

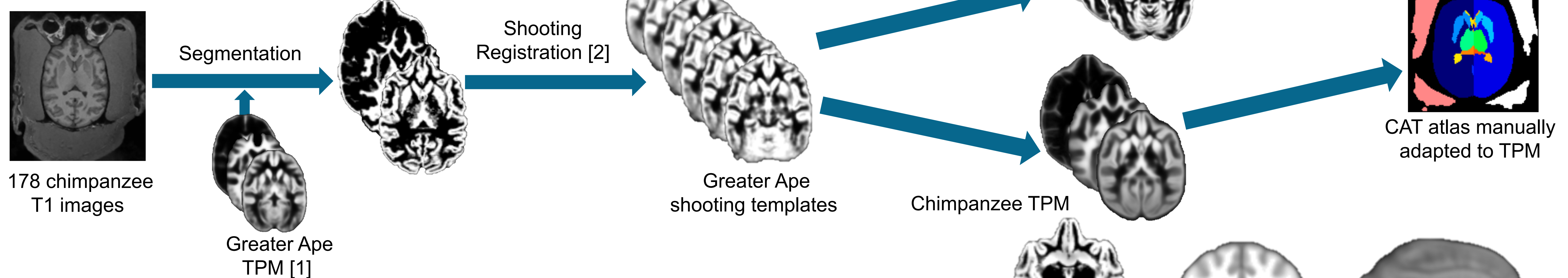
Introduction

- Chimpanzees are our closest extant primate relatives, therefore perfect for understanding the evolutionary development of the human brain
- But Human brains are significantly larger and with increased gyrification [1]
- Therefore, accurate image registration requires chimpanzee specific spatial normalization templates and tissue probability map (TPM).
- Chimpanzee specific templates and TPM are essential for reproducible group analysis
- A high contrast template makes macroanatomical annotations possible
- There is no publicly available chimpanzee atlas nor template so we provide an initial version of a chimpanzee template with macroscopic annotations and TPM for the Computation Anatomy Toolbox (CAT) (<http://dbm.neuro.uni-jena.de/cat>)

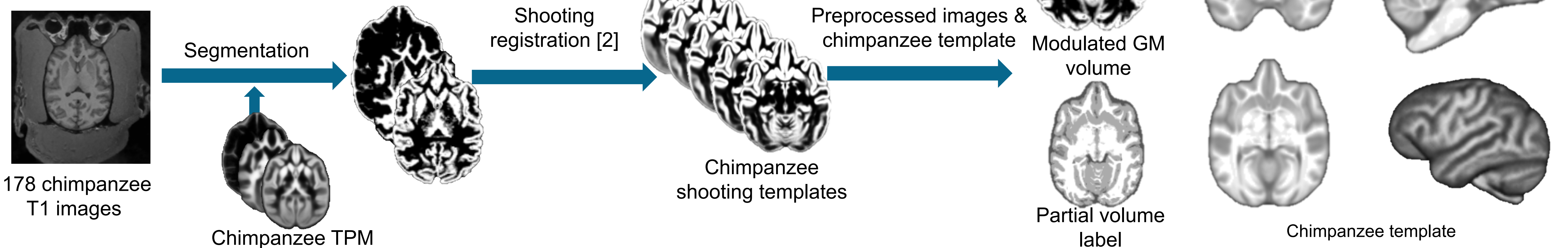
Methods

- 178 chimpanzees, 115 ♀, 63 ♂. (9 -54 y/o, 26.4±10.5) (<http://www.chimpanzeebrain.org>)
- Housed at Yerkes Primate Research Centre (YPRC) and the University of Texas M.D Cancer Centre (UTMDACC).

