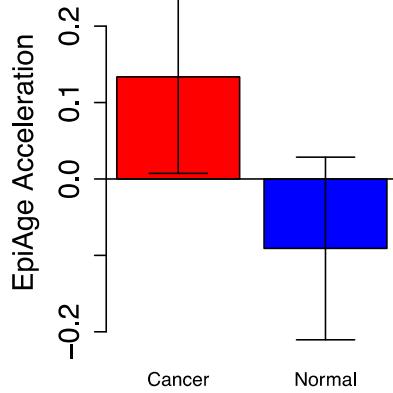
Normal Tumor

breast, colon, lung, pancreas, thyroid

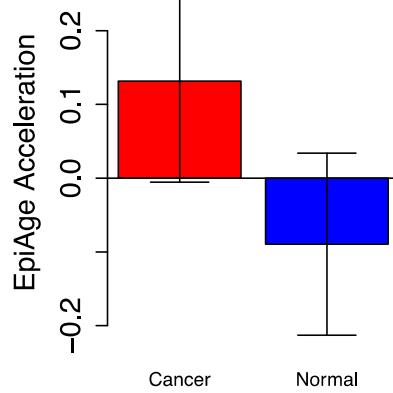


Epigenetic Clocks

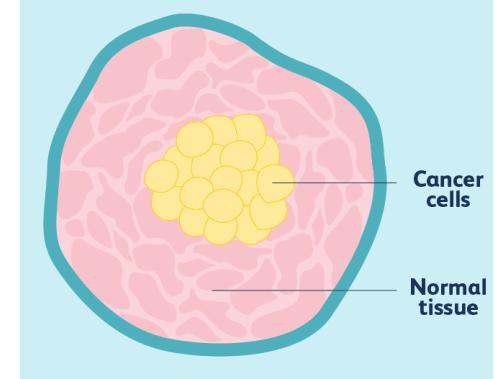
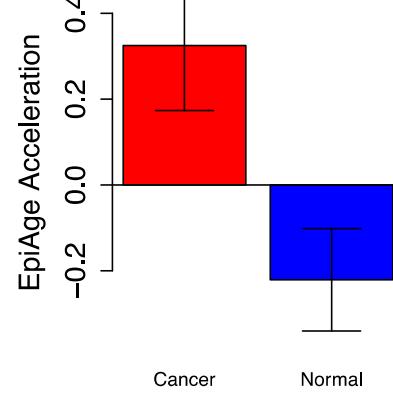
Horvath1 $p = 0.21$



