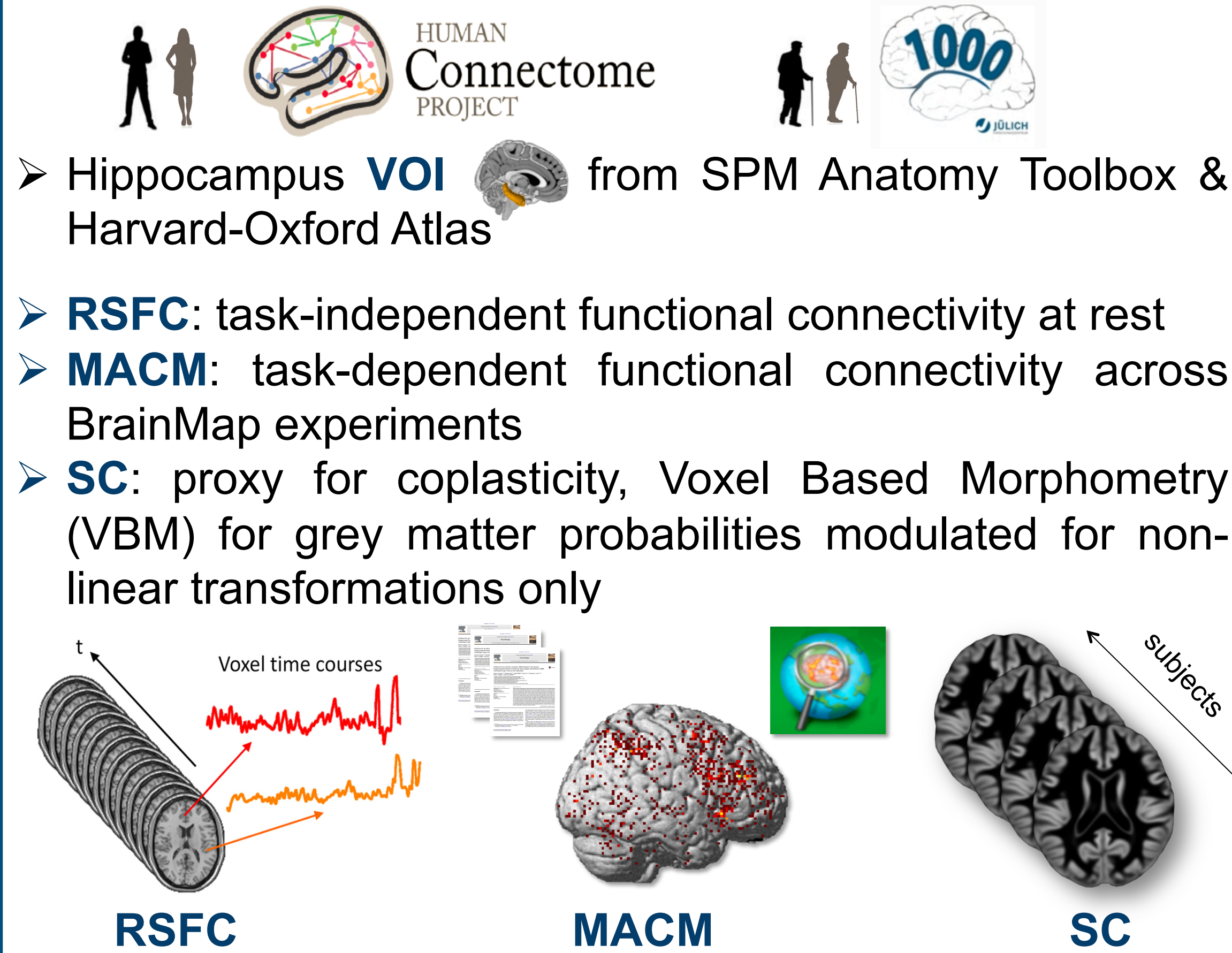


Introduction

- **Connectivity-Based Parcellation (CBP)** reveals brain organization [1]
- **Different markers of functional connectivity** can be used for CBP: task-(in)dependent functional connectivity (RSFC, MACM) and structural (SC)
- **RSFC-CBP** most widely used to examine brain organization but **denoising** and its effects on CBP has not yet been investigated
- Recent methods developments focused on cortical partitions, hence neglecting crucial subcortical structures [2,3]
- We examined **stability and reproducibility of RSFC-CBP** as a function of different denoising approaches in the hippocampus