Step 1: Initial segmentation & registration

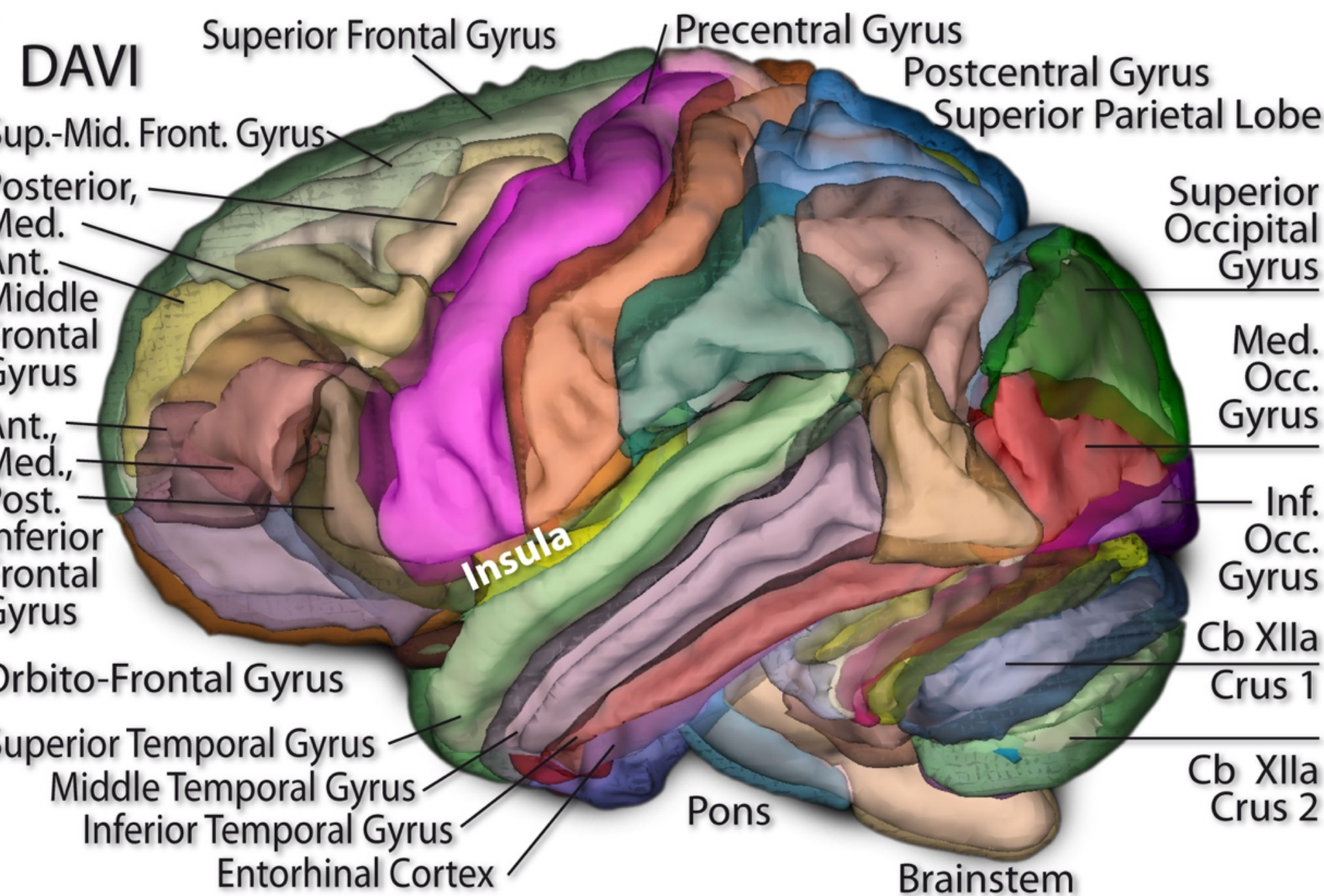
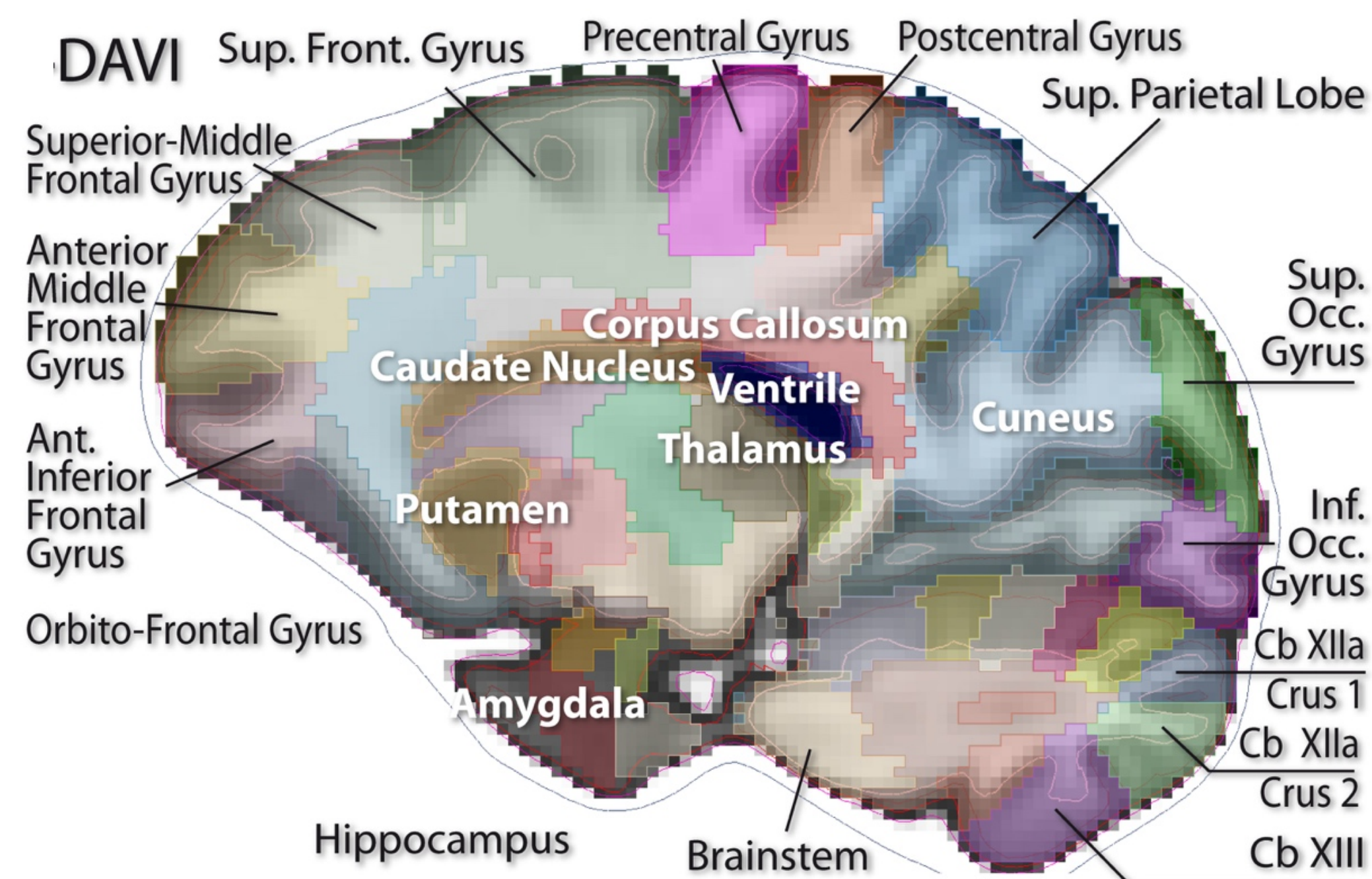


