

# Human cortex development is shaped by molecular and cellular brain systems

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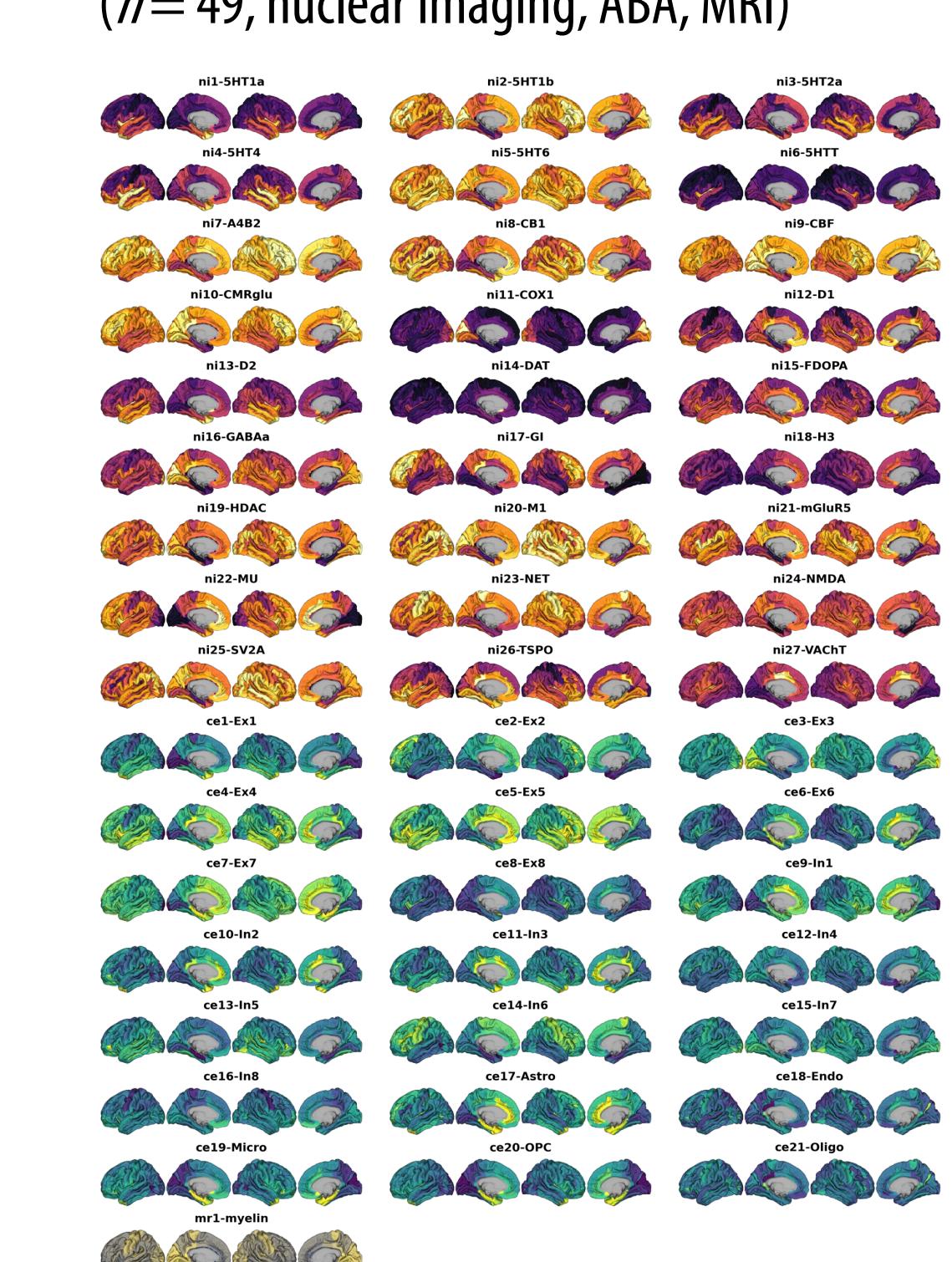
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## Research motivation & impact