Methods

- RSFC-CBP and SC-CBP performed on functional and structural MRI data of **two cohorts**:
HCP (n = 323) 1000BRAINS (n = 671)



RSFC denoising approaches:

- 24 Motion regressors (no denoising)
- Global Signal Regression (GSR)
- White Matter and CSF signal regression (WM/CSF)
- FMRIB's ICA-based X-noisifier (FIX)
- Combination of FIX and GSR (FIX+GSR)
- FIX and WM/CSF regression (FIX+WM/CSF)

Clustering:

- k-means for k=2-7, 500 repetitions, 255 iterations

Reliability assessment:

- Internal stability: 10 000 split-half resampling
- Reproducibility: across cohorts and modalities with 10 000 bootstrap resampling
- Consistency measure: adjusted Rand Index (aRI)

Statistics:

- Analysis of Variance (ANOVA) performed on split-half-, bootstrap samples, RSFC voxels' time-course similarity, voxels' connectivity profile dissimilarity

Results

- High stability with **GSR, WM/CSF and FIX+WM/CSF** (.79-.84 aRI; $p < .0001$)
- **Highest reproducibility** across samples (.52 aRI) and modalities ($< .45$ aRI) likewise with **FIX+WM/CSF** ($p < .0001$, Fig. 1A)
- FIX+GSR and FIX+WM/CSF significantly **reduced** the similarity between the **time-courses of the seed voxels** ($p < .0001$), and additionally **enhanced the dissimilarity of their connectivity profiles** ($p < .0001$, Fig. 1B,C)

