



Supervoxel parcellation of visual cortex connectivity

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Summary

- ▶ How is the cortex functionally and structurally organized?
- ▶ New tool: High-resolution connectivity matrices
 - ▷ Functional measure: Resting-state connectivity
 - ▷ Anatomical measure: Diffusion tractography
- ▶ How similar are functional and anatomical connectivity?
 - ▷ **Result:** Similar at voxel-scale, but depends on cortical location
- ▶ What is the spatial structure of these connectivity matrices?
 - ▷ Explore using spatially-informed clustering
 - ▷ **Result:** Reveals retinotopic and functional organization

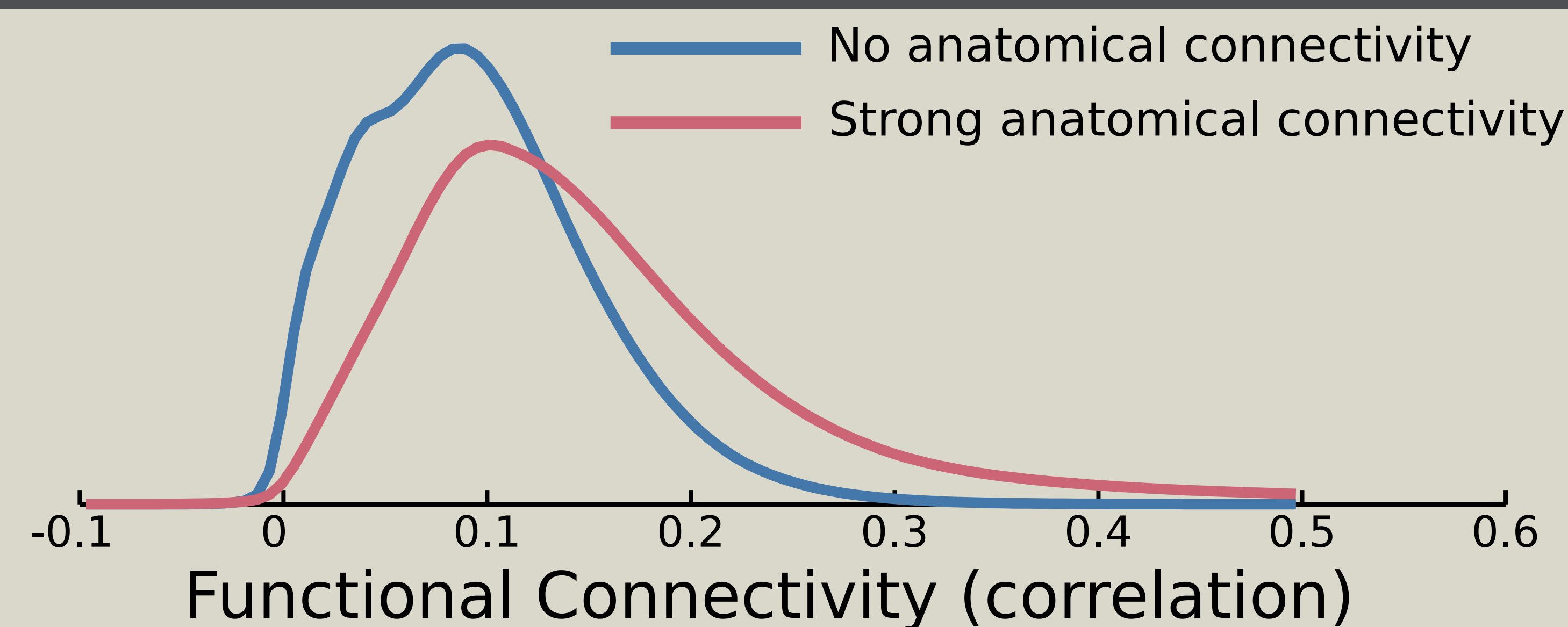
Previous Work

- ▶ Whole-brain comparisons of functional and anatomical connectivity are coarse, atlas-dependent^{1;2;3;4}
- ▶ Greedy clustering algorithms give only approximate solutions^{5;6;7}

