



UMass Chan
MEDICAL SCHOOL



Genomics &
Computational
Biology

Linking single-cell transcriptomic and genomic changes in the aging human brain

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Instructor

Umass Chan Medical School

ASHG

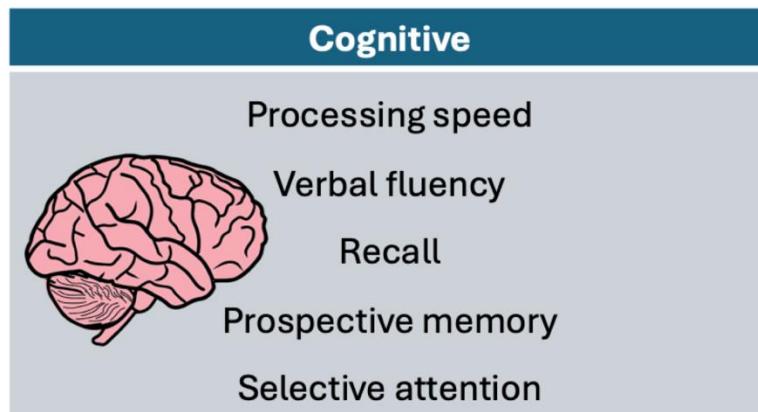
Oct, 2025

Financial Disclosure

I do not have any relationships to report within the last 24 months with ACCME defined ineligible companies.

Why study aging?

Physical	Molecular
Thinning skin and loss of elasticity	Telomere shortening
Graying and thinning hair	Changes in proteostasis
Loss of muscle mass	Epigenetic alterations
Reduction in height	Disrupted macroautophagy
Decreased joint mobility	Genomic instability



- Aging is the primary risk factor for neurodegenerative diseases, cancers, and metabolic disorders.
- Study aging help extending lifespan and developing therapies for prevalent age-related disorders.

Aging and DNA damage

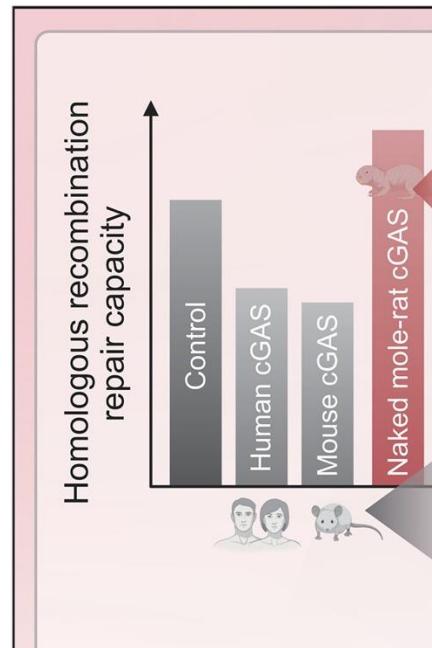
- Genetic disorders with defective DNA repair show premature aging, supporting the link between DNA damage and aging.



Cockayne Syndrome caused by ERCC6 or ERCC8 mutation

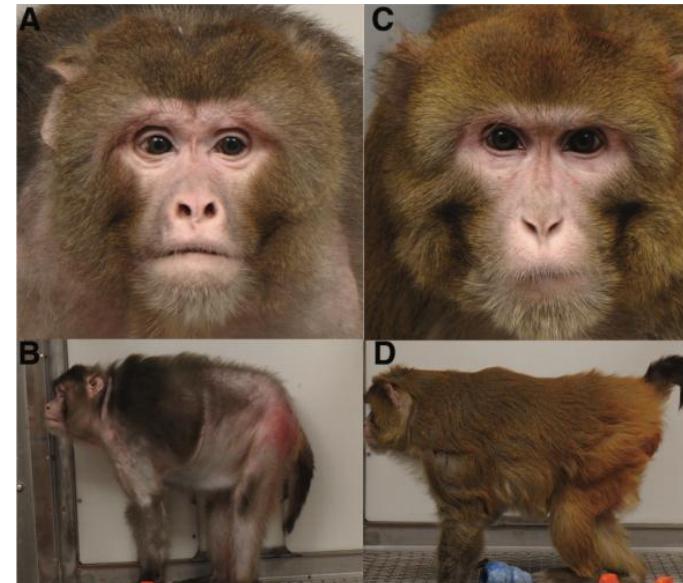
Aging and DNA damage

- Genetic disorders with defective DNA repair show premature aging, supporting the link between DNA damage and aging.
- Species with enhanced DNA repair mechanisms, like naked mole rats, exhibit slower aging and lower cancer rates.



Aging and DNA damage

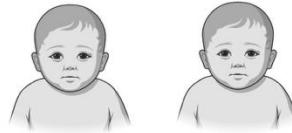
- Genetic disorders with defective DNA repair show premature aging, supporting the link between DNA damage and aging.
- Species with enhanced DNA repair mechanisms, like naked mole rats, exhibit slower aging and lower cancer rates.
- Caloric restriction can reduce oxidative stress and DNA damage, prolonging lifespan in various organisms.



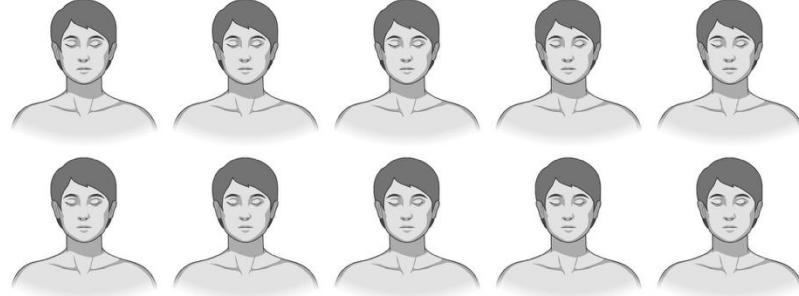
Genome and transcriptome in single-cell

19 fresh frozen human
prefrontal cortex

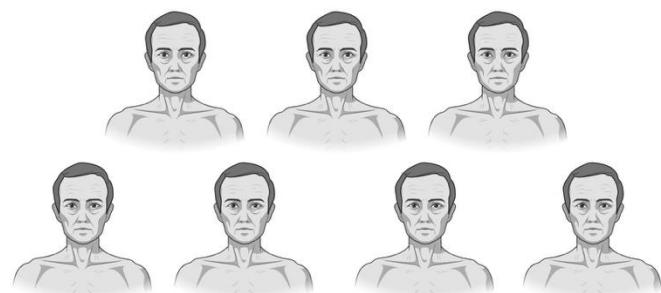
two
infants



ten
adults



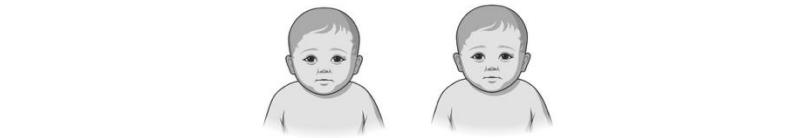
seven
elderly



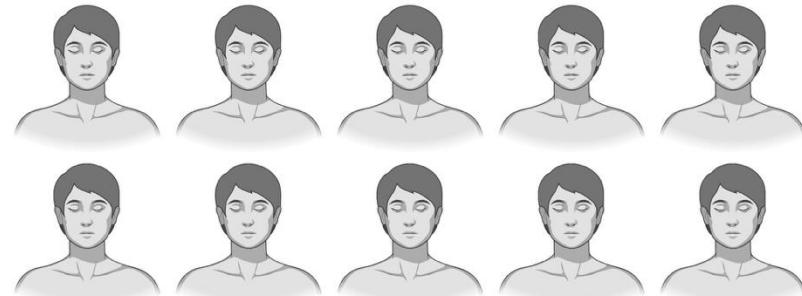
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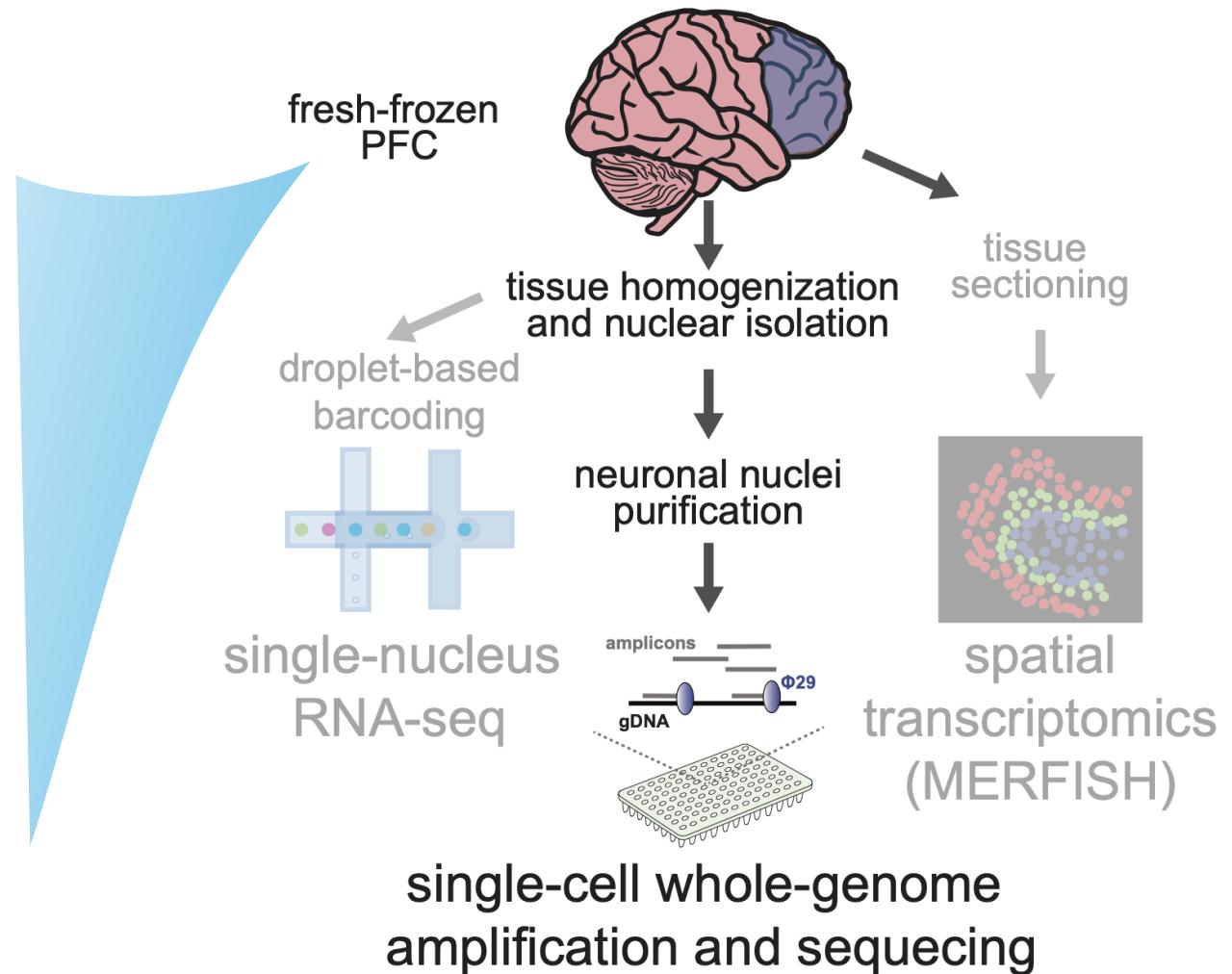
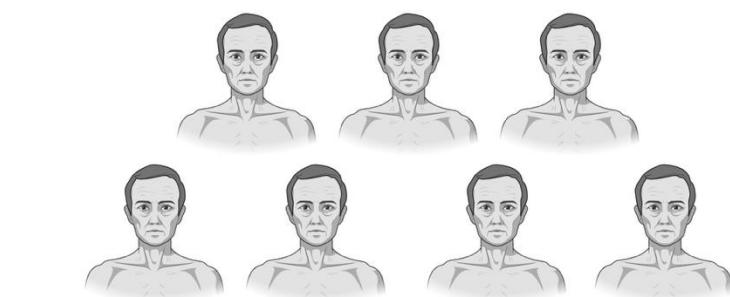
two infants