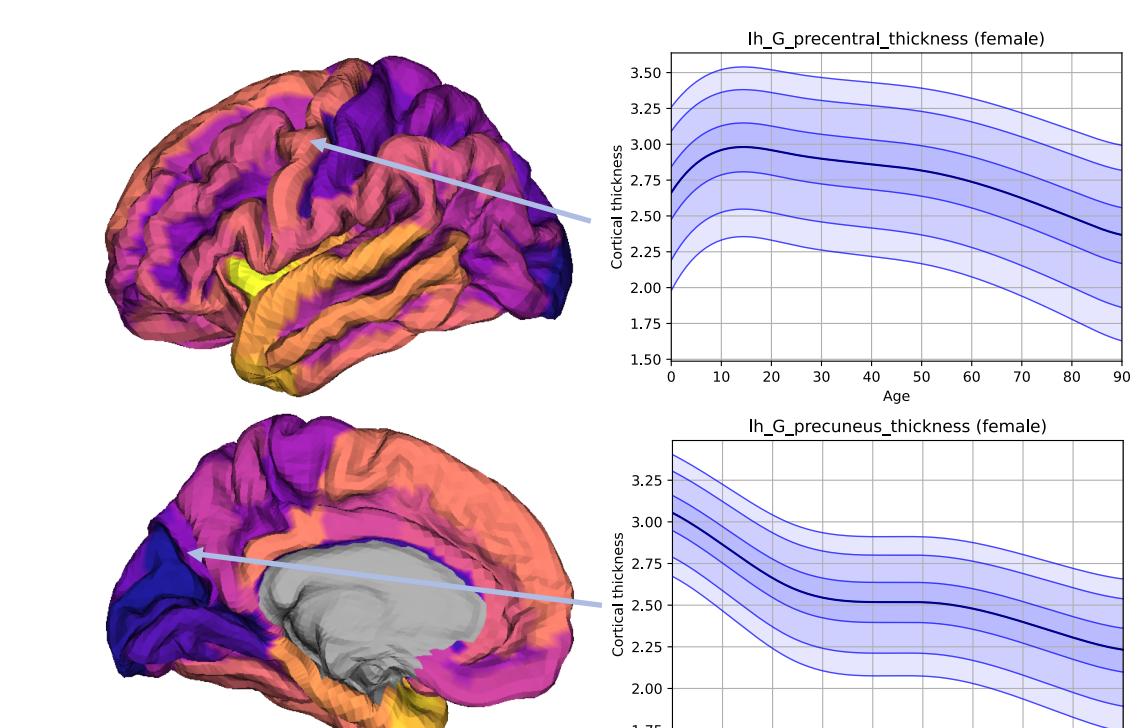
- Human cerebral cortex morphology is subject to complex and diverse changes over the lifespan<sup>1,2</sup>.
- Several factors might influence cortical thickness (CT) development and maturation, but human data are scarce.
- Through spatial correlation approaches<sup>3,4</sup>, recent advances in normative modeling of population-scale neuroimaging data<sup>1,2</sup>, and availability of multi-level atlases of molecular and cellular brain systems<sup>5-7</sup>, we identify potential mechanisms underlying human CT development.
- This work...
  - 1) provides new hypotheses on mechanisms involved in human cortex development,
  - 2) introduces a framework for studying neurodevelopmental mechanisms *in vivo* on the individual level, promising new insights into typical and atypical neurodevelopment alike, and
  - 3) further emphasizes the value of normative modeling frameworks in neurodevelopmental research.

## Data sources

"Multilevel" maps of molecular and cellular brain systems<sup>5-7</sup>  
(n = 49, nuclear imaging, ABA, MRI)



Normative models of "representative" CT development<sup>2</sup>  
(n = 58,836; ~2 to 100 years; 148 regions)



ABCD longitudinal CT data<sup>8</sup>  
(n = 6,789; ~10 and 12 years)

