

Unique features of functional connectivity fingerprints during aging

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Sample



N ≈ 550

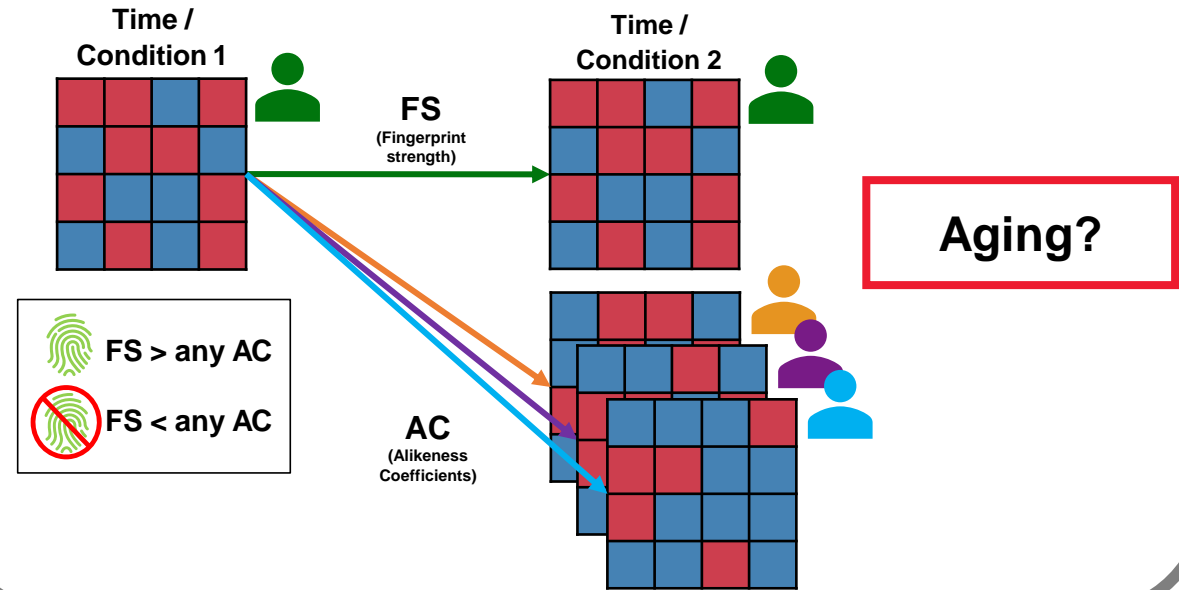


51% female

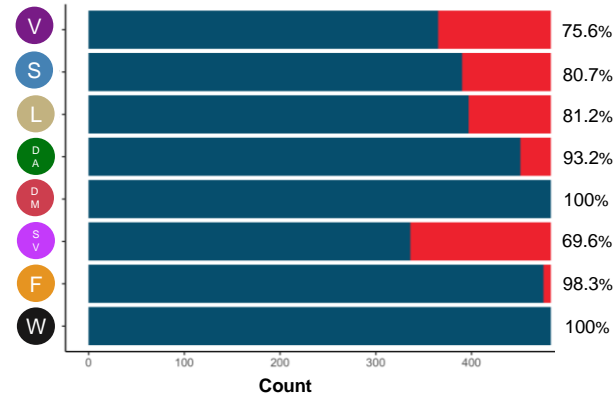


18-87 years
 (50.39 avg)

Fingerprinting methodology



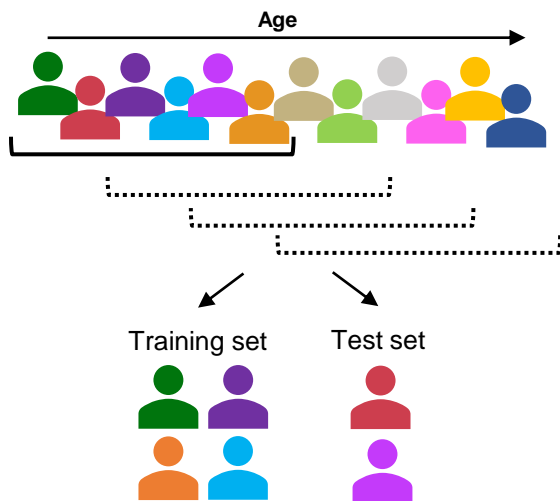
Rest – Task (n = 483)