ten adults



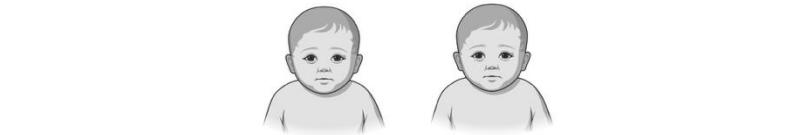
seven elderly



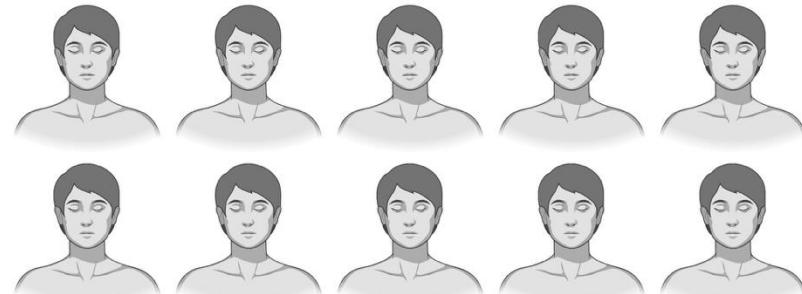
Genome and transcriptome in single-cell

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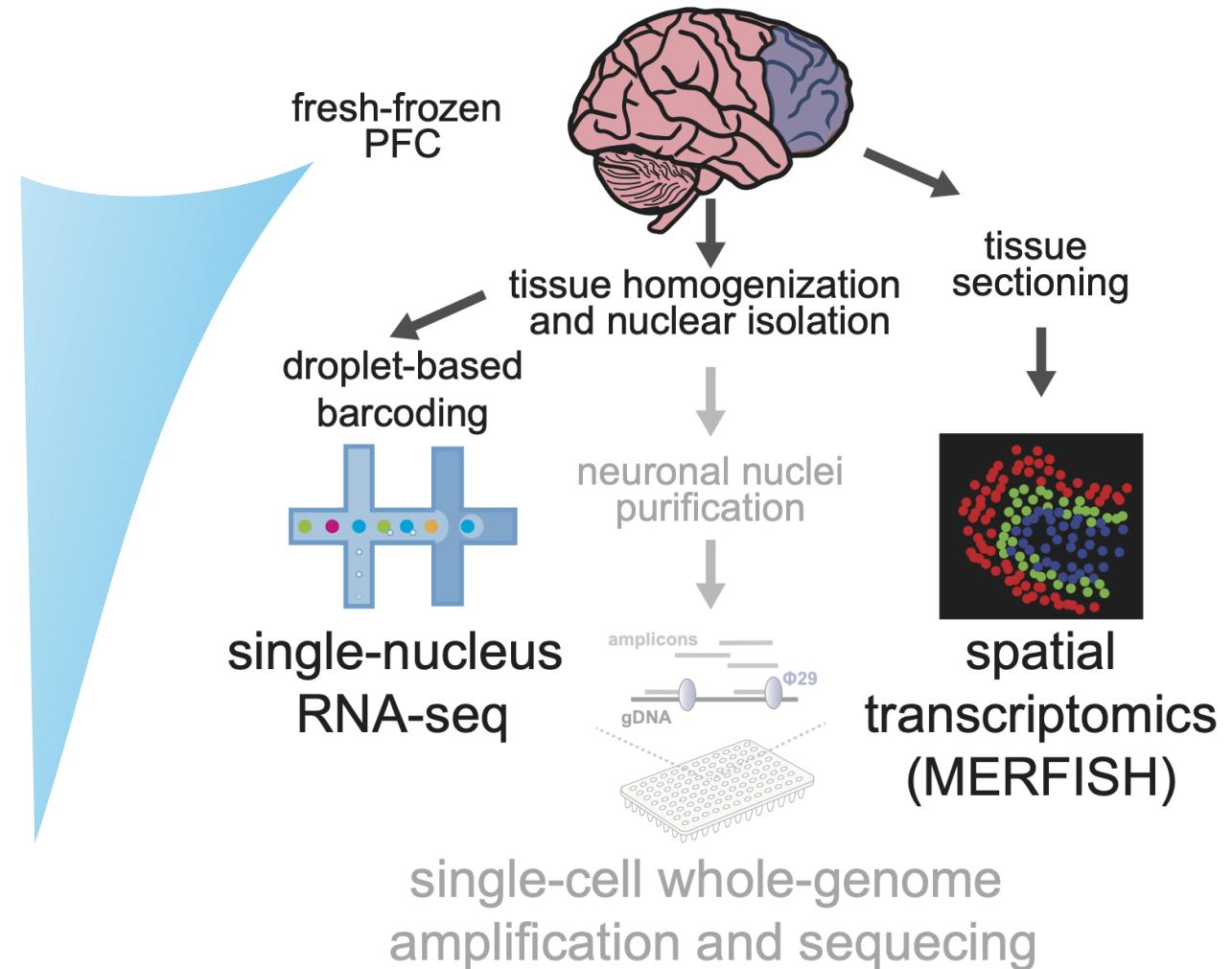
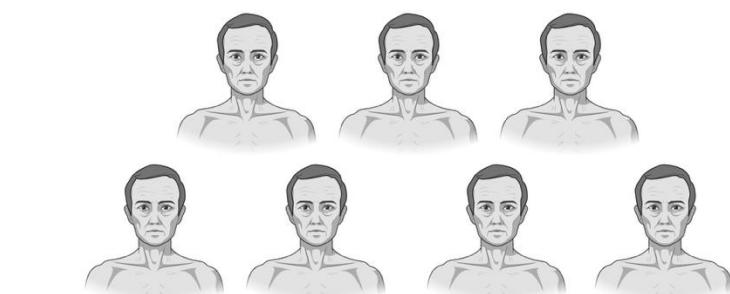
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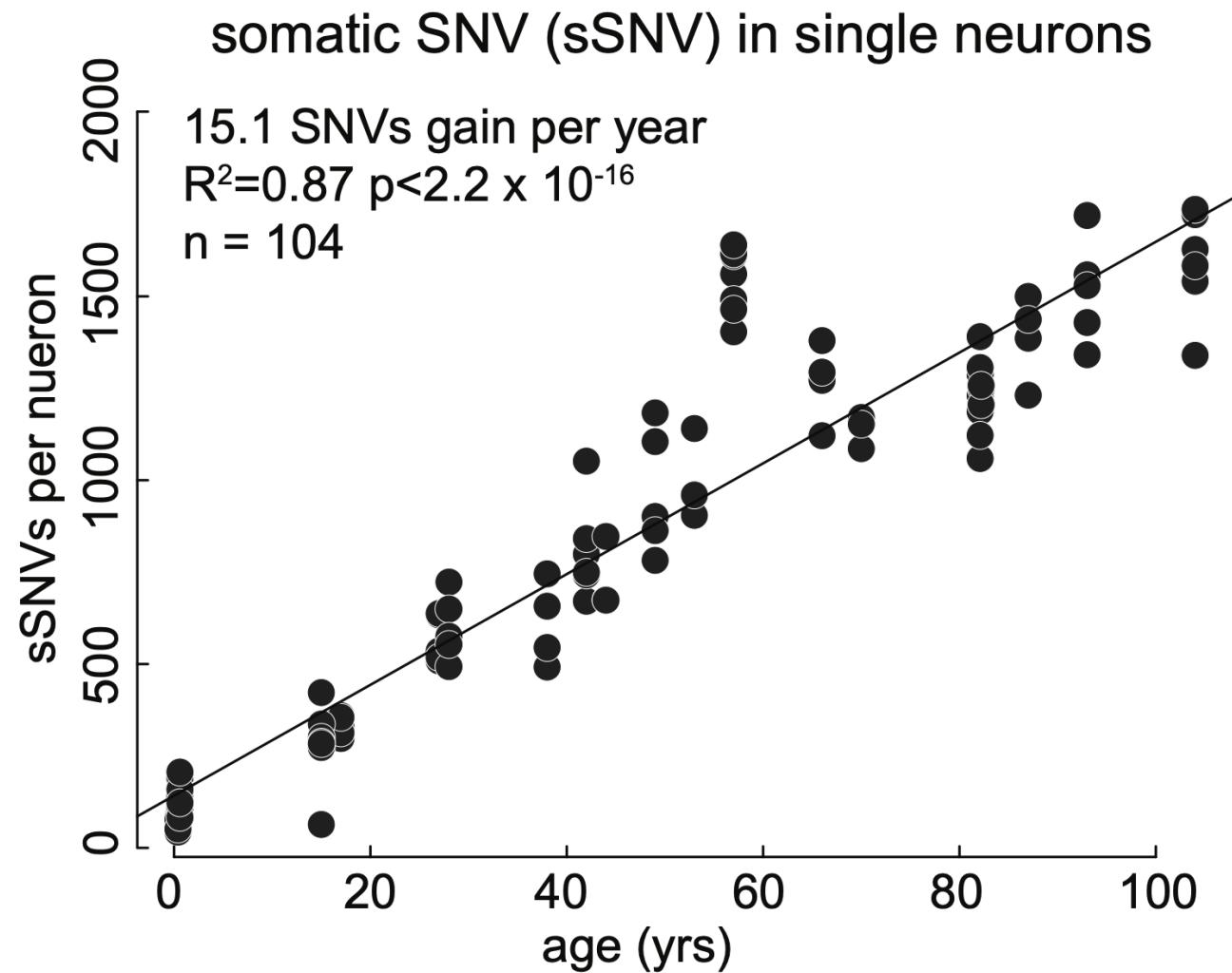