Step 2: Final segmentation & registration



Results

Davi Labelling



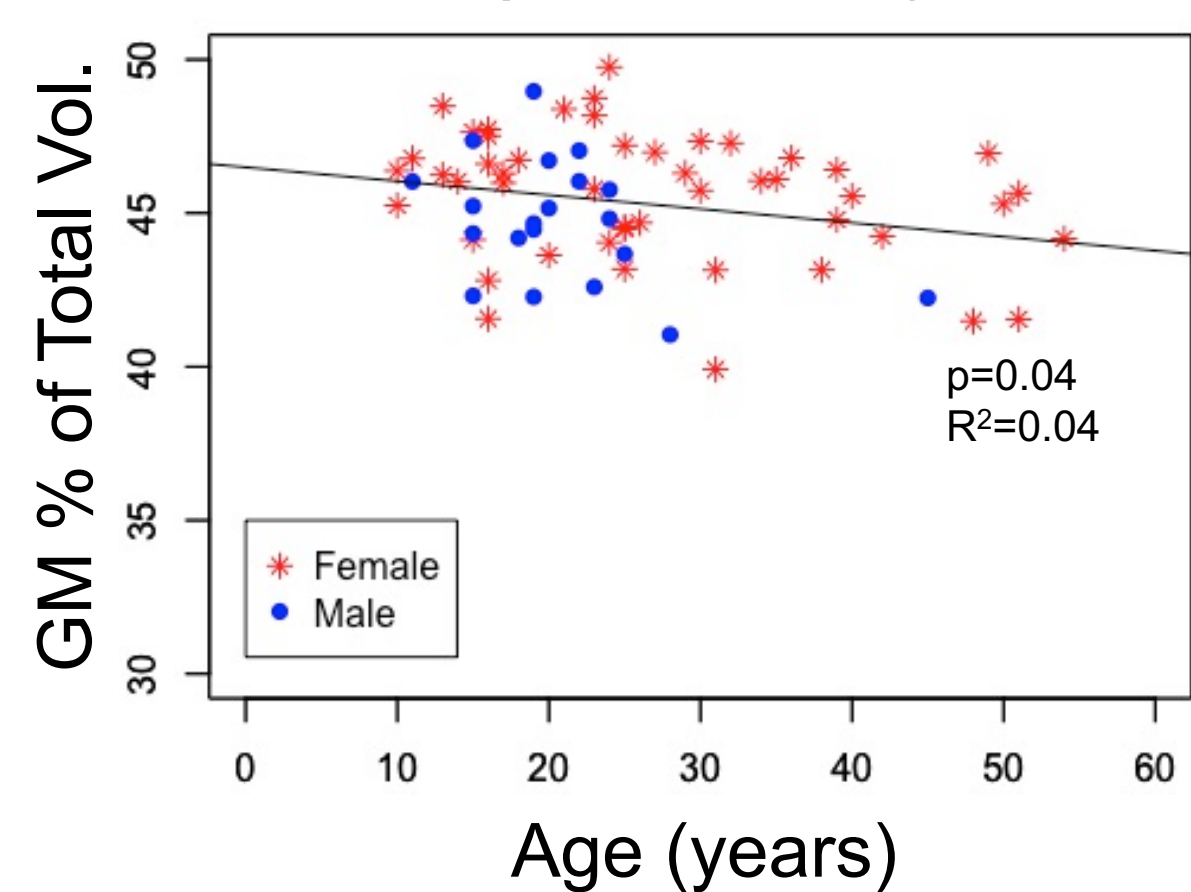
Davi Labelled ROI Age Regression

ROI	3T	1.5T	Total
Sup. Frontal G.	n.s.	☆☆☆	☆
Mid. Frontal G.	n.s.	☆☆	n.s.
Pars Orbitalis	☆	☆	☆☆
Pars Triangularis	n.s.	☆☆	☆
Pars Opercularis	n.s.	n.s.	n.s.
Orbitofrontal Cortex	☆☆	☆☆☆	☆☆☆
Med. Orbitofrontal G.	☆☆☆	☆☆	☆☆☆
Pre-central G.	n.s.	☆☆	n.s.
Post-central G.	n.s.	☆☆	n.s.
Paracentral Lobule	☆	n.s.	n.s.
Cingulate G.	n.s.	☆☆☆	☆☆
Sup. Temporal G.	☆	☆☆☆	☆☆☆
Mid. Temporal G.	n.s.	☆☆	☆
Inf. Temporal G.	☆☆	☆☆☆	☆☆☆
Entorhinal gyrus	n.s.	☆☆☆	n.s.
Insula	n.s.	☆☆	☆