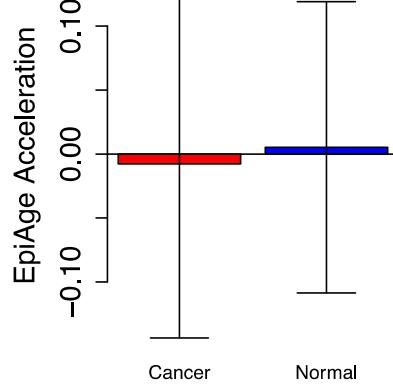
Hannum $p = 0.23$



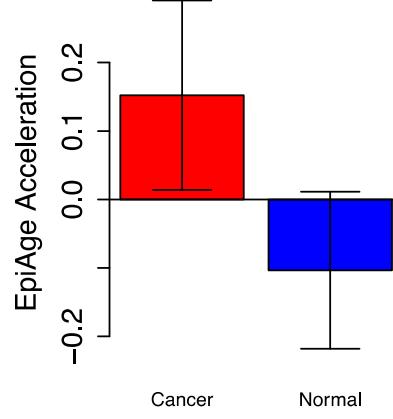
Levine $p = 0.0023$



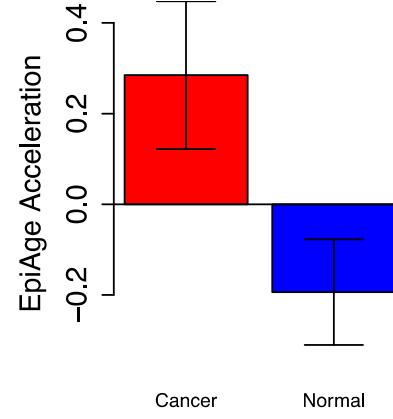
Lin $p = 0.88$



Horvath2 $p = 0.19$



Yang $p = 0.01$



Epigenetic Clocks



alarm clock

taking apart

© photo by Gabriel Menashe | 06/2013
takingapart.com

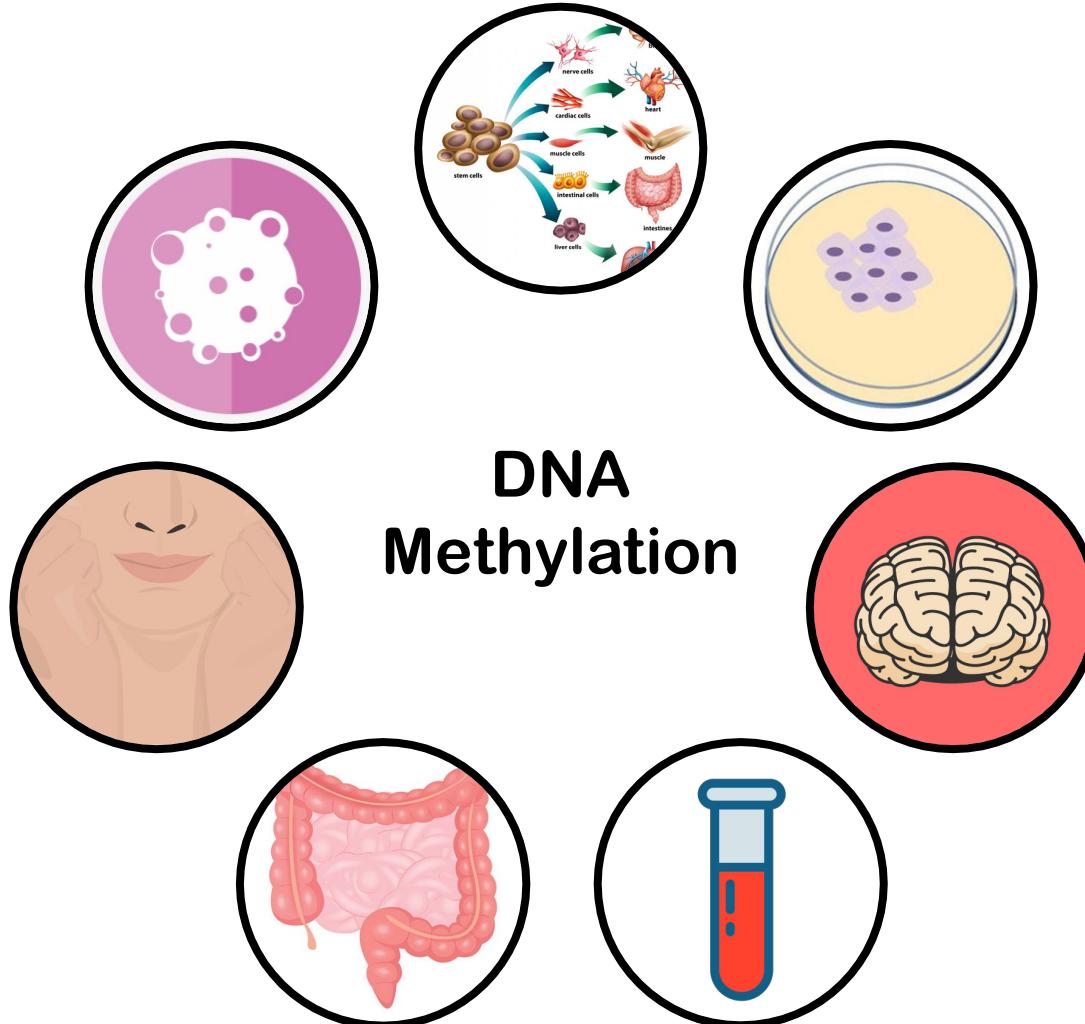


DNAm can capture a lot of cellular/molecular changes.

What are the core signals (parts) being captured by the clocks?

Are there shared signals across aging phenomena and/or tissues?