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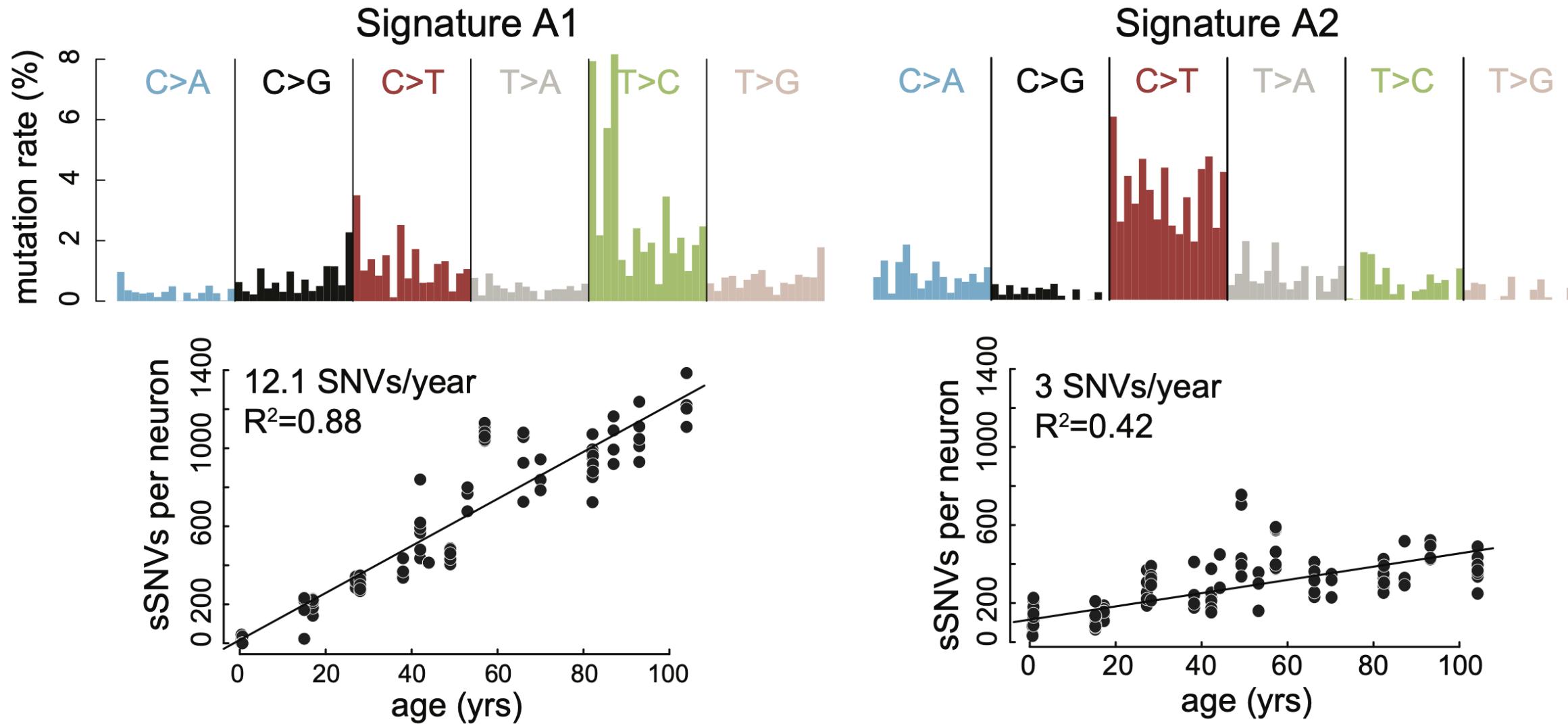


How does somatic mutations accumulate in neurons during aging?

Somatic SNV accumulate in neurons



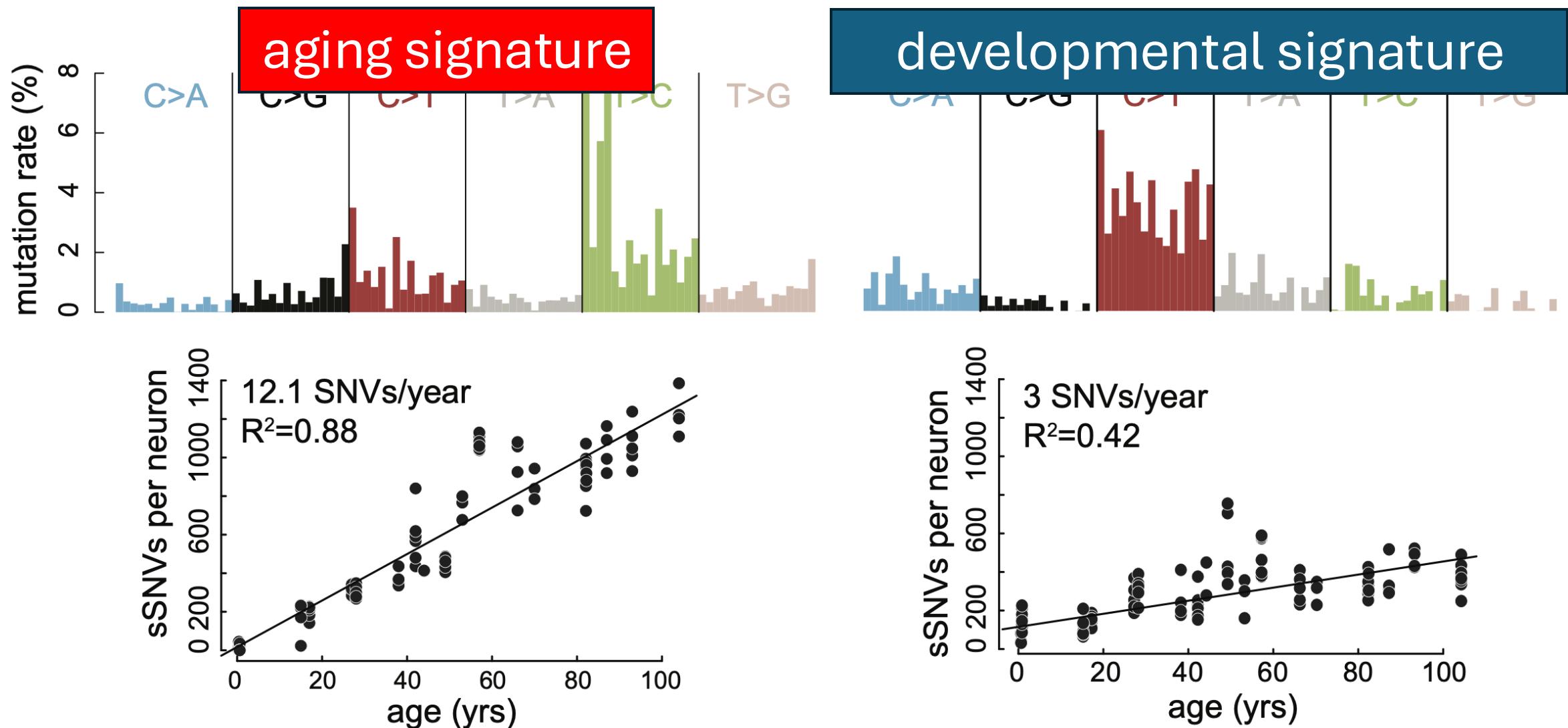
A signature associated with aging



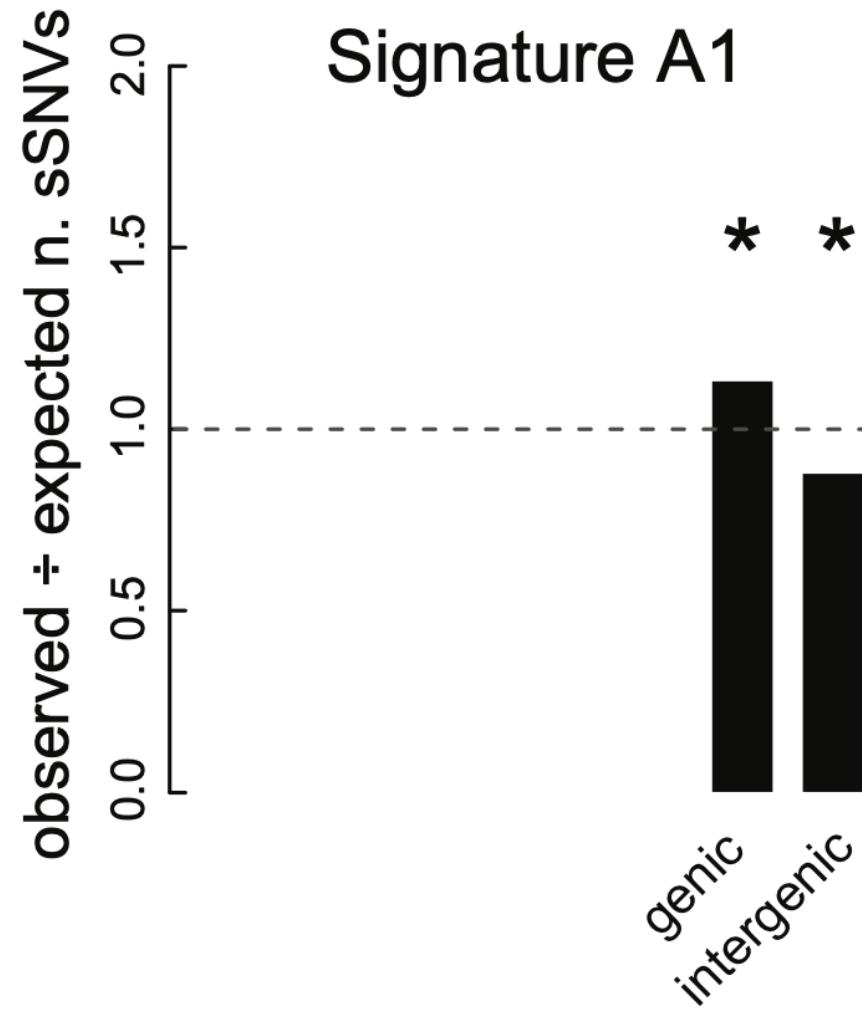
Signatures are fingerprints of certain mutagens