Parahippocampal G.	n.s.	n.s.	☆☆
Fusiform G.	☆	☆☆	☆☆☆
Amygdala	n.s.	☆☆☆	☆☆
Hippocampus	n.s.	n.s.	n.s.
Sup. Parietal Lobule	n.s.	☆	n.s.
Supramarginal G.	n.s.	☆☆	☆☆
Angular G.	n.s.	☆	☆
Sup. Occipital G.	n.s.	☆☆	n.s.
Mid. Occipital G.	n.s.	n.s.	n.s.
Inf. Occipital G.	n.s.	n.s.	n.s.
Precuneus	n.s.	☆☆	☆☆
Cuneus	n.s.	n.s.	n.s.
Lingual G.	☆☆☆	☆☆	☆☆
Caudate Nucleus	☆☆☆	☆☆	☆☆☆
Putamen	☆	☆☆	☆☆☆
Globus pallidus	☆☆☆	n.s.	☆☆
Thalamus	n.s.	☆☆	n.s.
Hypothalamus	n.s.	n.s.	n.s.

☆- p ≤ 0.05, ☆☆☆- p ≤ 0.001.
☆☆- p ≤ 0.01, n.s. = not significant.

Age Regression

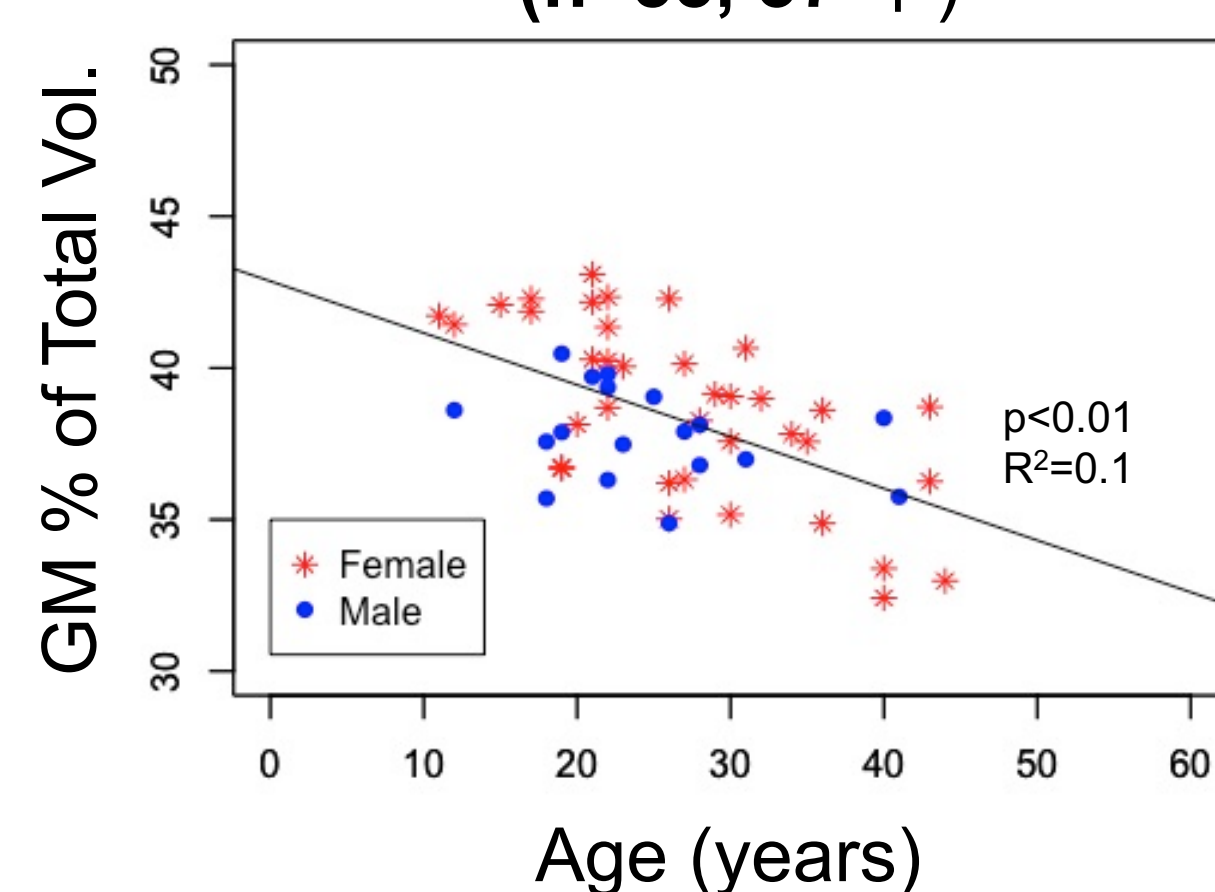
3T GM atrophy in aging (n=70, 49 ♀)