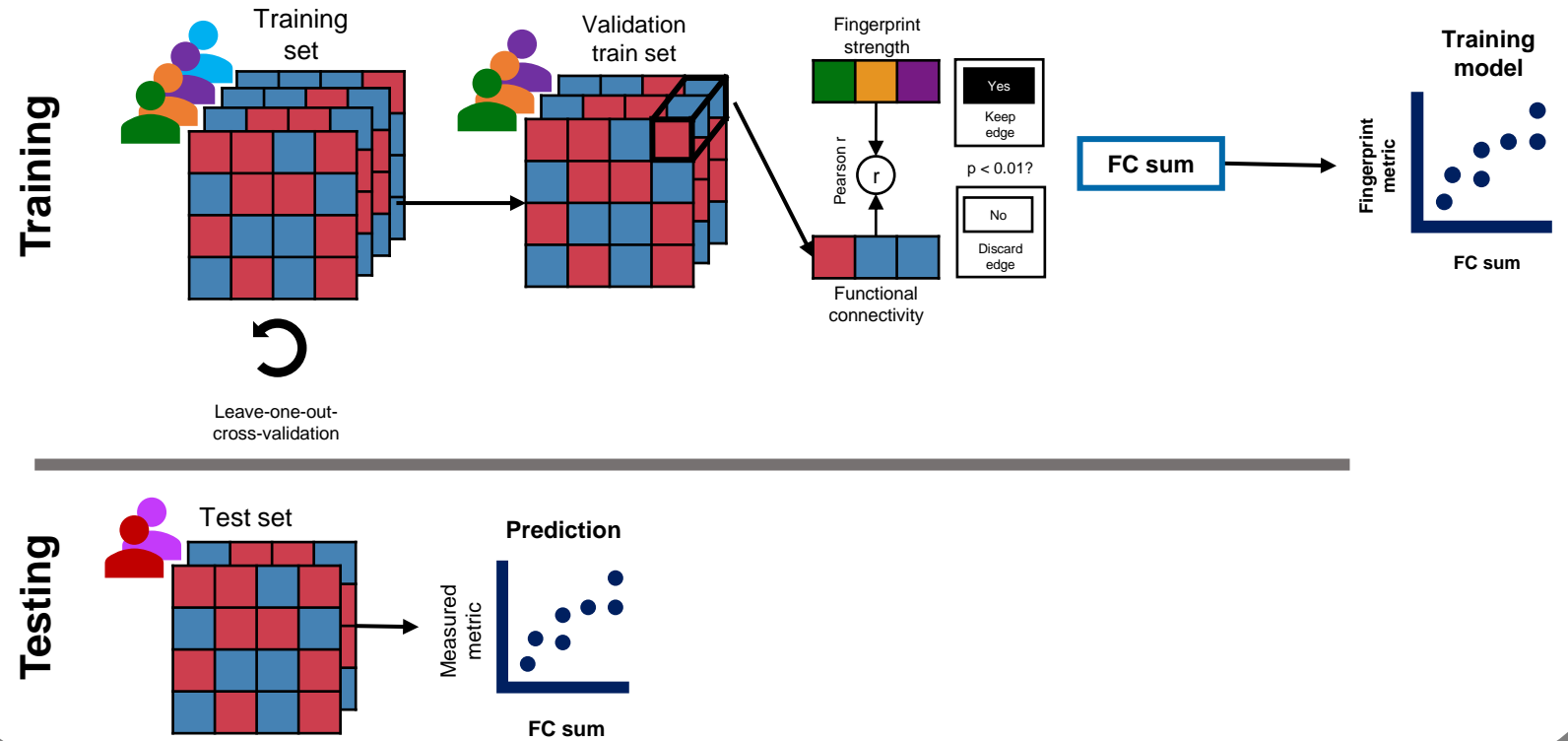
Functional connectivity patterns are highly unique during aging

Which brain regions are unique?
Do they change with age?