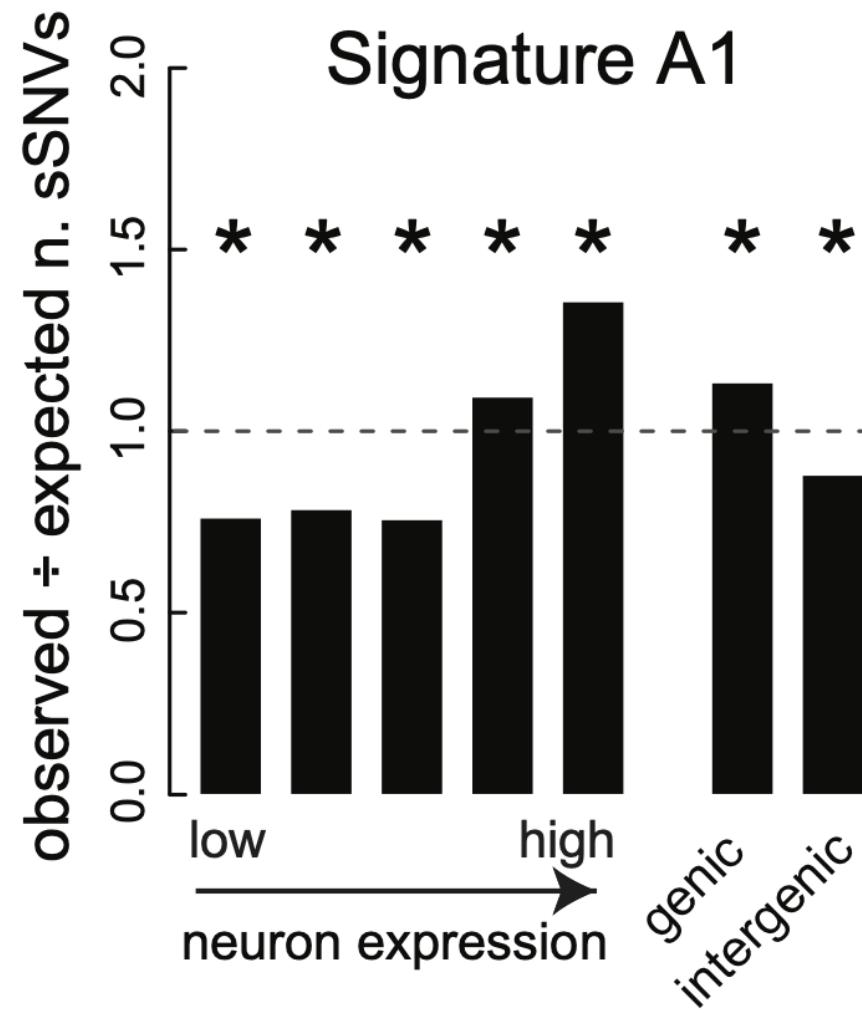
A signature associated with aging



Aging signature enrich at active genes

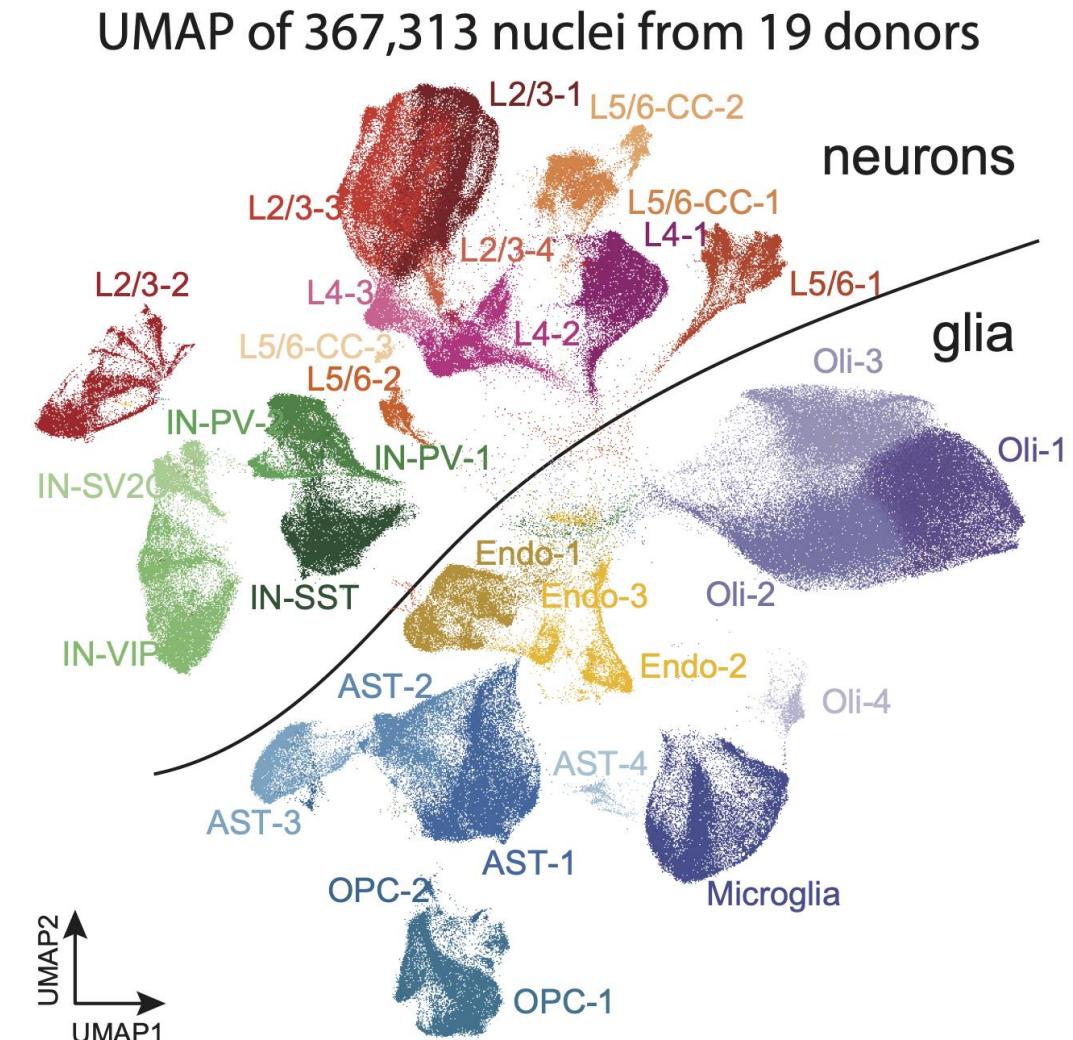


Aging signature enrich at active genes

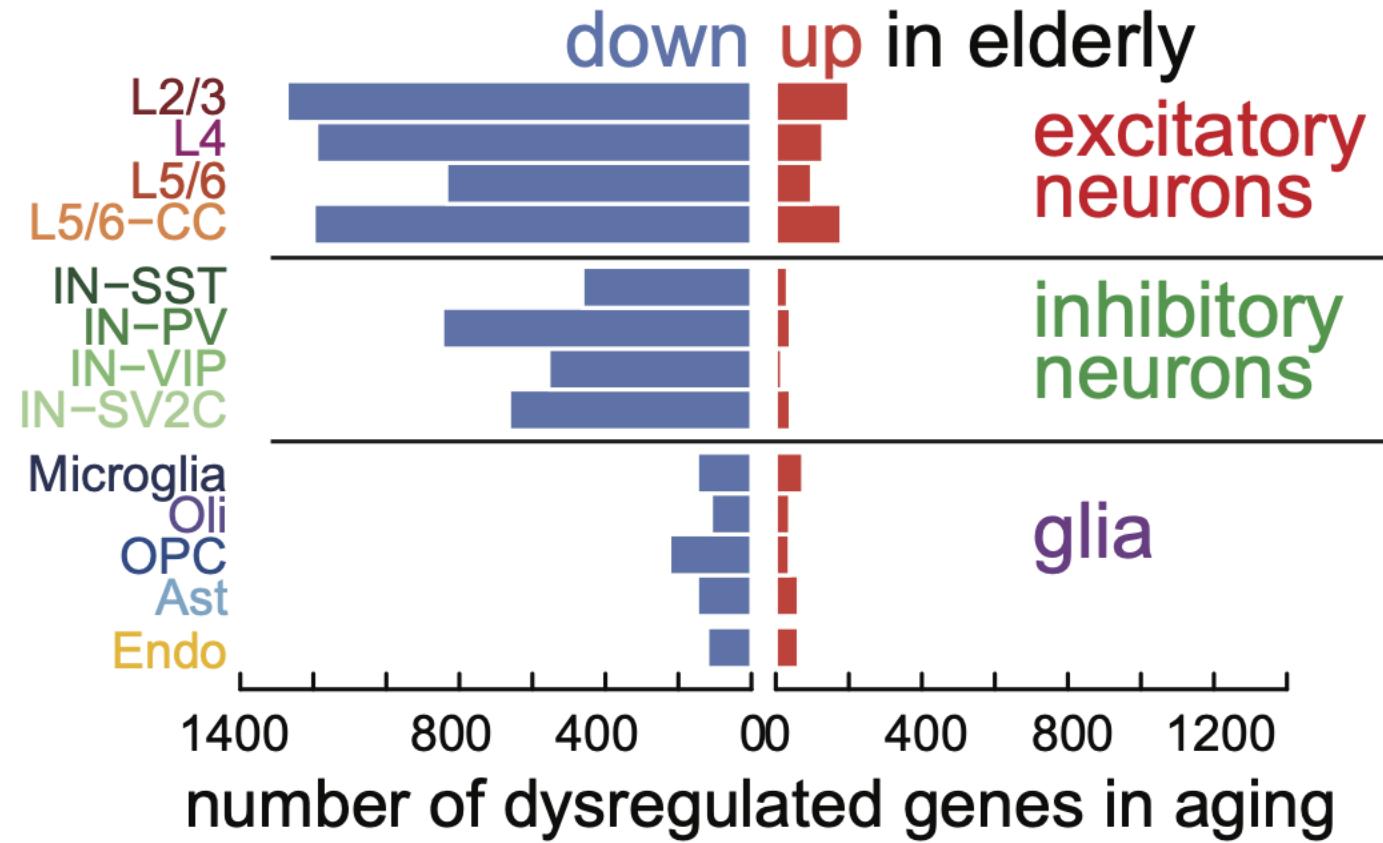


How does transcriptome change in PFC
during aging?