VBM age analysis, 6mm smoothing, TFCE corrected (FWE<0.05)



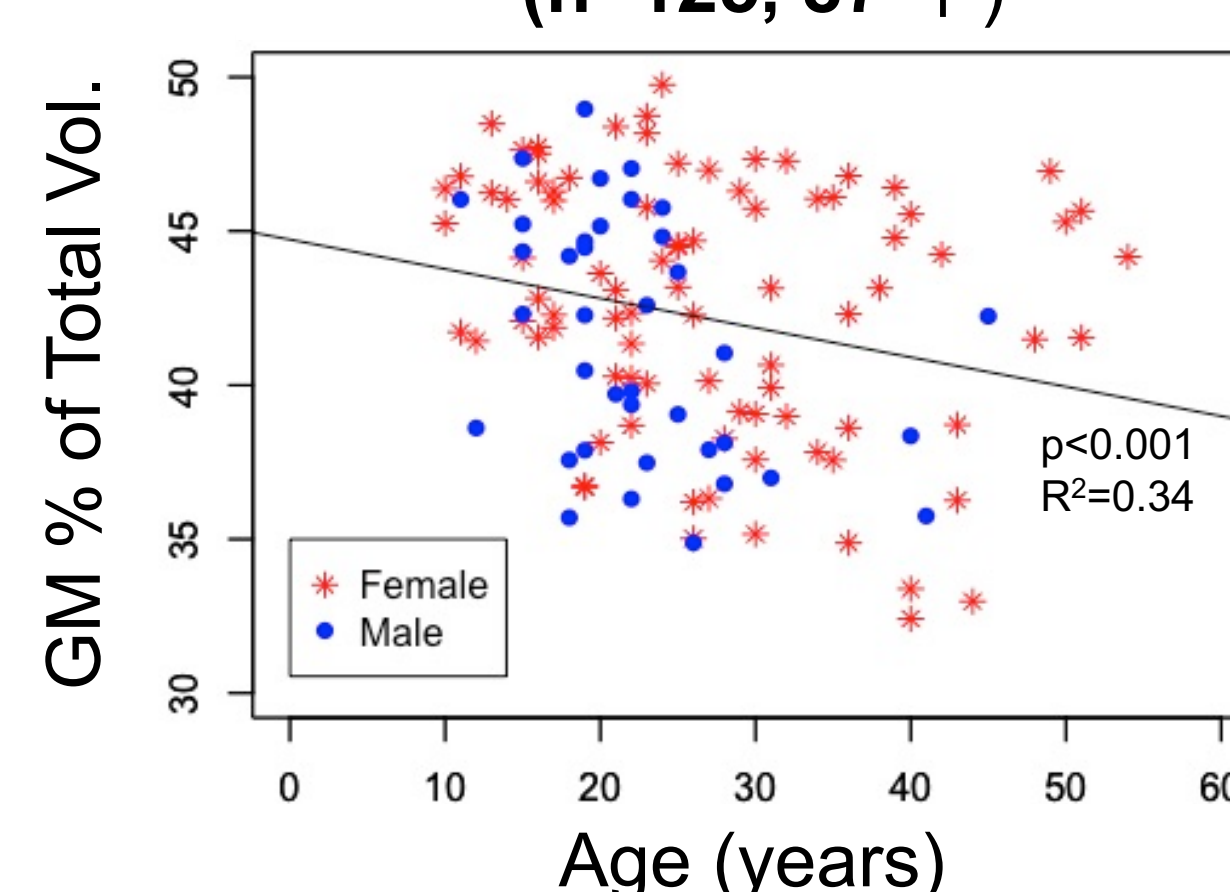
1.5T GM atrophy in aging (n=55, 37 ♀)