Data: Human Connectome Project

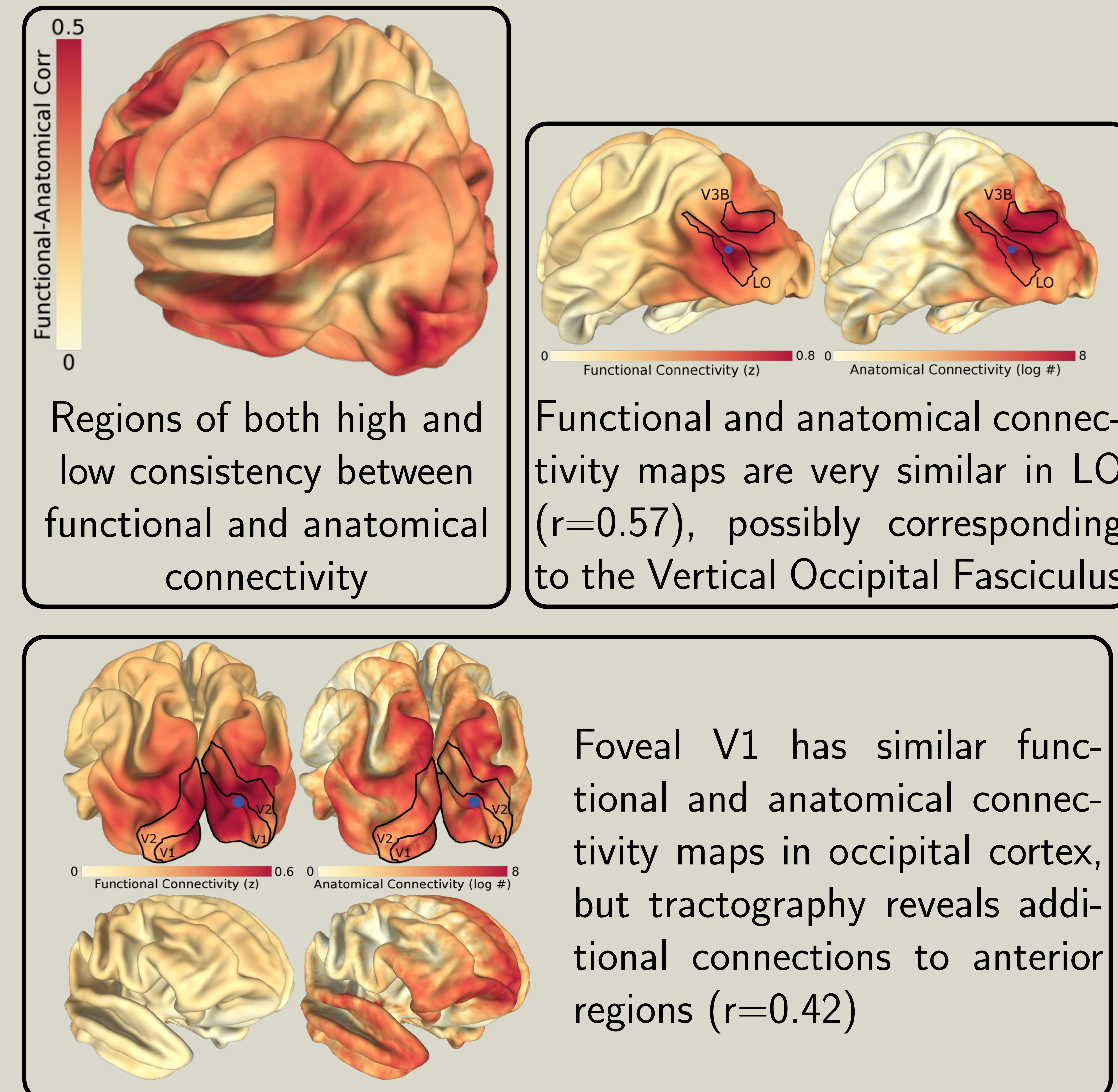
- ▶ Resting-state fMRI - 40 subjects (2mm isotropic)
 - ▷ Connectivity = correlation between timecourses
- ▶ Diffusion Tractography - 10 subjects (1.25mm isotropic)
 - ▷ Sampled 33 billion tracts using FSL
 - ▷ Connectivity = log number of fibers between voxels

Voxel-level Multimodal Comparison



Anatomical connectivity is consistently predictive of functional connectivity for individual voxel pairs ($r=0.25$)