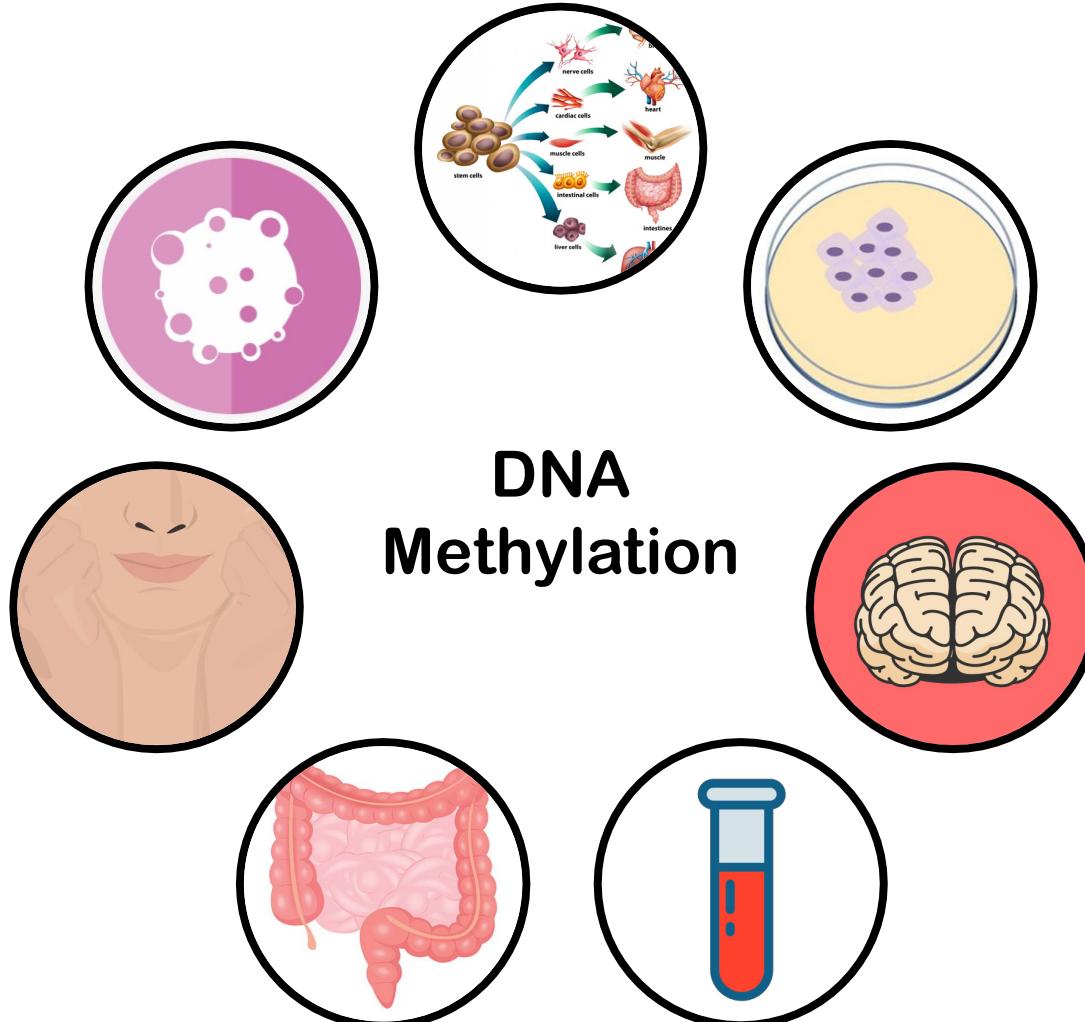
Shared DNAm Signals



PCA in 9 Datasets

Whole blood
Adult Brain
Developmental Brain
Dermis
Epidermis
Senescence
iPSC/Reprogramming
Tumor/Normal