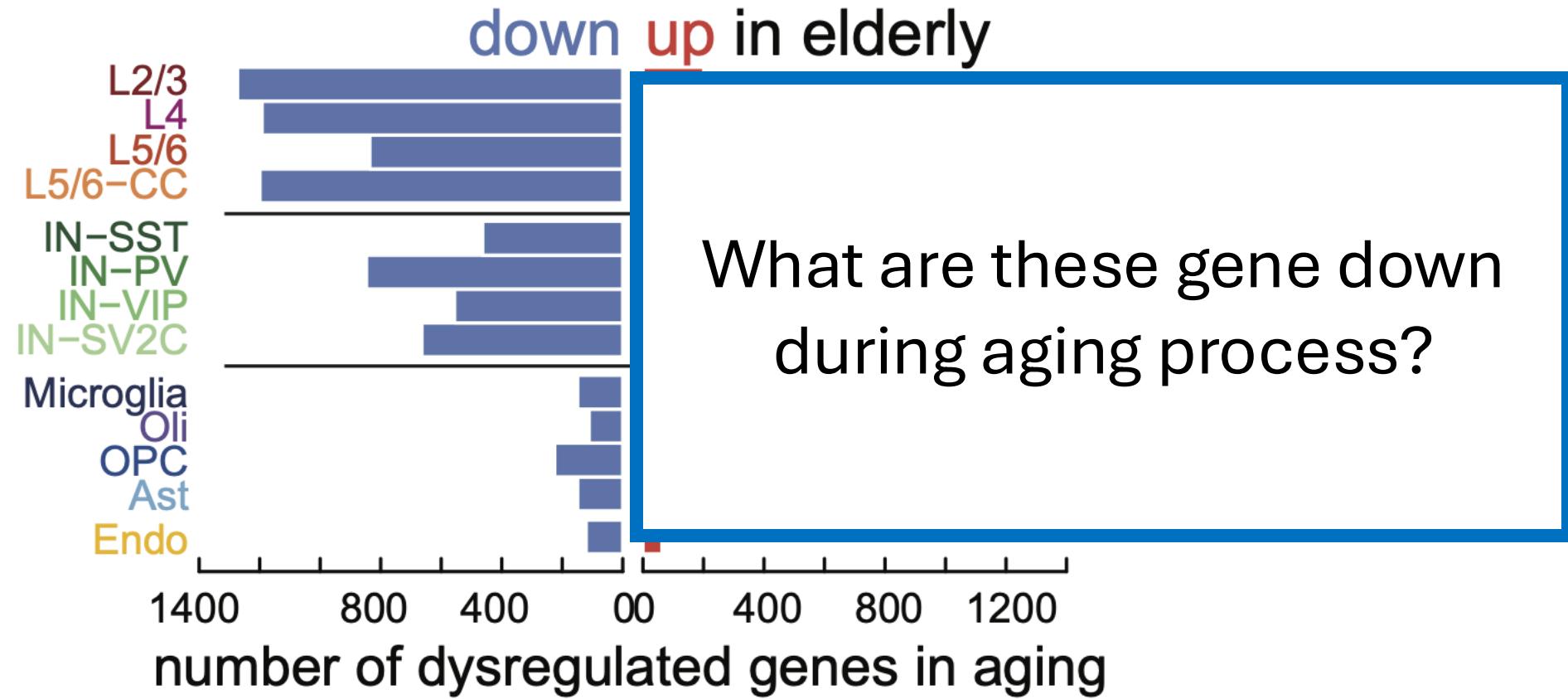
snRNA-seq captured cells in PFC



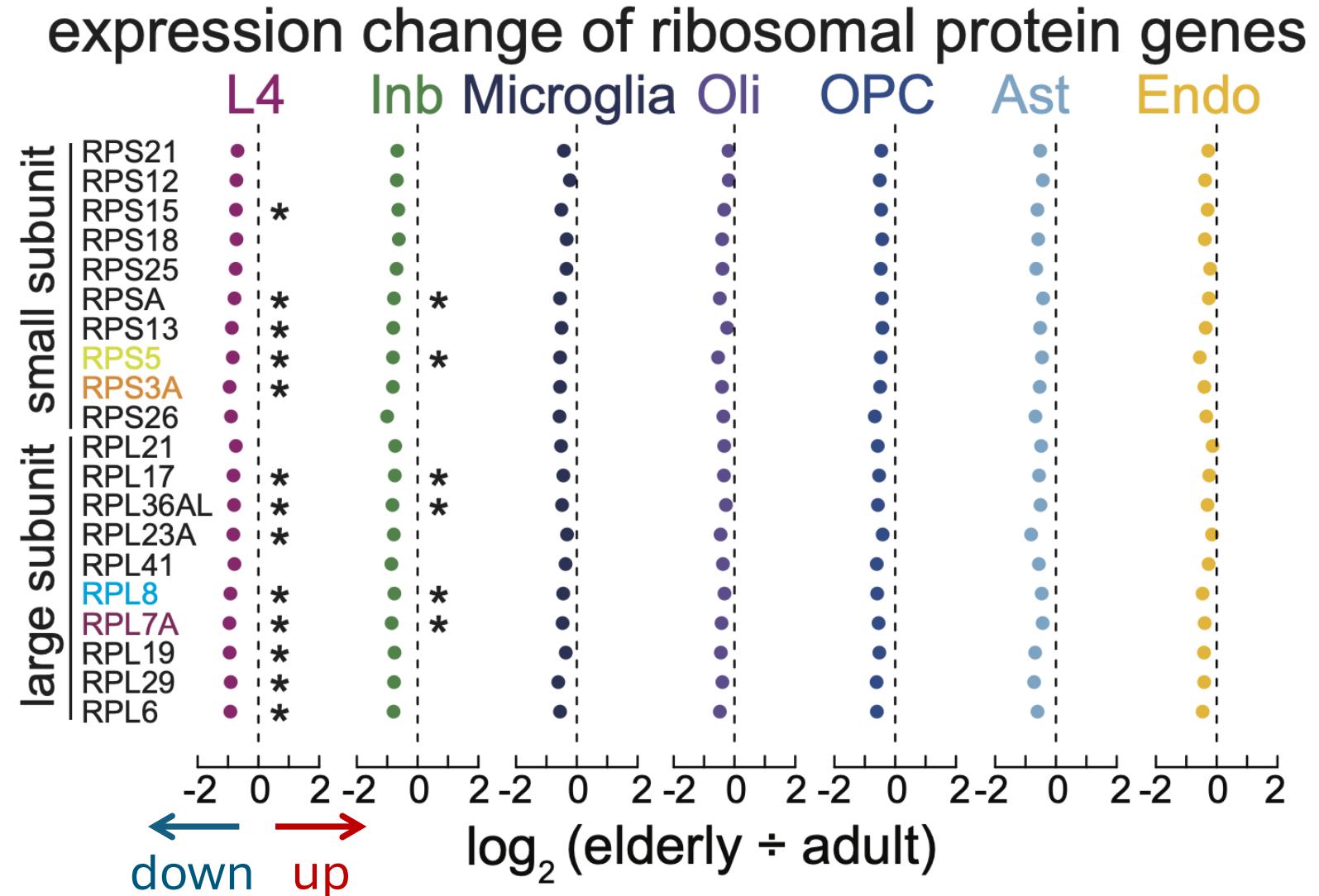
Many genes are commonly down during aging



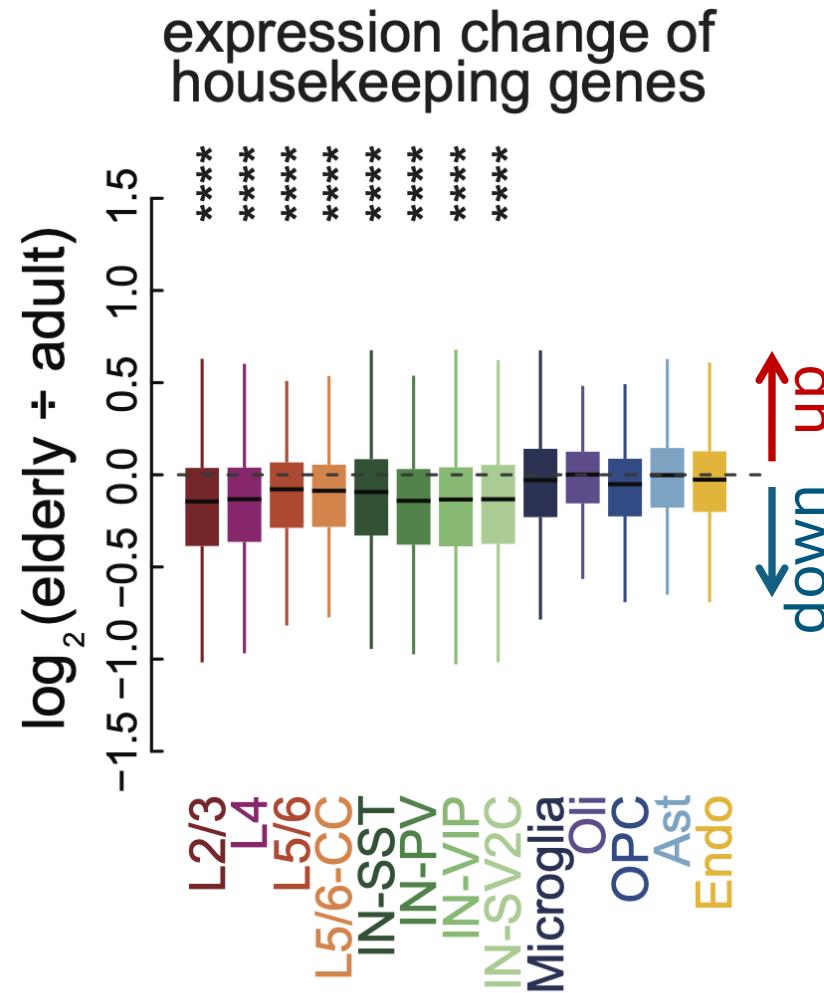
Many genes are commonly down during aging



