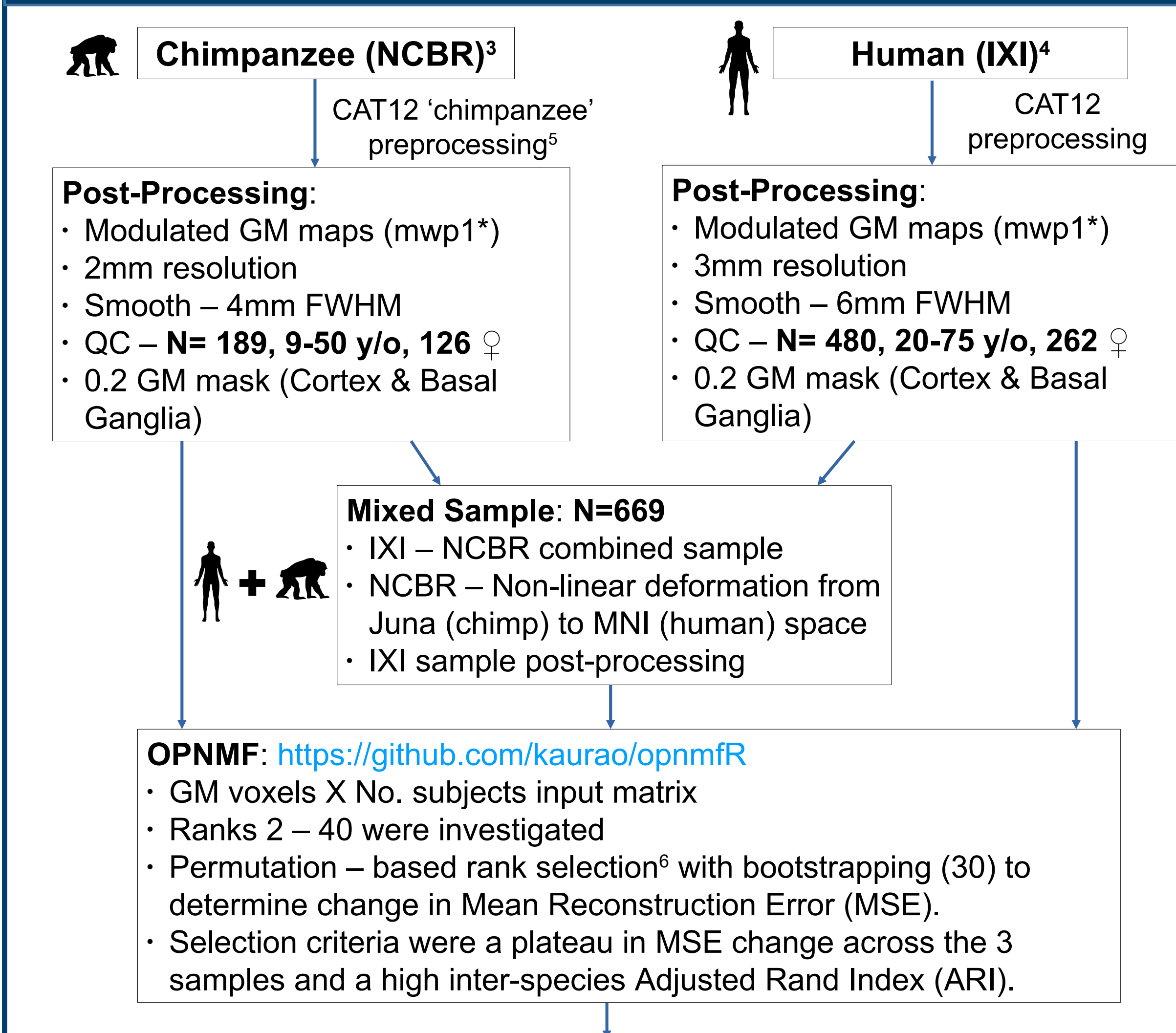


INTRODUCTION

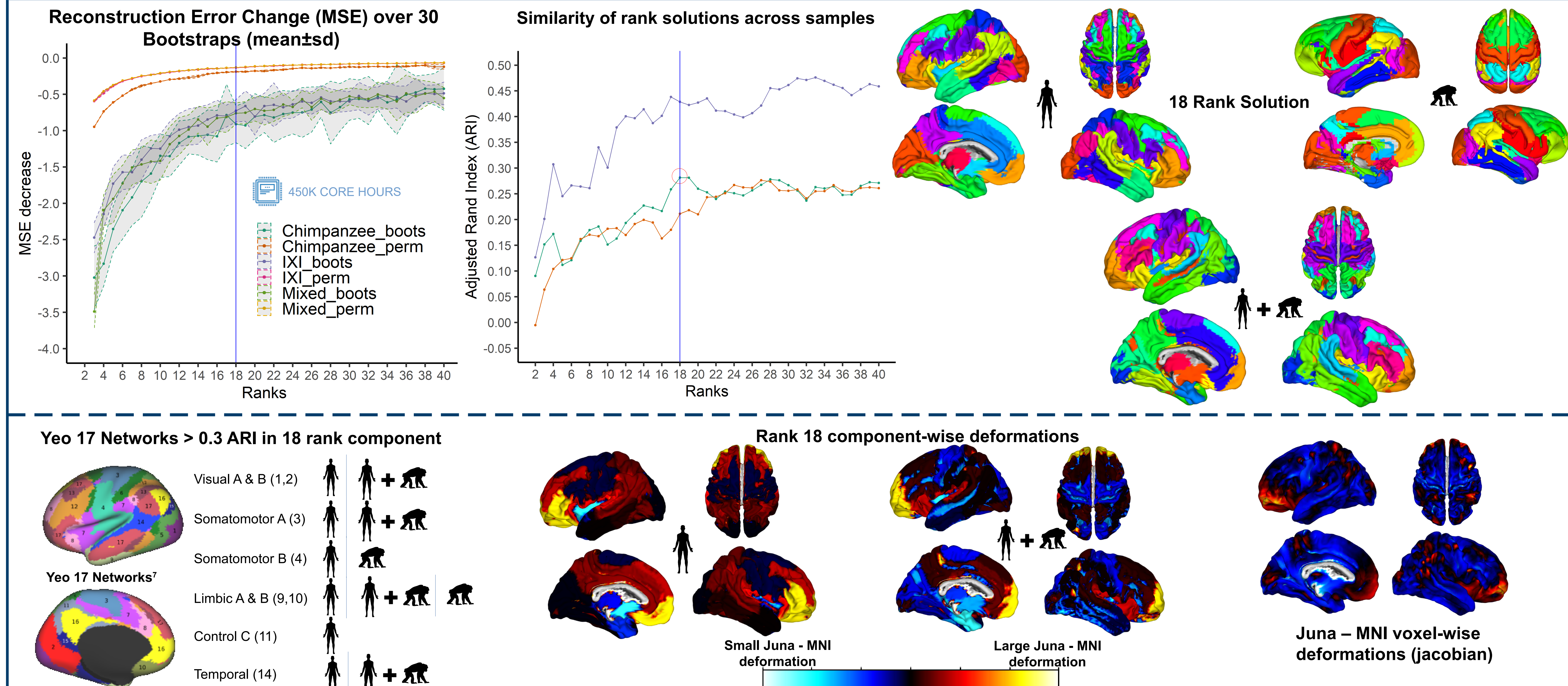
- Grey matter (GM) structural covariance (SC) uncovers regions that share connectivity, genetic similarities, and/or development¹
- Orthogonal Projective Non-Negative Matrix Factorisation (OPNMF)² creates data-driven SC regions with non-negative loadings improving interpretability and biological meaning to the components
- **Aim:** To present and compare OPNMF SC regions in humans and chimpanzees

METHODS



- Comparison:**
- Similarity of selected rank components were compared (ARI) to function connectivity networks (Yeo 17 networks)⁷.
 - Present mean deformation (Juna – MNI) of selected rank solution components.

RESULTS



DISCUSSION

- All 3 samples reveal a similar structure of reconstruction error change.
- Evolutionary deformations are captured within the 18 rank solution.
- The human rank solutions shows similarity to more higher-order networks as compared to the chimpanzee.
- The mixed sample contains SC features from both species and represents an extant greater ape (*homo* & *Pan*) parcellation.
- 18 rank solution contains similar cross-species regions that may represent conserved structural organisation in human evolution

- References**
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