Shared DNAm Signals

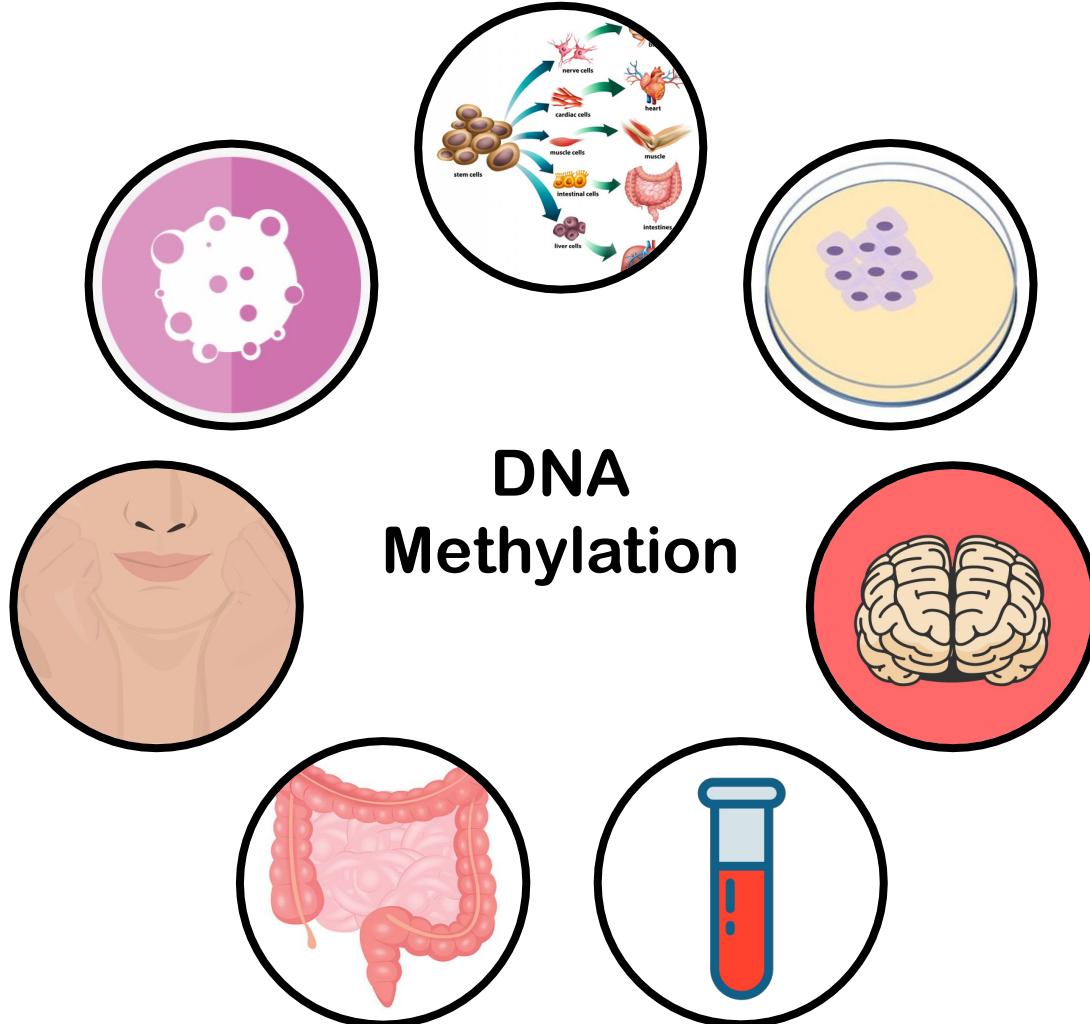


PCA in 9 Datasets

Whole blood
Adult Brain
Developmental Brain
Dermis
Epidermis
Senescence
iPSC/Reprogramming
Tumor/Normal