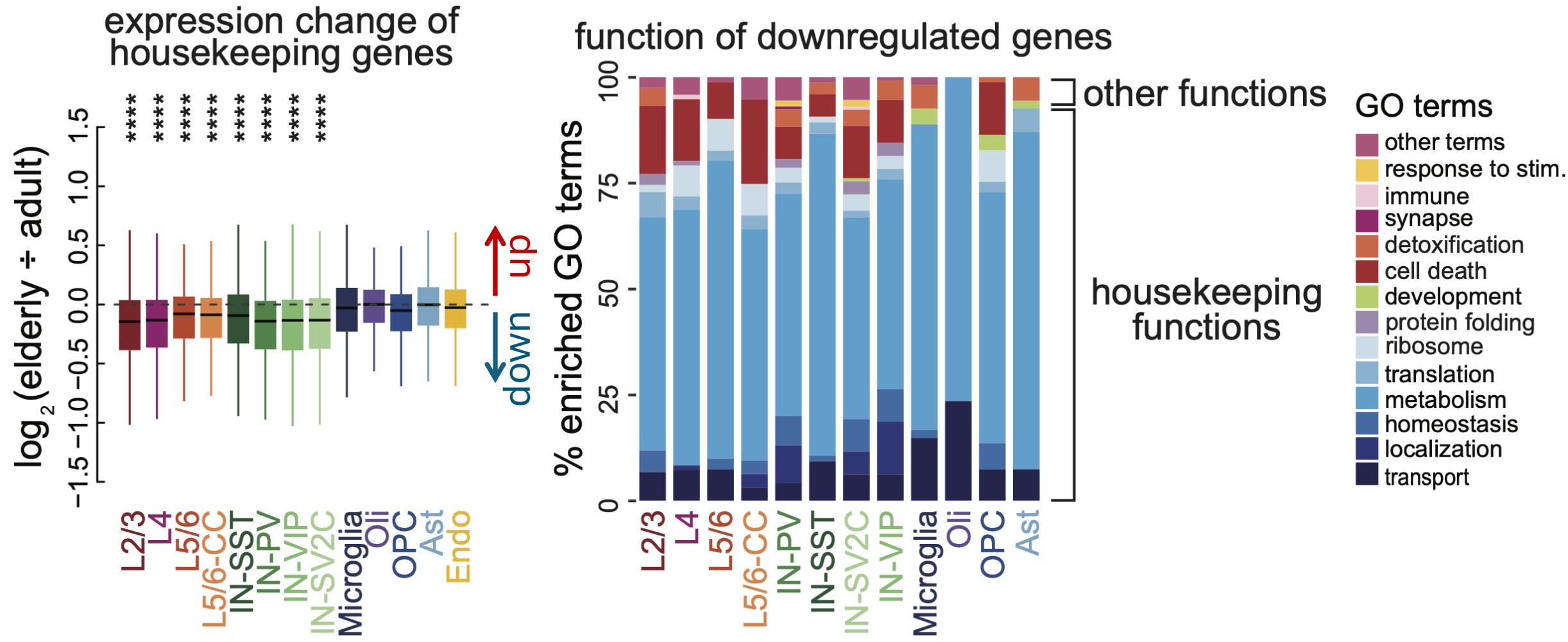
Common decrease of housekeeping program



Common decrease of housekeeping program

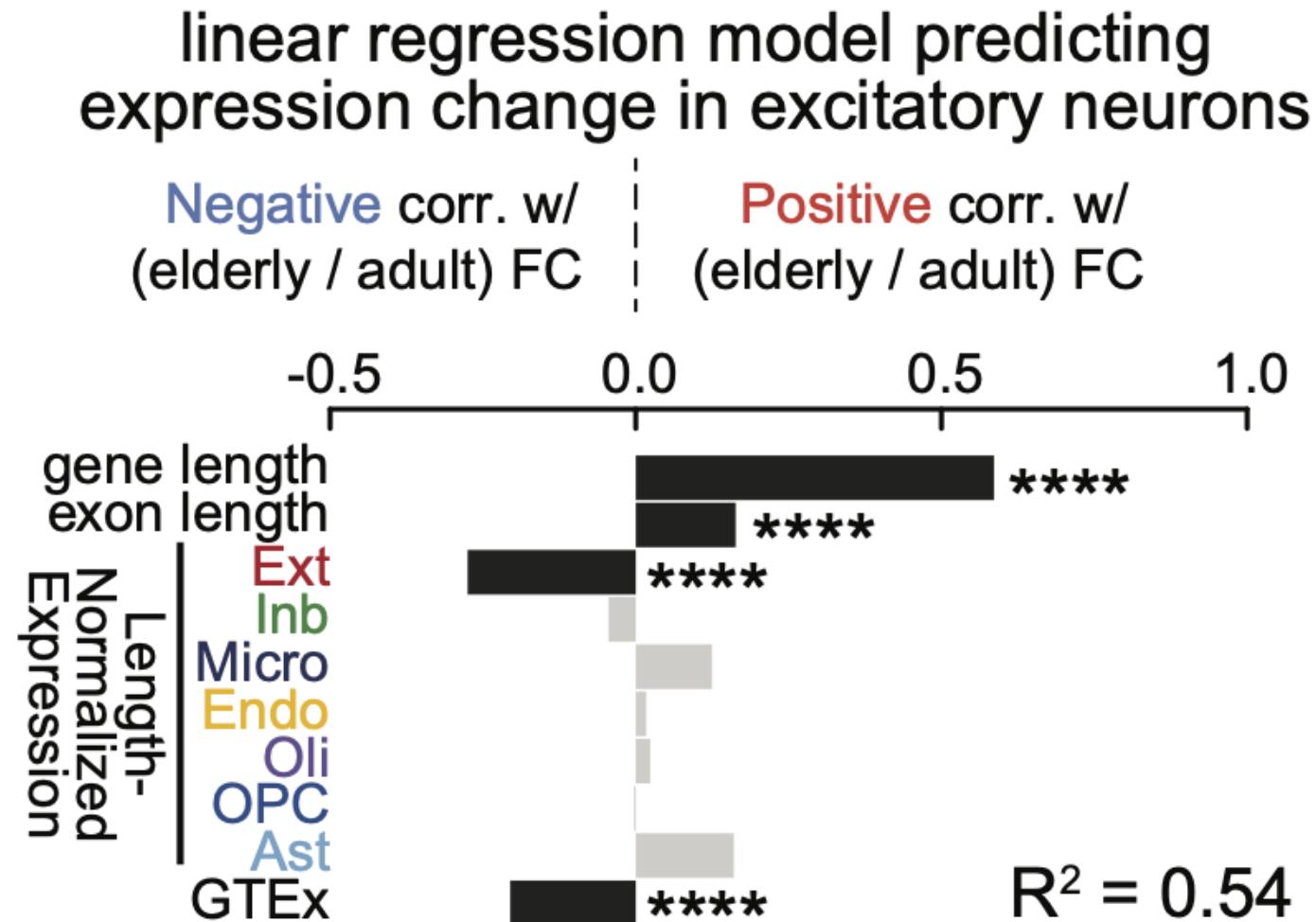


Common decrease of housekeeping program

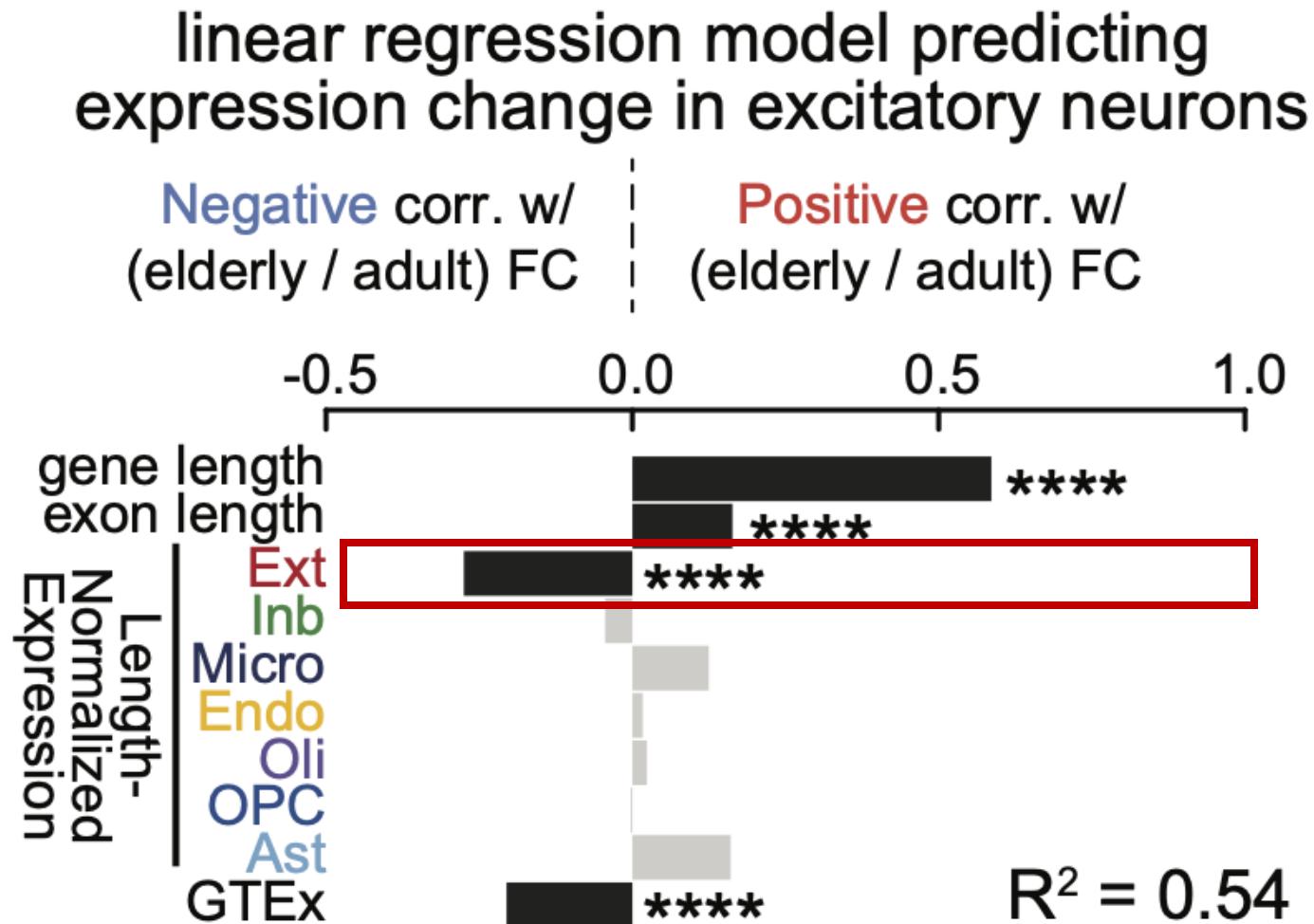


How does genomic changes link with
transcriptomic changes during PFC aging?

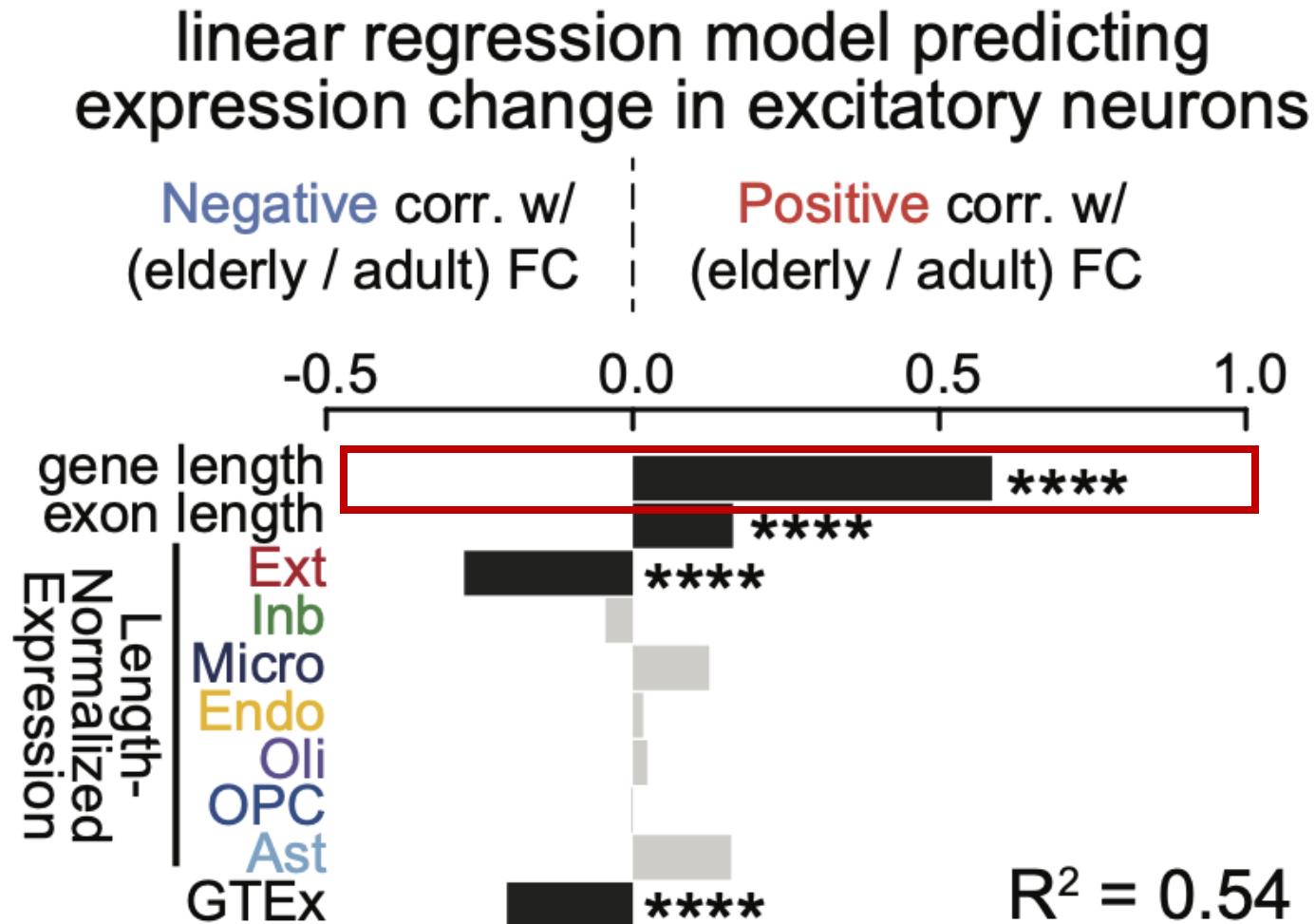
Short and active genes go down during aging