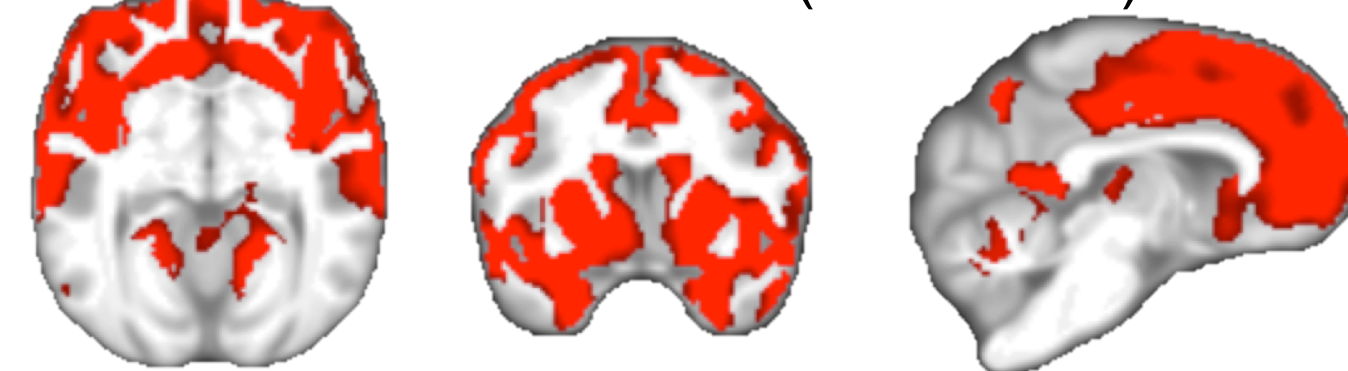
VBM age analysis, 6mm smoothing, TFCE corrected (FWE<0.05)



All GM atrophy in aging (n=125, 87 ♀)



VBM age analysis, 6mm smoothing, TFCE corrected (FWE<0.05)



Discussion

- Here we present a chimpanzee specific preprocessing pipeline with required templates for the CAT toolbox that enables automated processing and analysis.
- We validated our template by showing similar age effects as [3,4] and VBM illustrated a widespread GM volume decrease with age, similar to that seen in humans [5]
- We understand that macroscopic parcellation is limited in general [6]. However, our ROI and VBM analysis showed that this pipeline and macroscopic labelling can help to improve our understanding of primate brain development and aging [7].