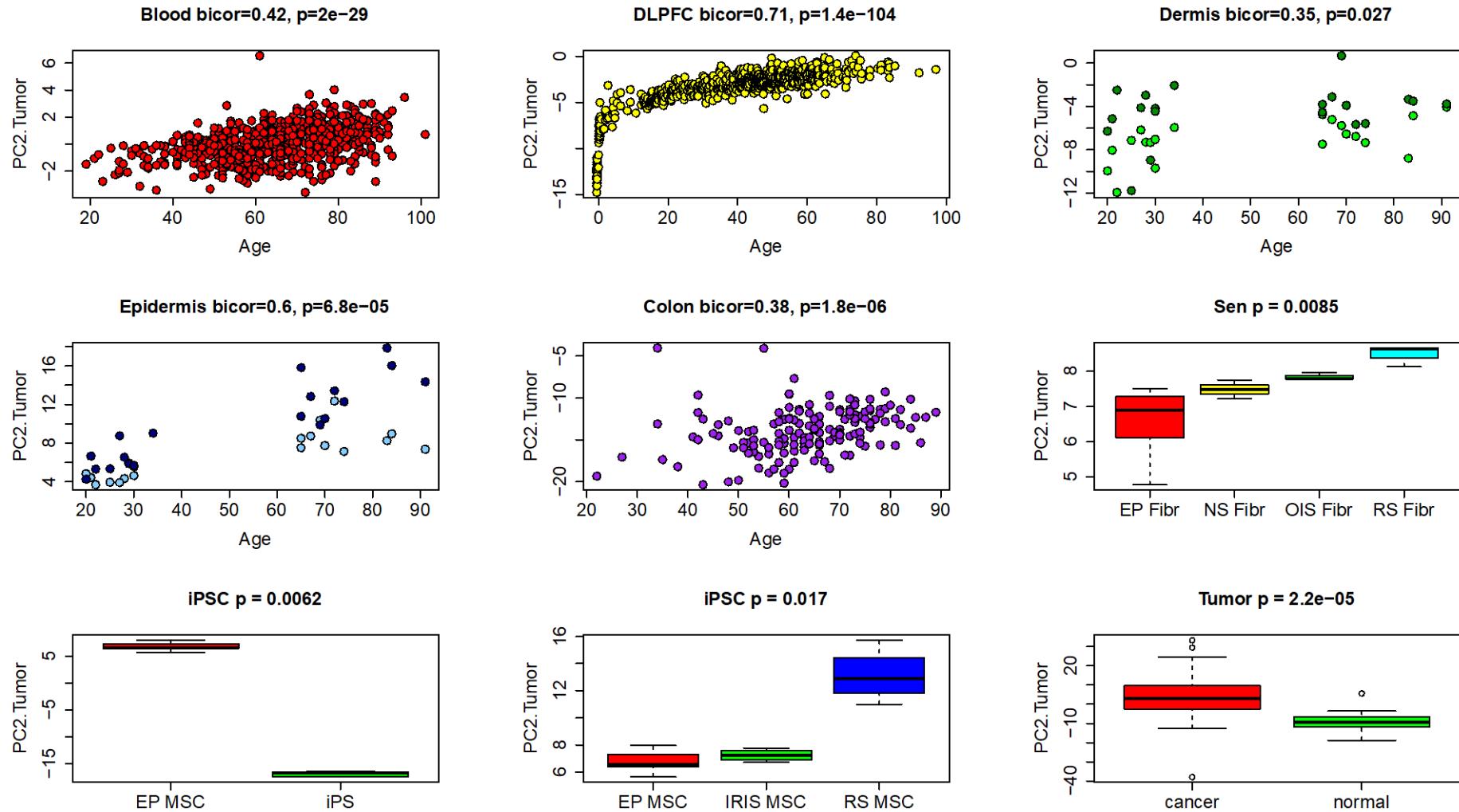
Estimate top 10 PCs in all datasets (n=90 variables in each dataset)