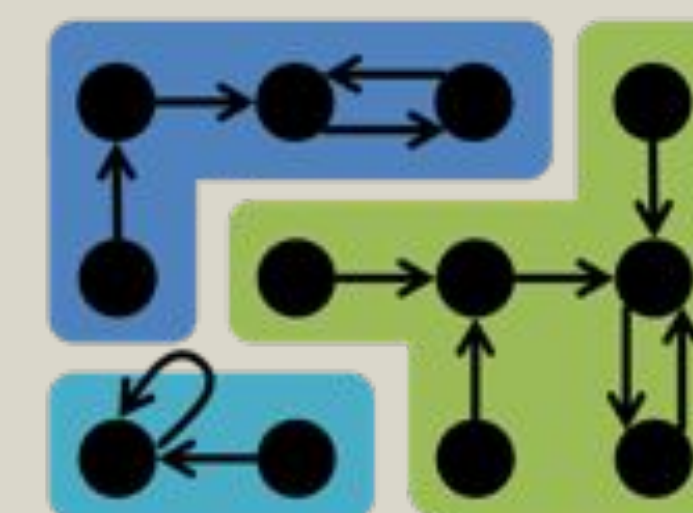
Multimodal Comparisons Across Cortex



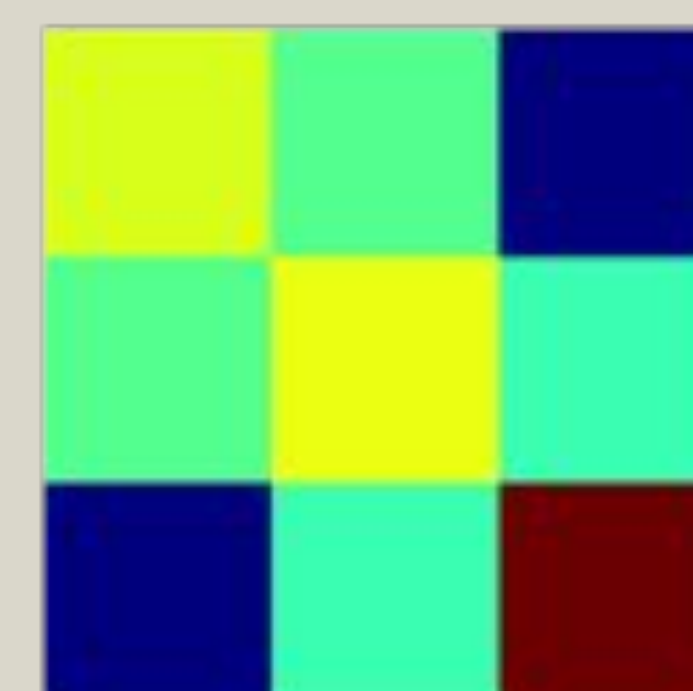
Generative Clustering Model

- ▶ Produces spatially-contiguous "supervoxels"
- ▶ Refines clustering with multiple passes
- ▶ Uses data statistics to help set number of clusters

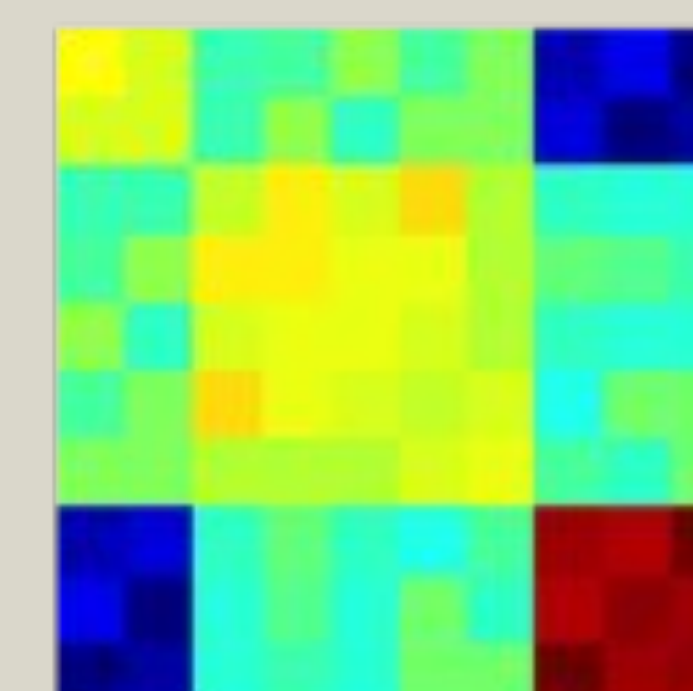
① Each voxel selects a neighbor to cluster with