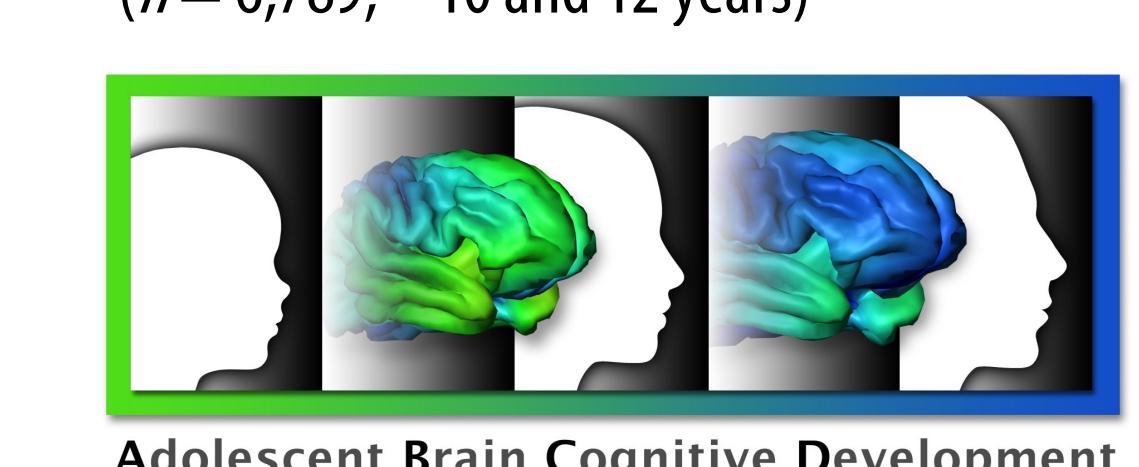
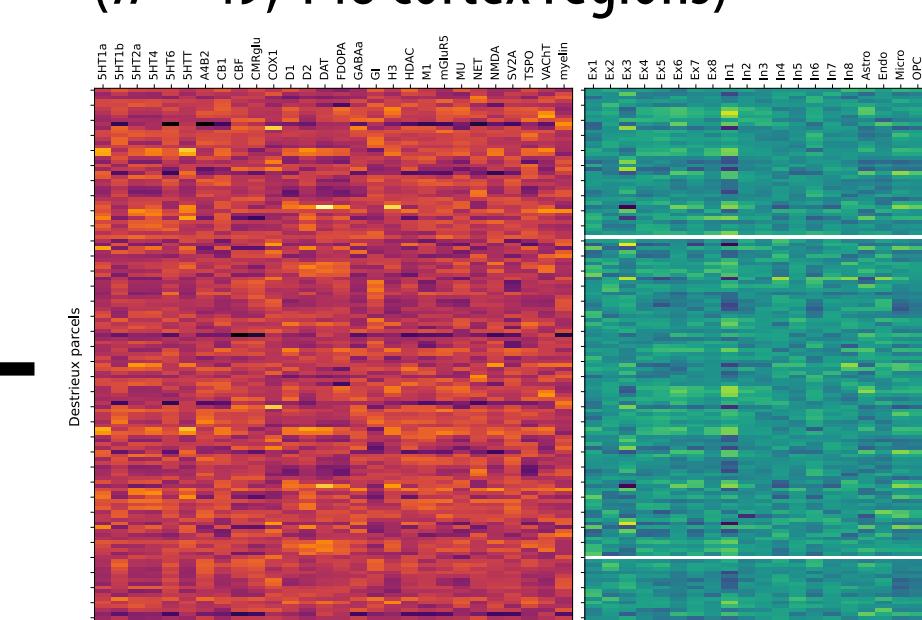


IMAGEN longitudinal CT data<sup>9</sup>  
(n = 995 to 1,278; ~14, 19, and 22 years)