Shared DNAm Signals

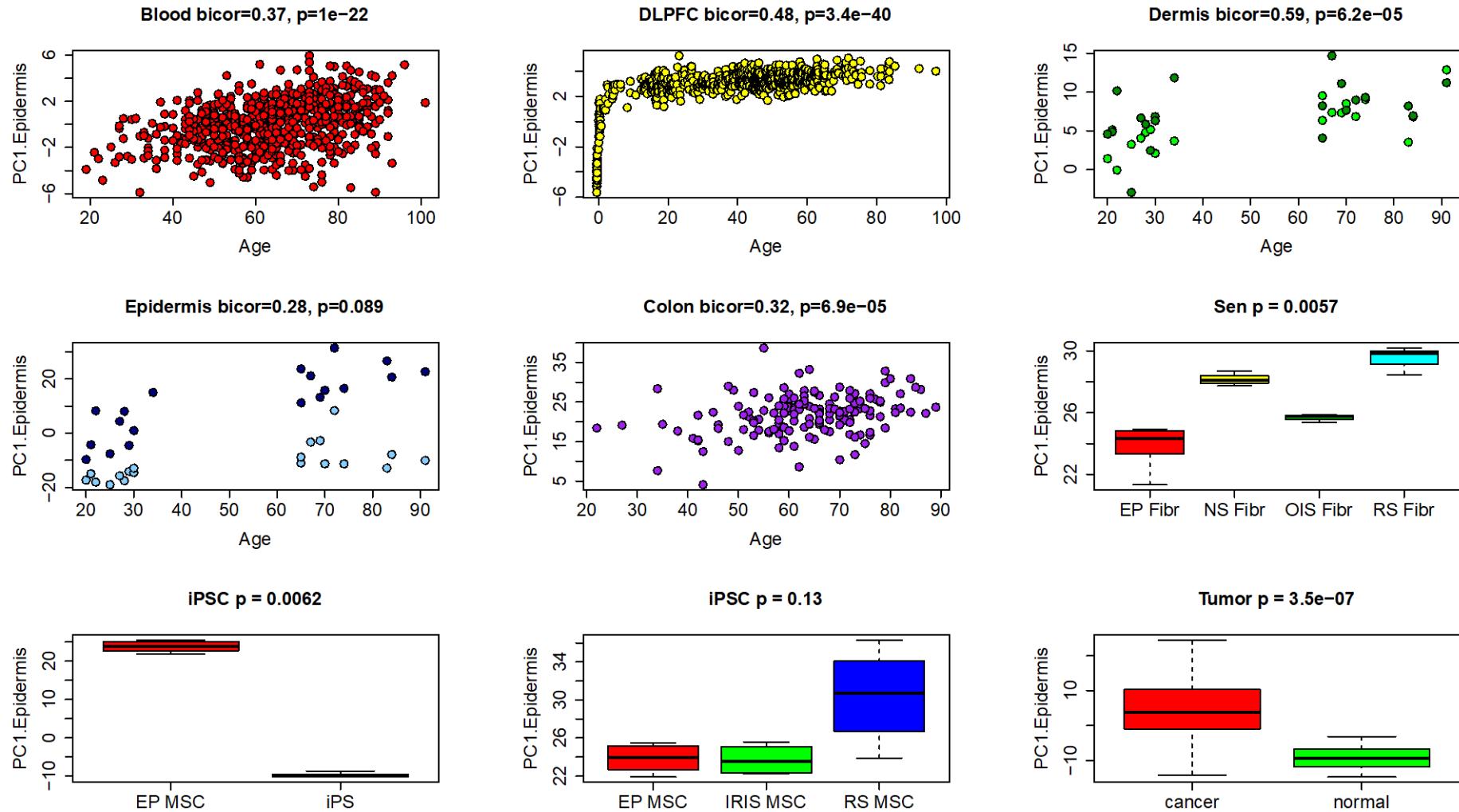


10 of the 90 PCs
had consistent
age and/or aging-
outcome
associations
across all 9
datasets

Shared DNAm Signals



Shared DNAm Signals



Conclusions

Does biological aging in a tissue predispose it to tumorigenesis?

- Aging/Cancer: probability with time vs. causal driver

Preliminary Evidence for Aging → Cancer

1. One can estimate “aging” in various tissues using DNAm.
2. For most clocks, tissues show different rates of aging.
3. DNAMAge can differentiate tumor versus normal tissue (acceleration in cancer).
4. DNAMAge can differentiate normal breast tissue in women with history of breast cancer versus controls.
5. DNAm patterns in cancer apply to other tissues.
 1. Correlate with age in blood, brain, skin, colon
 2. Accelerated in skin exposed to sun
 3. Accelerated in senescent cells (oncogene induced and replicative)

Acknowledgements

ALIS

Laboratory for
AGING IN LIVING SYSTEMS

Current Members

Kyra Thrush (PhD Student Computational Biology)
Albert Higgens-Chen (Post Doc)
John Gonzalez (PhD Student in Molecular Medicine)
Margarita Meer (Assistant Research Scientist)
Diana Leung (PhD Student Computational Biology)
Chris Minteer (PhD Student in Molecular Medicine)

Former Members

Zuyun Liu (Former Post Doc)



National Institut
on Aging



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Human Immunology Project Consortium

Q&A

Save the date for the next webinar:

- September 14, 1-2 p.m. ET
 - Speakers: Dr. Hyman Muss and Dr. Grant Williams
- Send speaker suggestions and other feedback to:
NCIDCCPSagingwebinar@mail.nih.gov



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