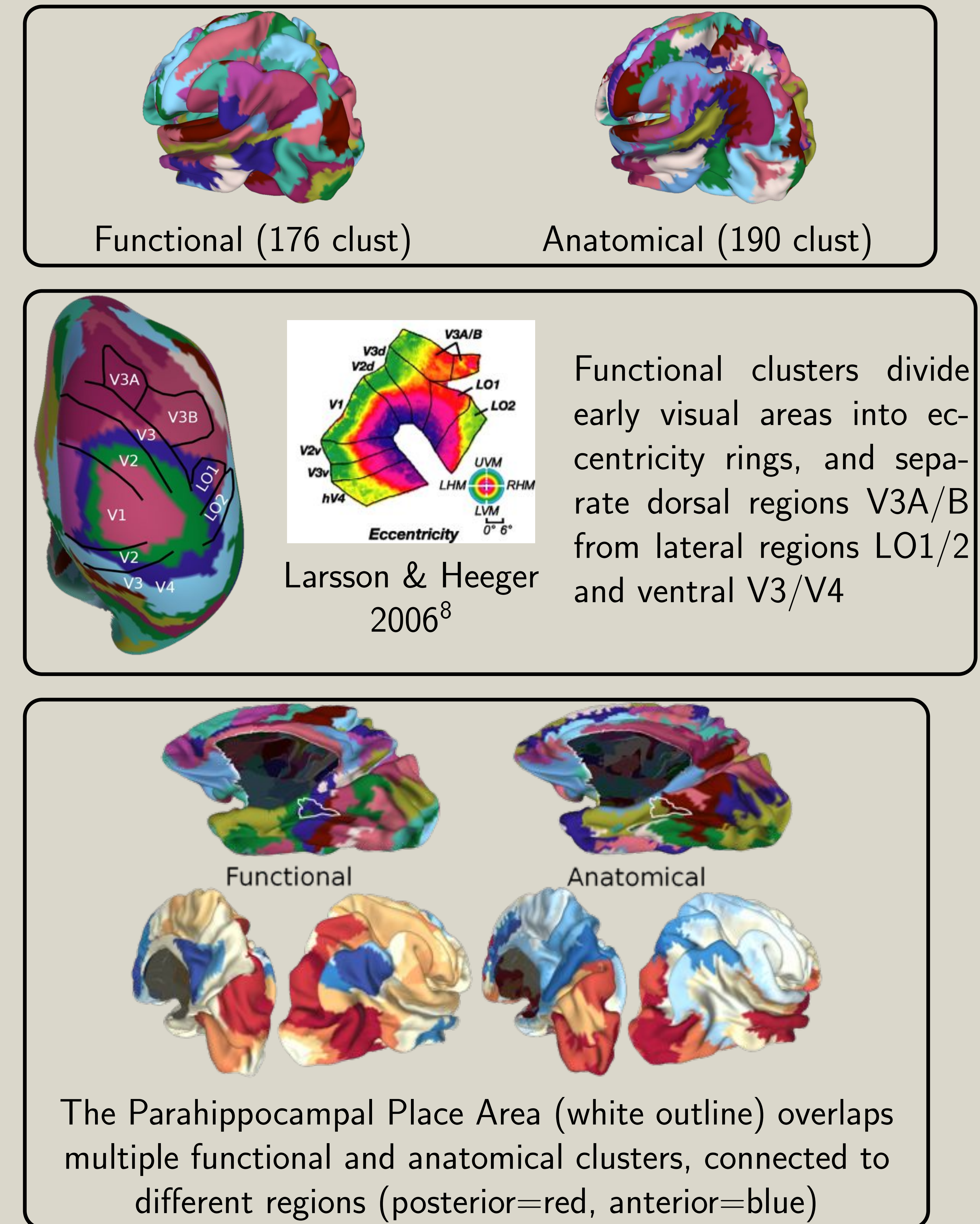
② Latent connectivity between supervoxels



③ Observed connectivity is noisy estimate of supervoxel connectivity



Supervoxel Clustering



Funding and References

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