RSFC-CBP's stability and reproducibility dependent on denoising

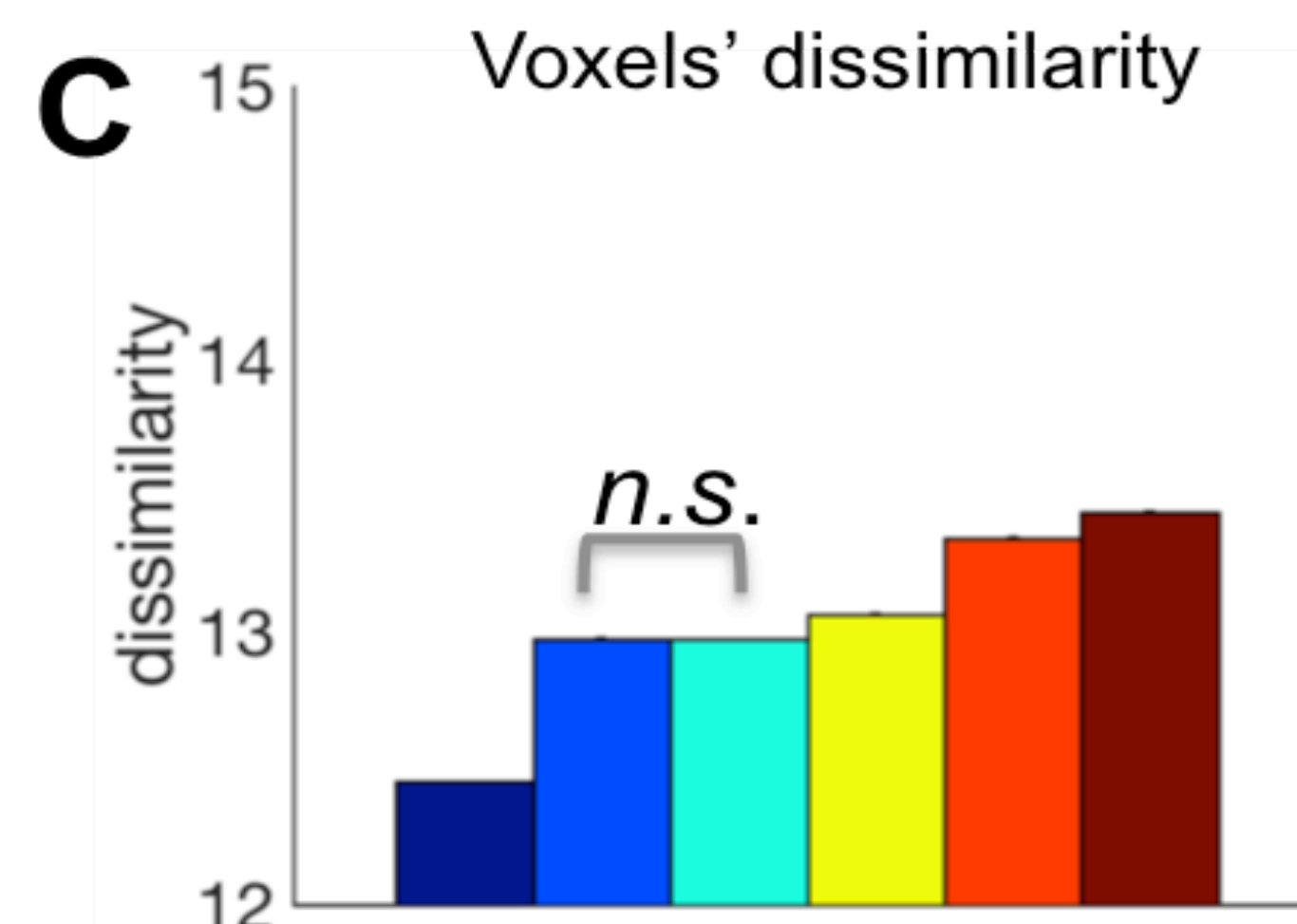
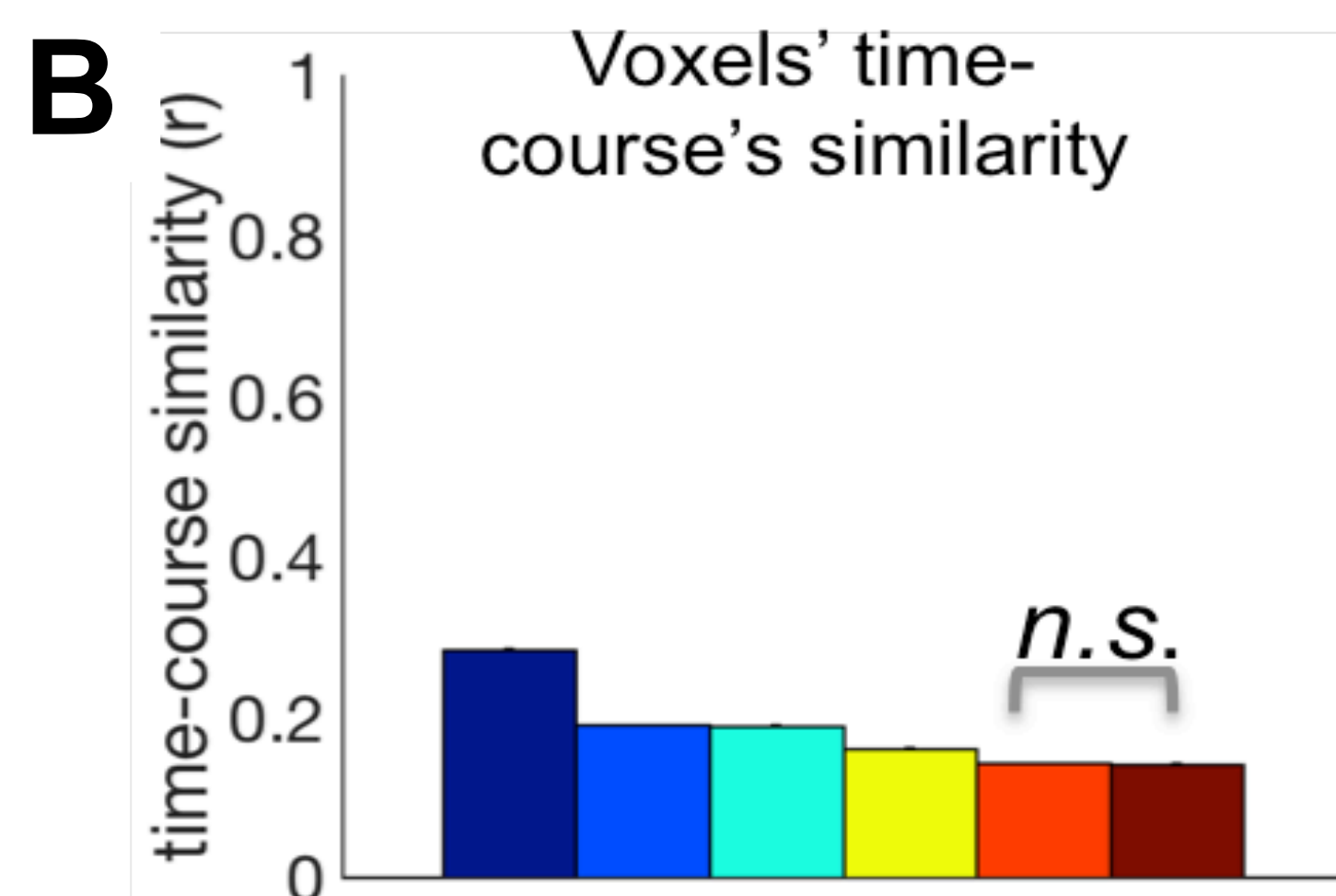