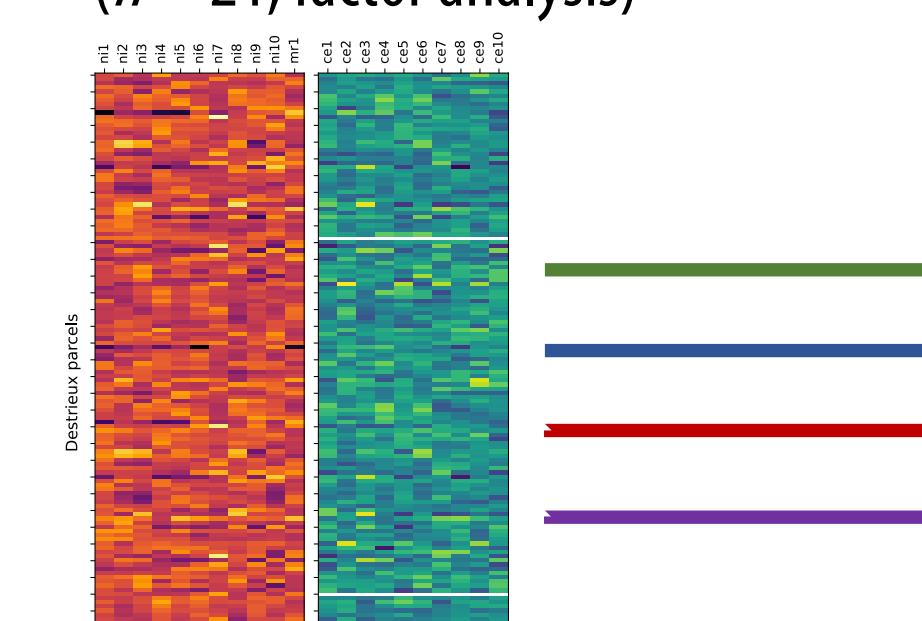


## Data processing

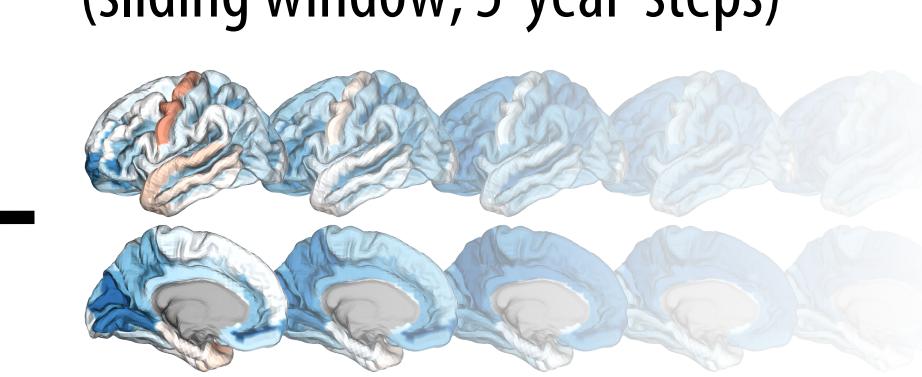
Parcellation  
(n = 49; 148 cortex regions)



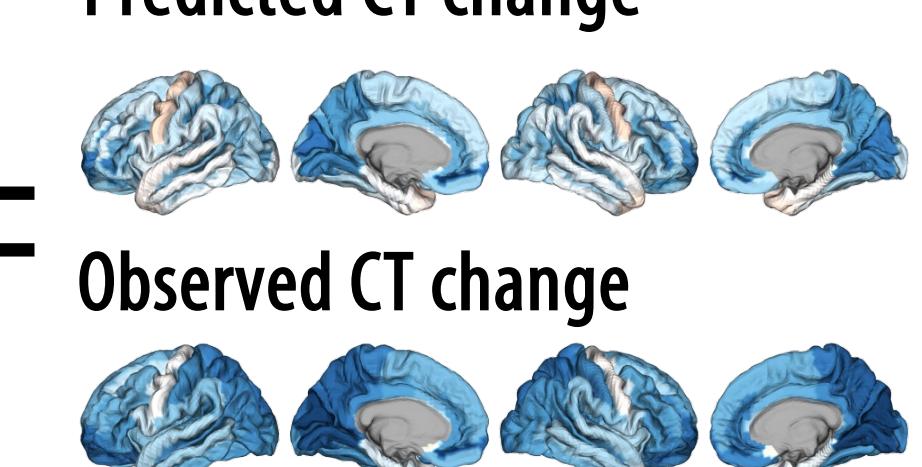
Dimensionality reduction  
(n = 21; factor analysis)



Timestep-wise CT change  
(sliding window, 5-year-steps)