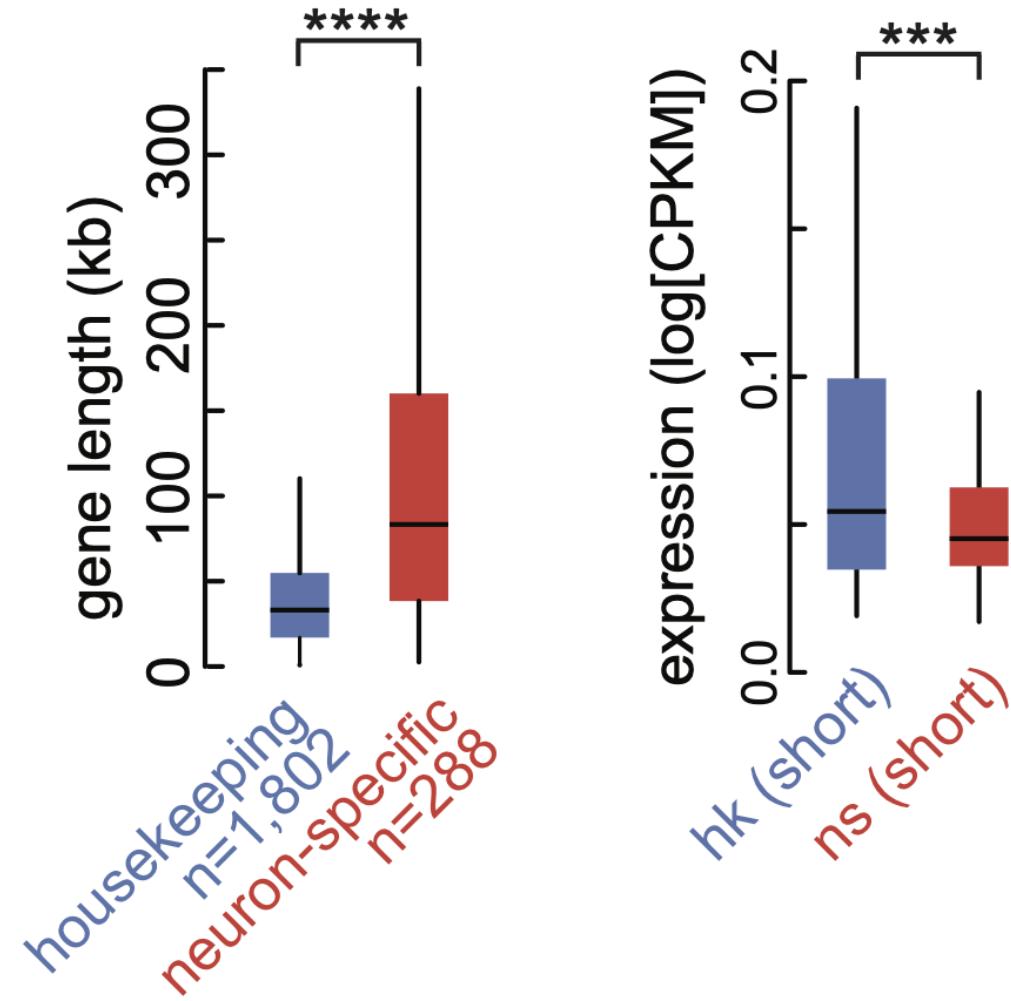
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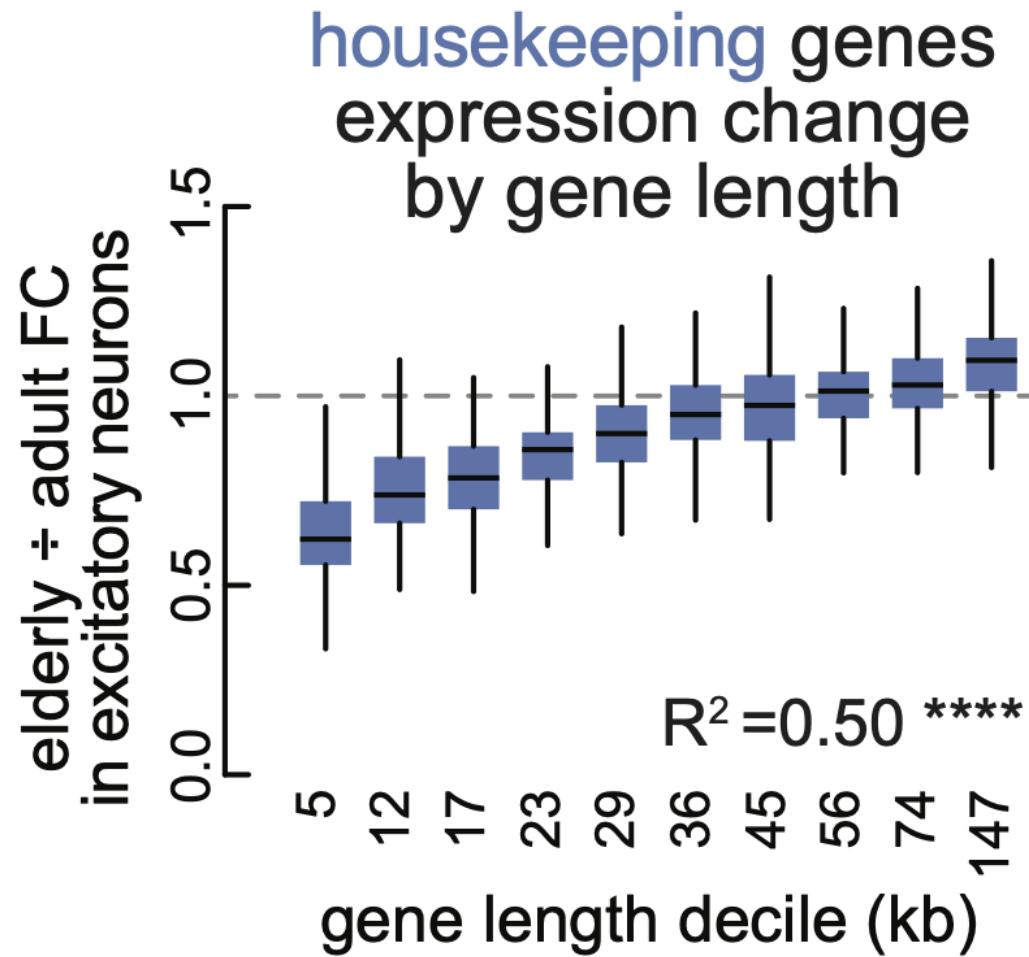
Short and active genes go down during aging



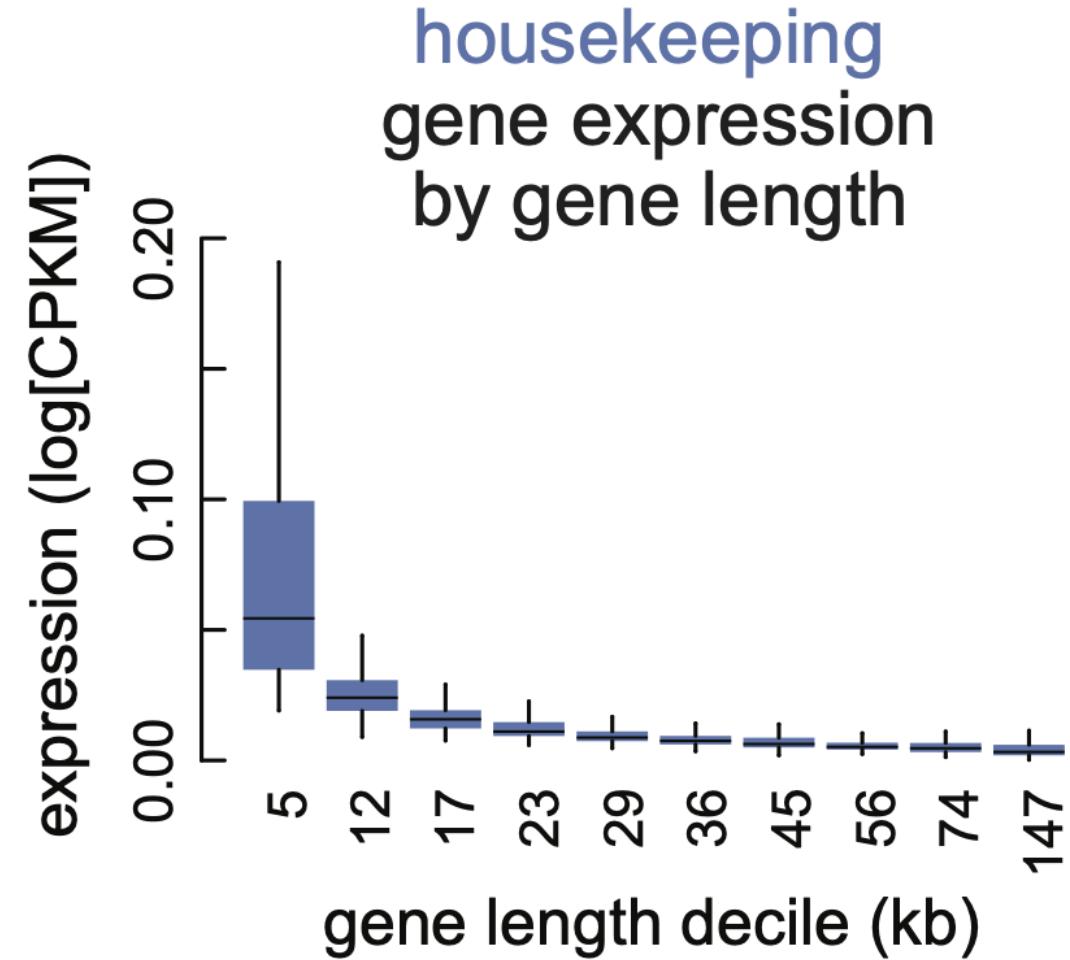
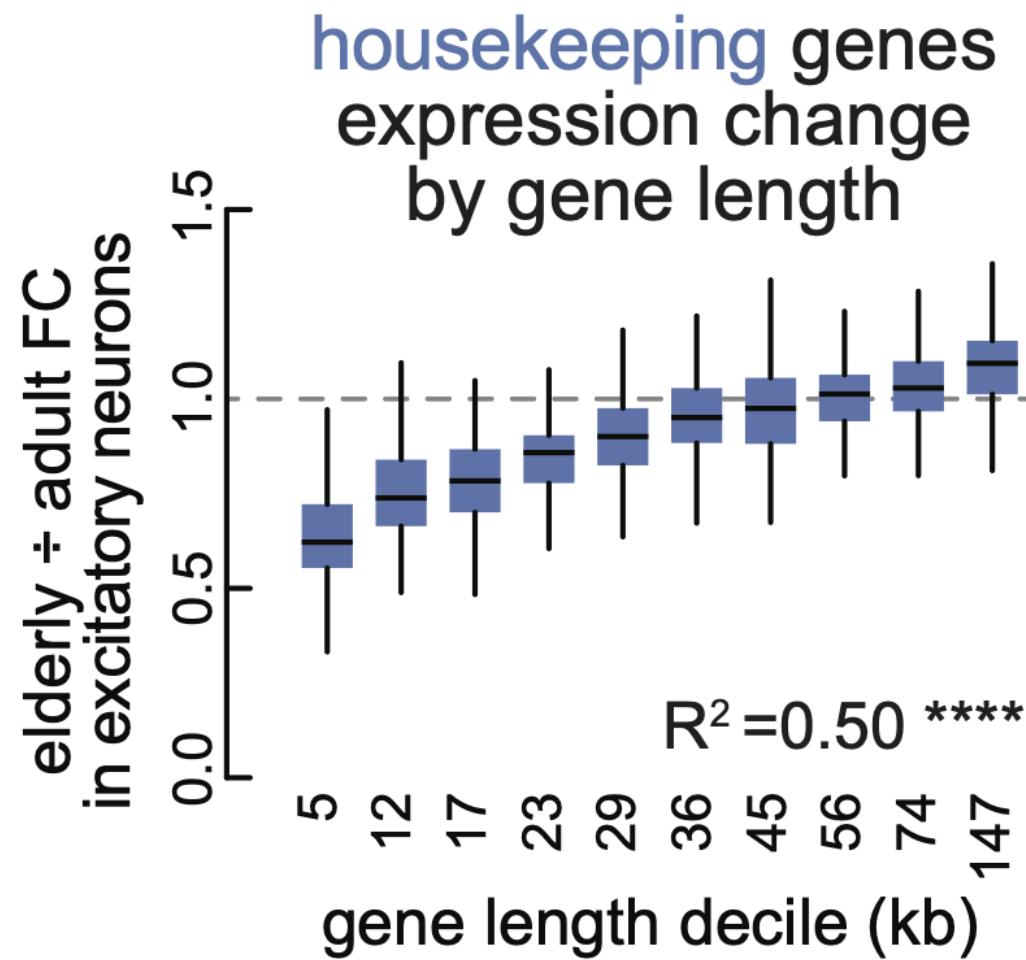
Housekeeping genes are short and active