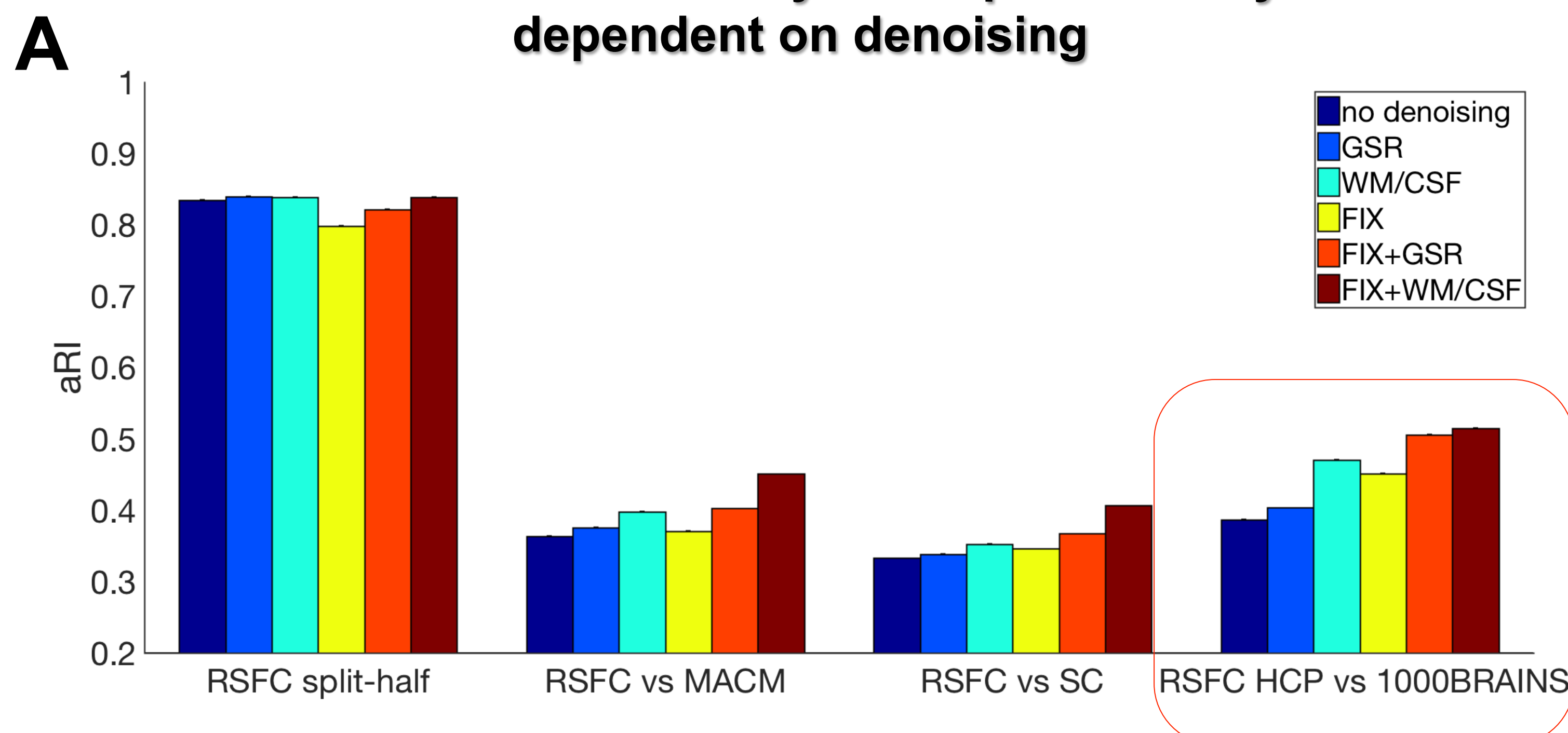
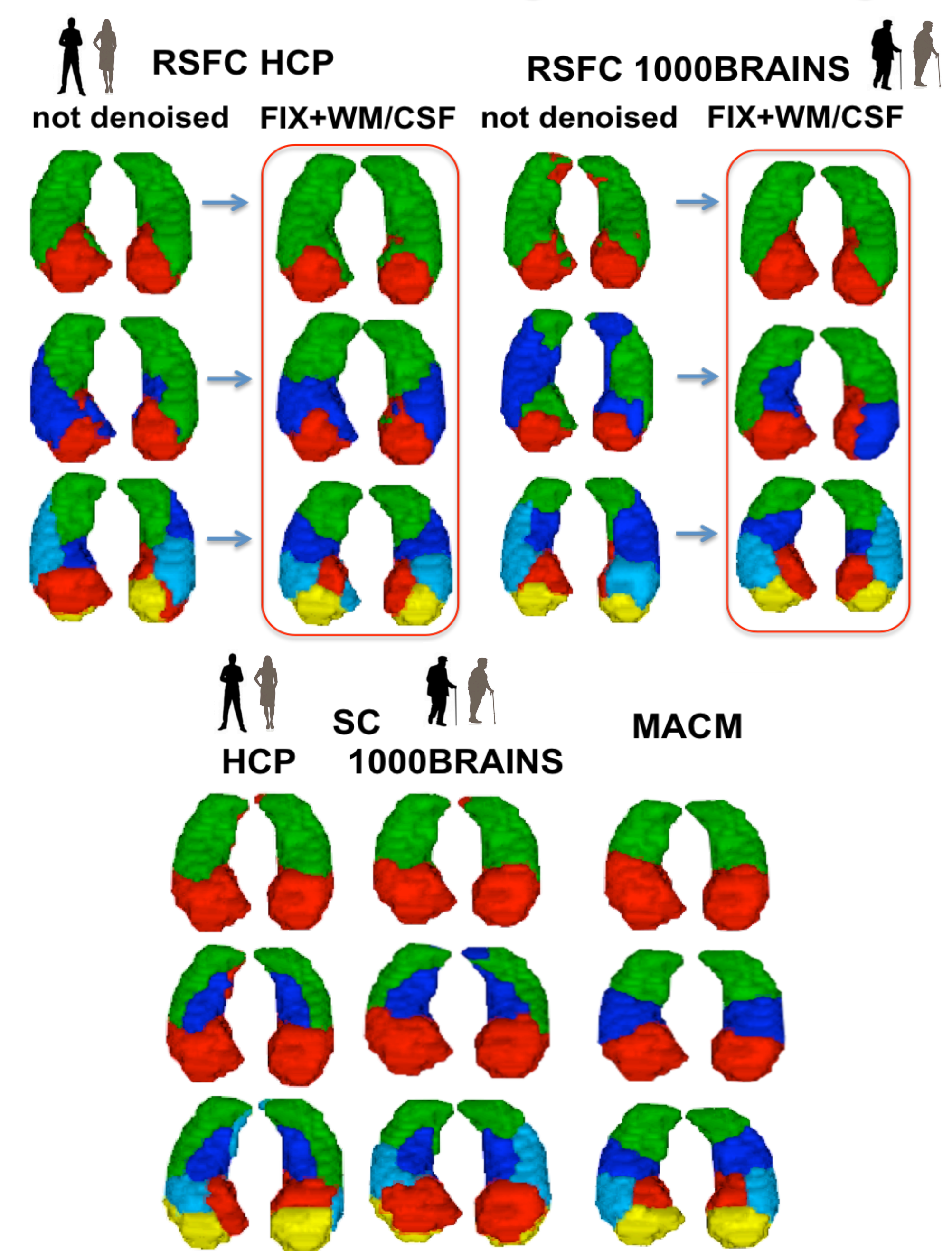