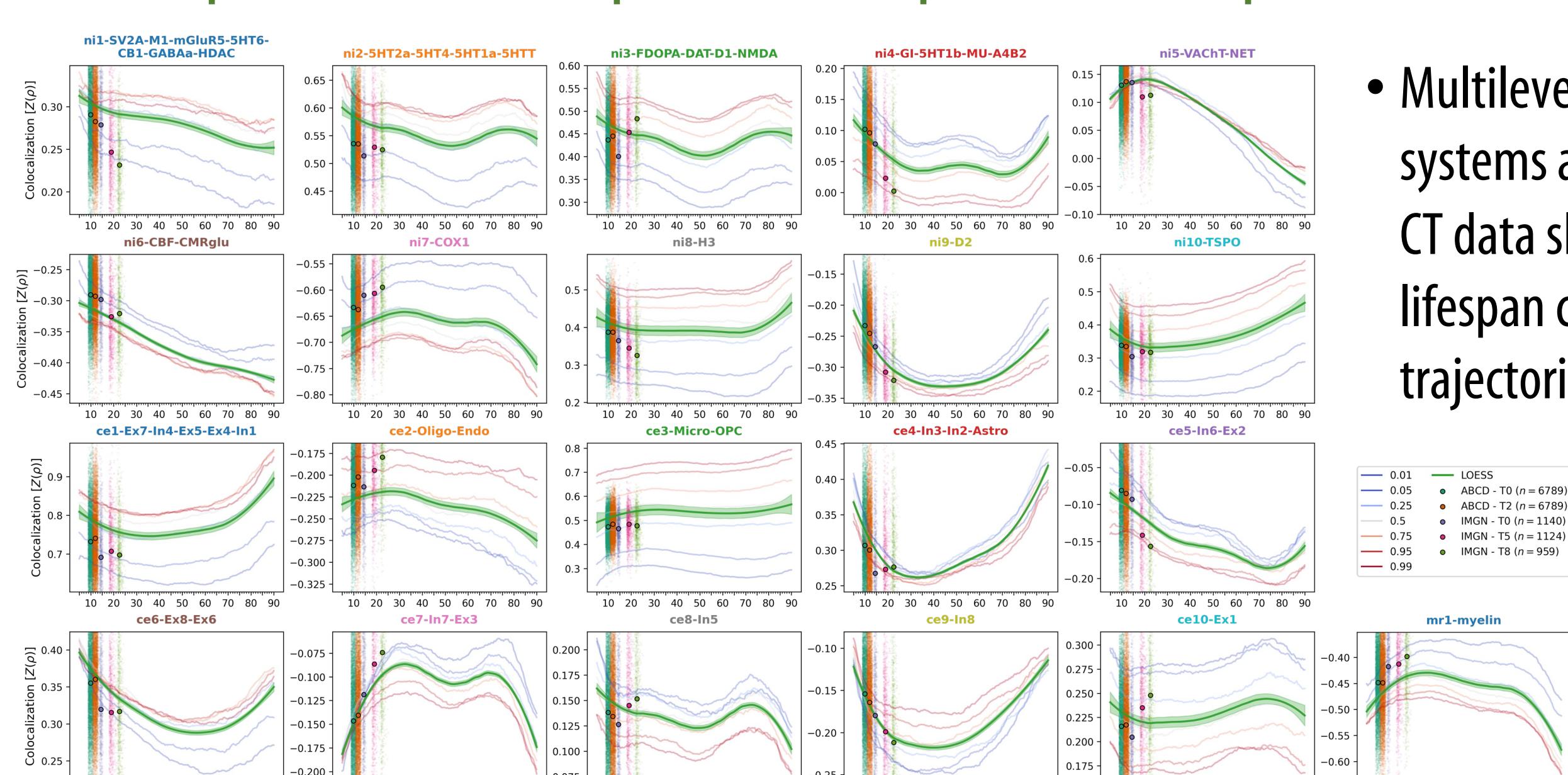
Subject-wise:  
Predicted CT change



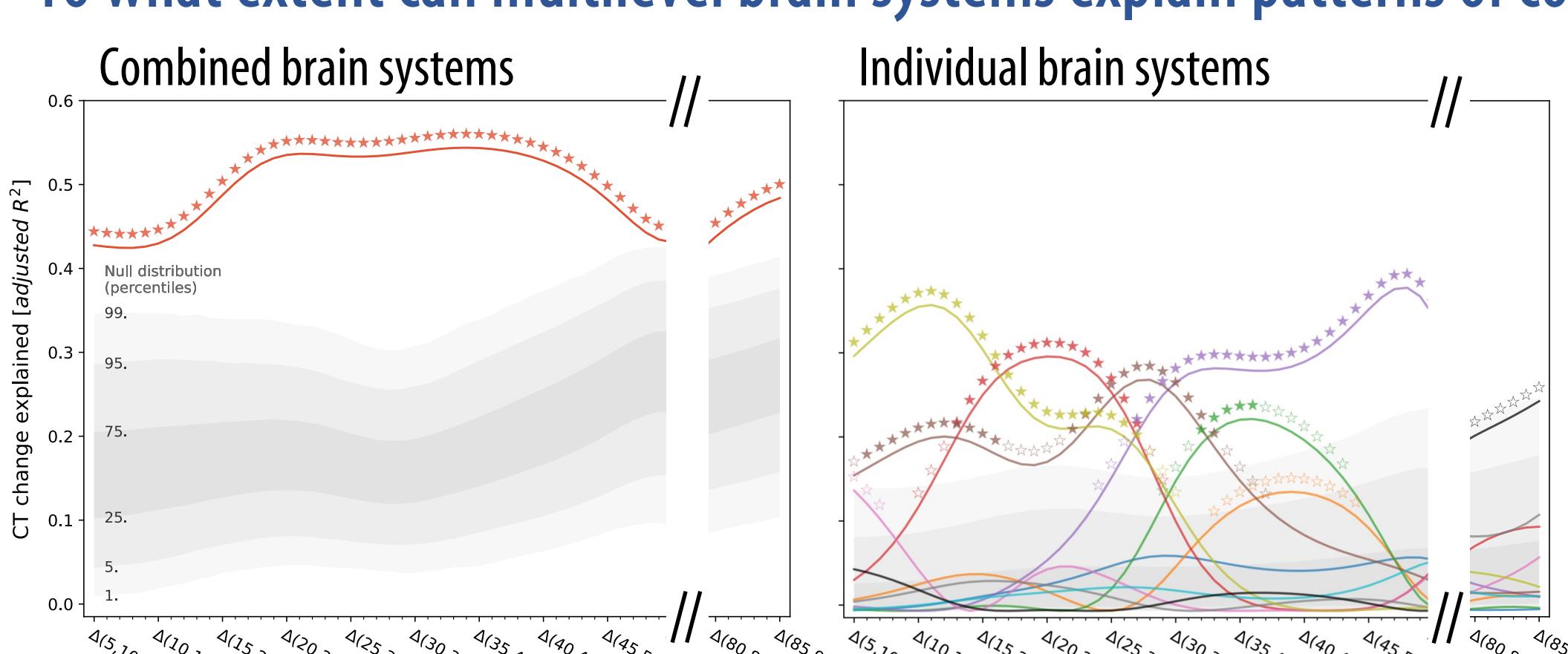
Observed CT change