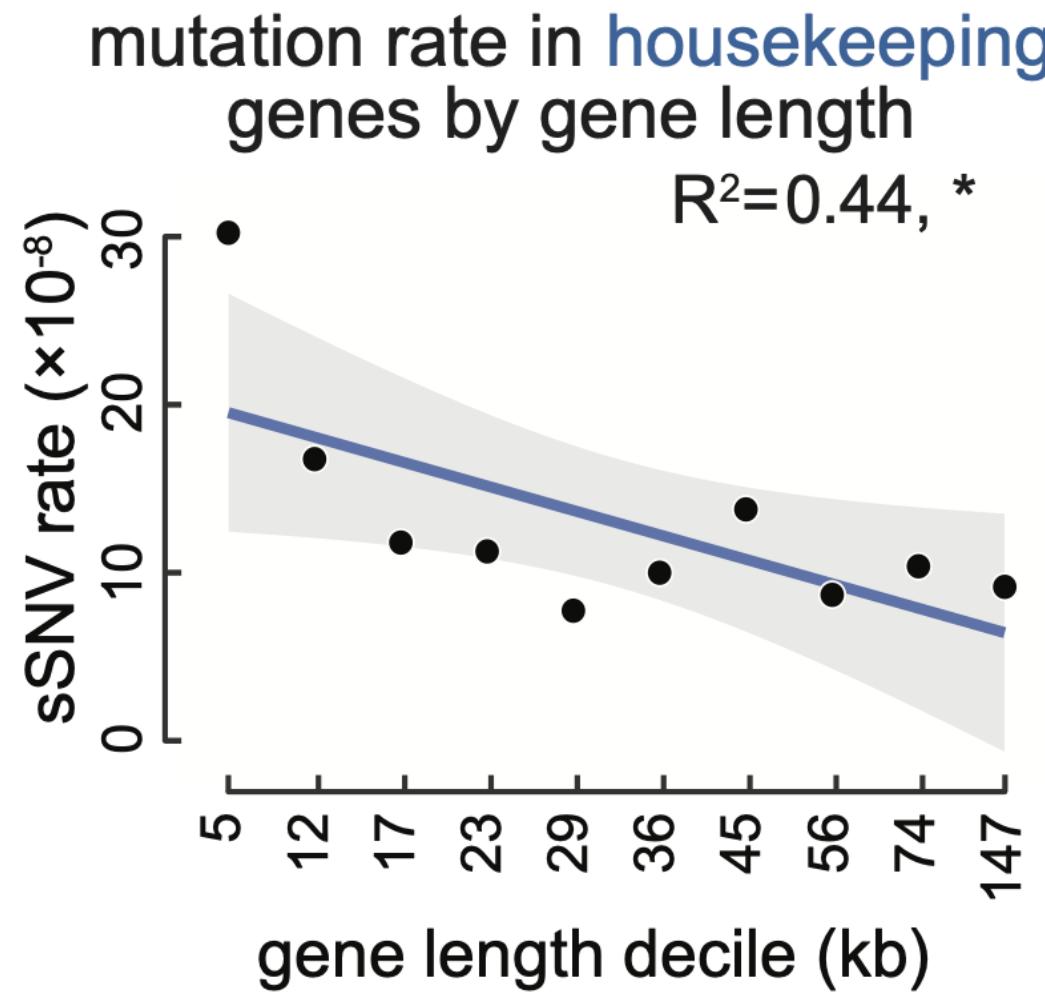
Shortest housekeeping genes down most



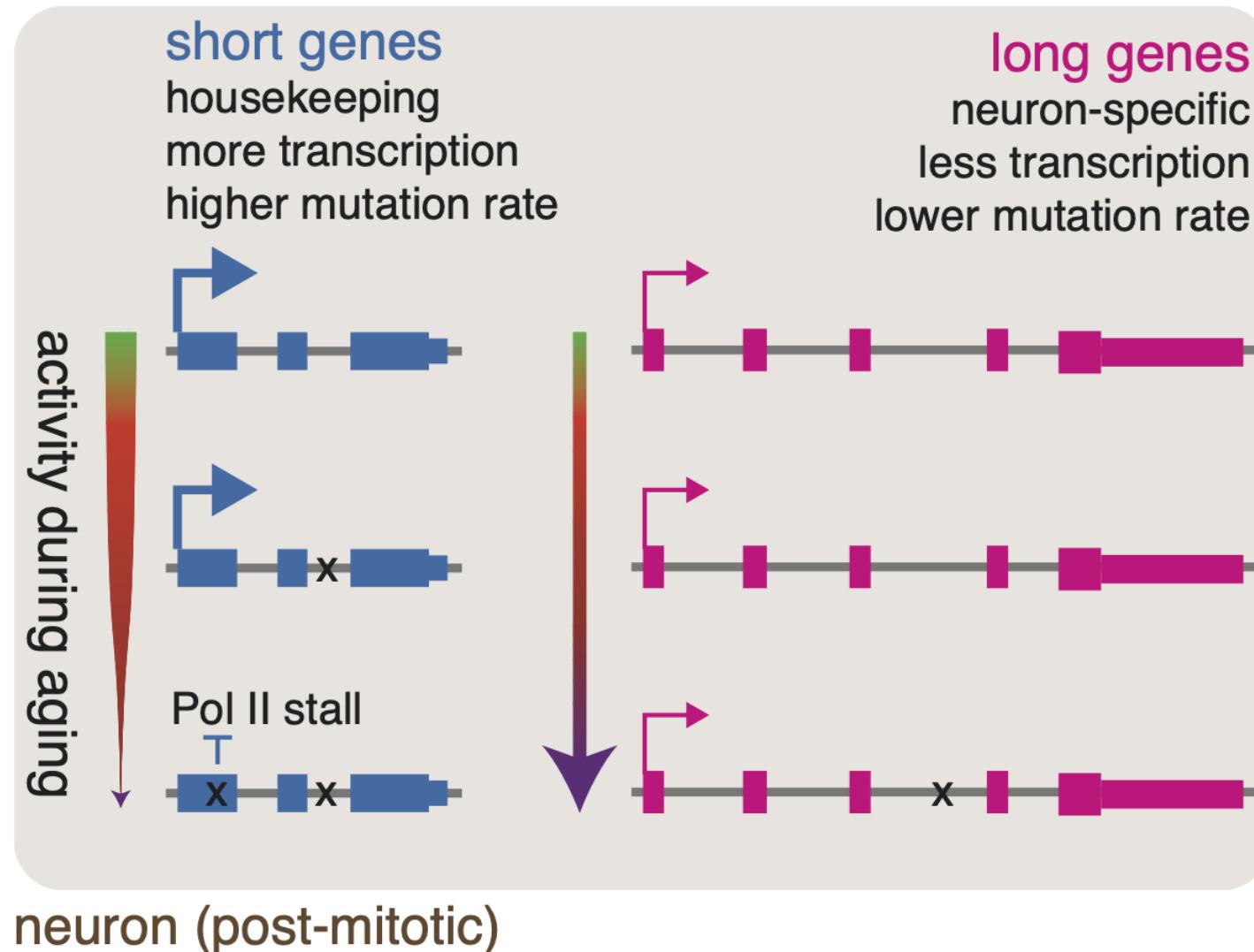
Shortest housekeeping genes down most



Short housekeeping genes bear more mutations

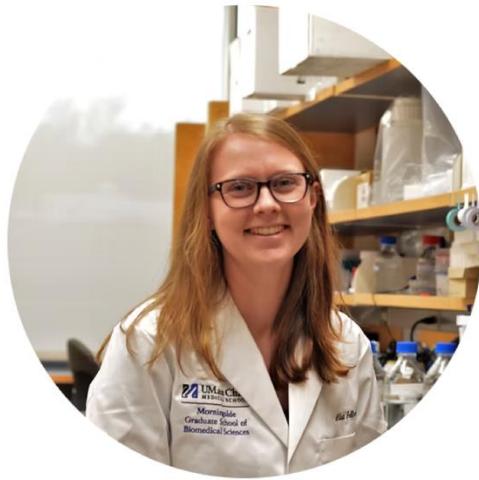


Proposed model



- Somatic SNVs accumulates in neurons with aging, faster in short and active genes.
- Short and highly active housekeeping genes are commonly downregulated in elderly human brain.
- **The more you use it, the more it wears out!**

Acknowledge



Ailsa Jeffries



Michael Lodato



Zhiping Weng

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Allie Tolles
Christina Baer
Cesar Sotelo
Yerin Kim

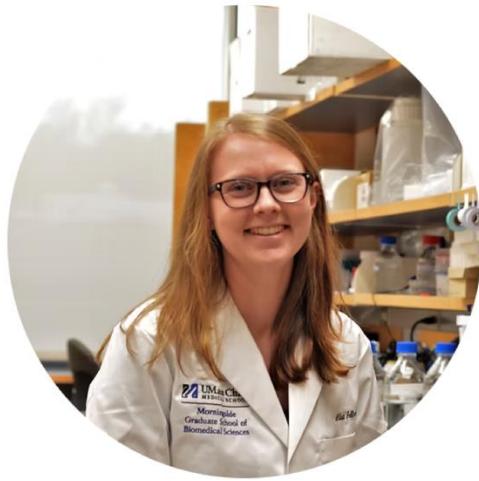


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