Sliding-window approach



Connectome predictive modelling



Model performance

Predicted brain regions

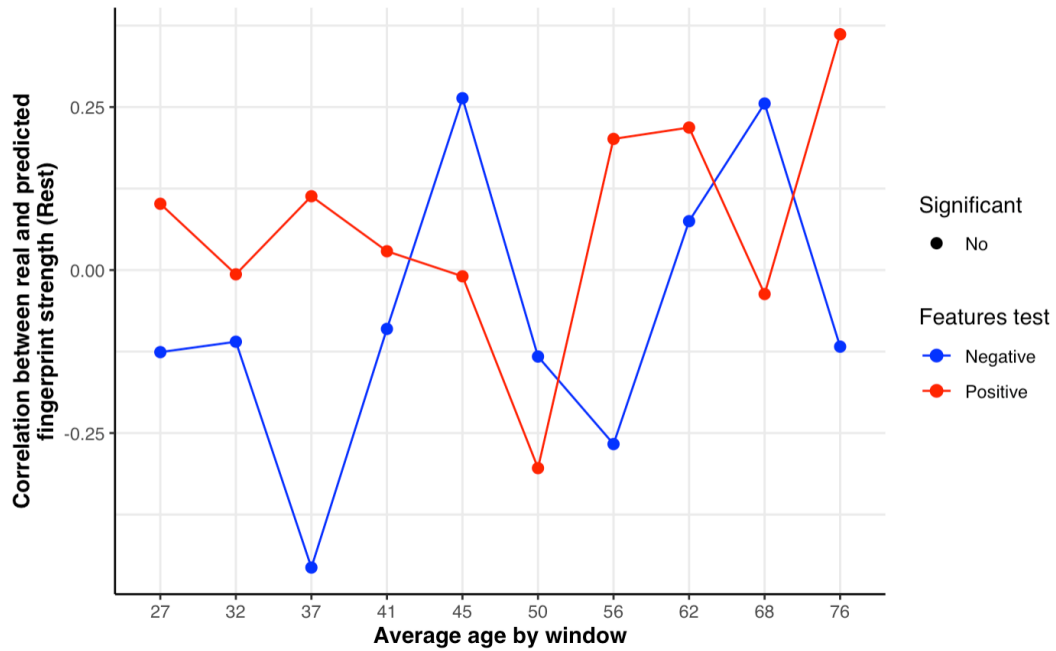


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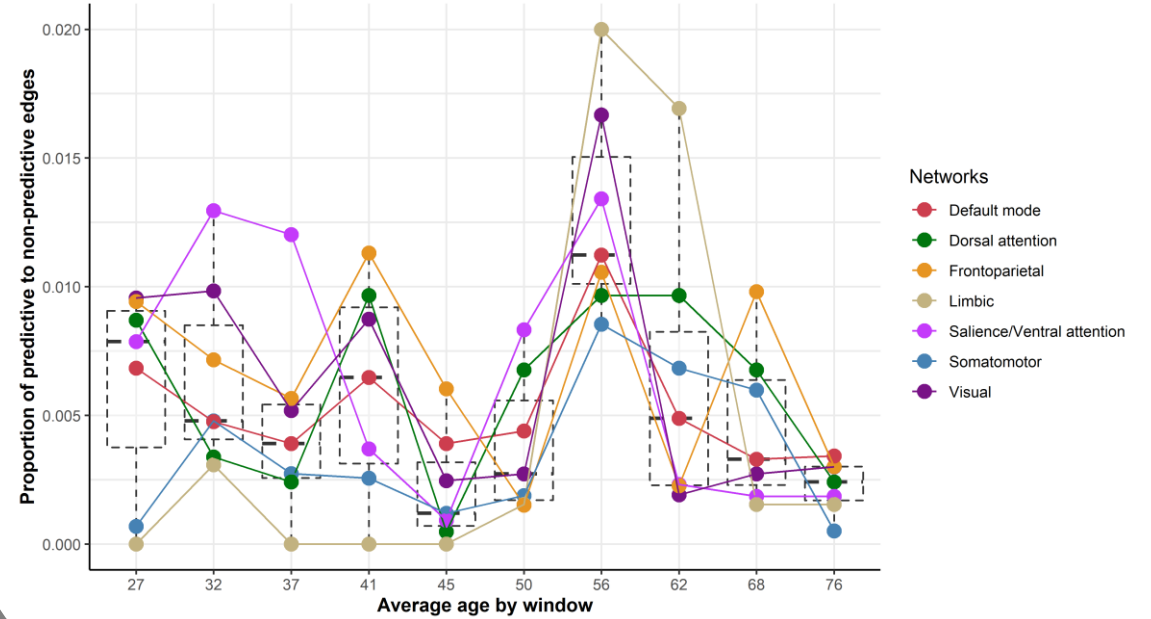


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Model performance



Associated brain regions



Not only are fingerprints unique,
but there is **no “one-region-fits-all”** that can predict fingerprints
across individuals AND across the lifespan

Acknowledgements



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