## Analyses & results

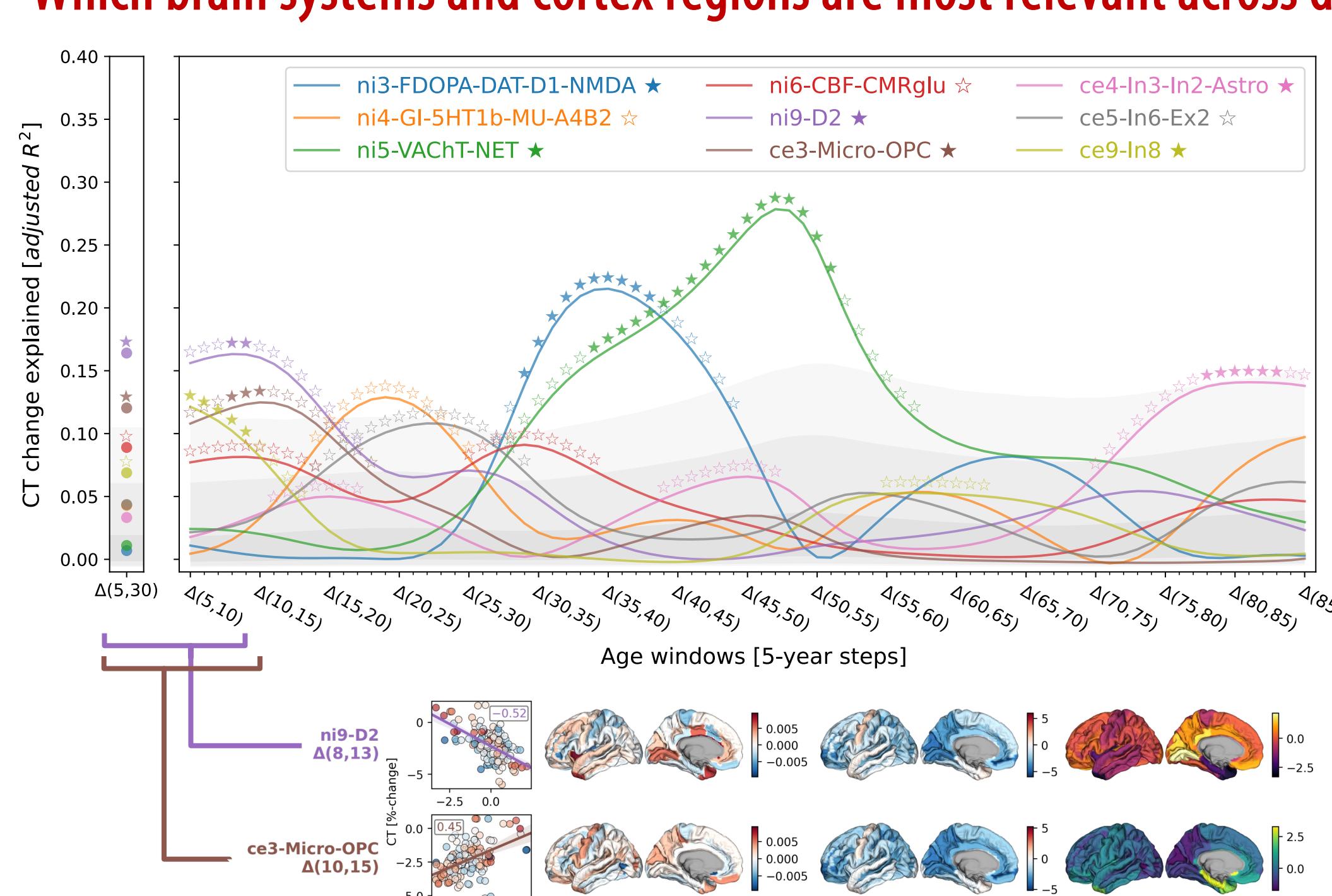
### How do spatial colocalization patterns develop across the lifespan?



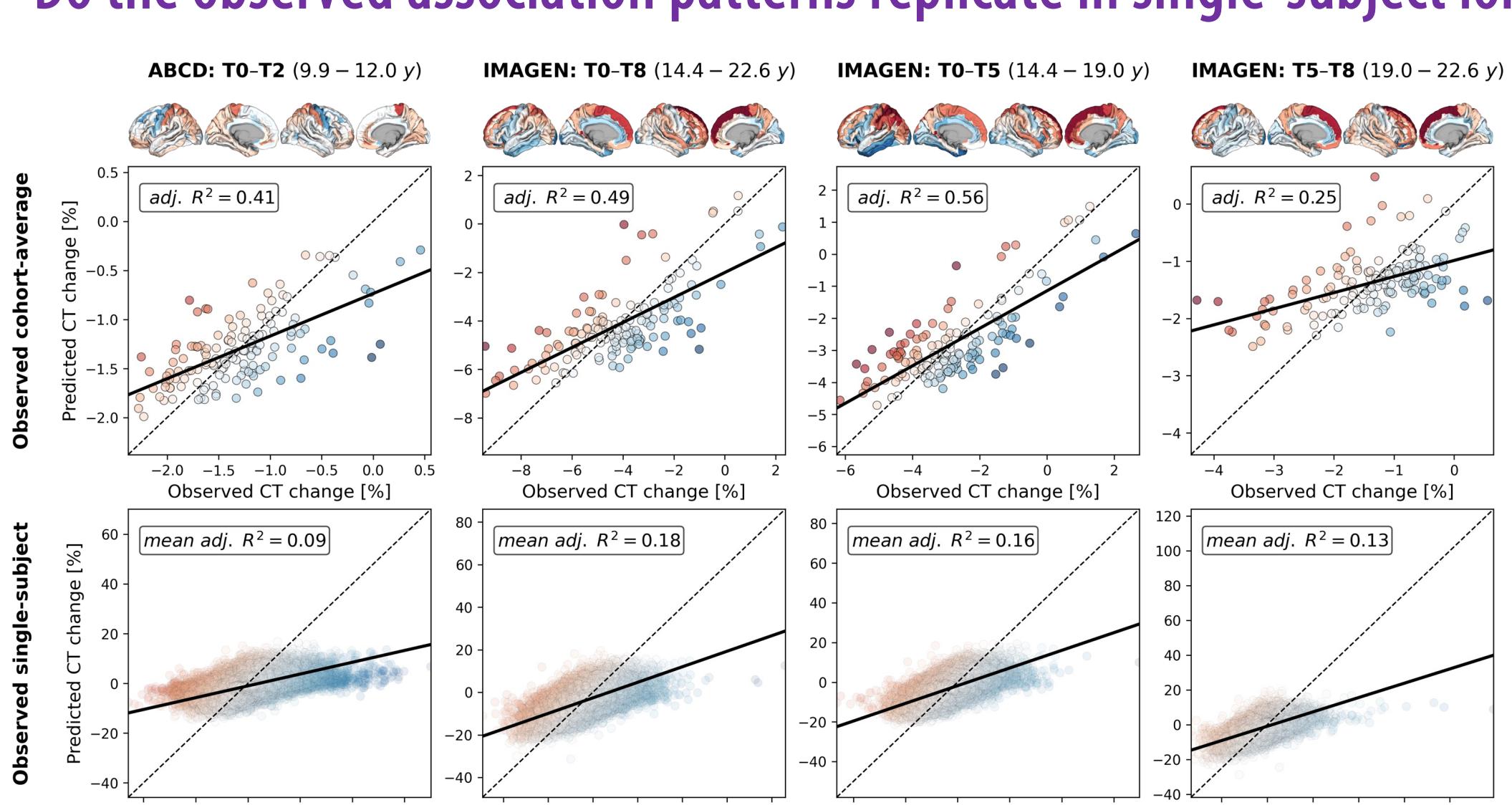
### To what extent can multilevel brain systems explain patterns of cortical development?



### Which brain systems and cortex regions are most relevant across developmental stages?



### Do the observed association patterns replicate in single-subject longitudinal data?



- Multilevel brain systems and modeled CT data show diverse lifespan colocalization trajectories.

- Molecular (Fig.) and cellular systems explain up to 55% of modeled lifespan CT change.
- Individual systems explain up to 40%.

- Only 9 systems can explain up to 57% of modeled CT change.
- D1/2 dopaminergic receptors, microglia, ST interneurons, and brain metabolism explain early CT development.
- Cholinergic and glutamatergic receptors explain later CT change.

- Modeled results replicate in independent longitudinal data.
- 6 biological systems explain up to 59% of cohort-average and 18% of single-subject CT development.

<sup>1</sup>Bethlehem, R. A. I. et al. (2022). Brain charts for the human lifespan. *Nature*, *604*(7906), Article 7906

<sup>2</sup>Rutherford, S. et al. (2022). Charting brain growth and aging at high spatial precision. *eLife*, *11*, e72904.

<sup>3</sup>Dukart, J. et al. (2021). JuSpace: A tool for spatial correlation analyses of magnetic resonance imaging data with nuclear imaging derived neurotransmitter maps. *Hum. Brain Mapp.*, *42*(3), 555–566.

<sup>4</sup>Vidal-Pineiro, D., et al. (2020). Cellular correlates of cortical thinning throughout the lifespan. *Sci. Rep.*, *10*(1), Article 1.

<sup>5</sup>Hansen, J.Y. et al. (2022). Mapping neurotransmitter systems to the structural and functional organization of the human neocortex. *Nat. Neurosci.*, *25*, 1569–1581.

<sup>6</sup>Hawrylycz, M.J. et al. (2012). An anatomically comprehensive atlas of the adult human brain transcriptome. *Nature*, *489*(7416), 391–399.

<sup>7</sup>Lake, B.B. et al. (2016). Neuronal subtypes and diversity revealed by single-nucleus RNA sequencing of the human brain. *Science*, *352*(6293), 1586–1590.

<sup>8</sup>Casey, B.J. et al. (2018). The Adolescent Brain Cognitive Development (ABCD) study: Imaging acquisition across 21 sites. *Dev. Cogn. Neuro.*, *32*, 43–54.

<sup>9</sup>Schumann, G. et al. (2010). The IMAGEN study: Reinforcement-related behaviour in normal brain function and psychopathology. *Mol. Psychiatry*, *15*(12), Article 12.

<sup>10</sup>Lotter, L.D. and Dukart, J. (2022). JuSpice - a toolbox for flexible assessment of spatial associations between brain maps. *Zenodo*.