Figure 1. Note: bars represent mean and +/- standard error.

- With FIX-WM/CSF 2-cluster solution demonstrated highest convergence across samples and between RSFC and SC ($p < .0001$)
- **Highest consistency between RSFC and MACM for 5-cluster solution** ($p < .0001$) featuring a differentiation into tail, body and a three-way subdivision of the hippocampus head into a ventral, dorsolateral, and dorsomedial part (Fig. 2)

Effect of denoising on clustering



RSFC-CBP's convergence dependent on granularity

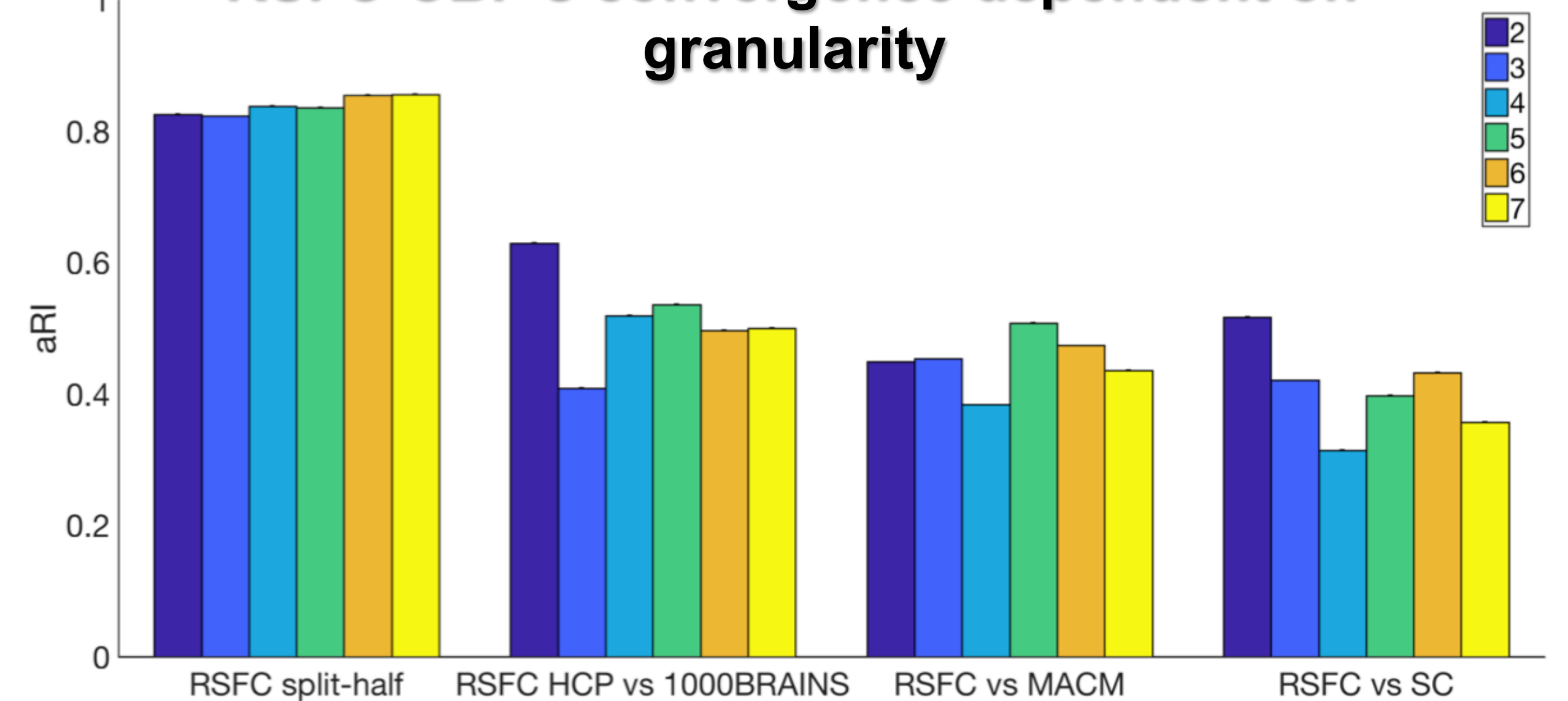


Figure 2. Note: bars represent mean and +/- standard error.

Discussion

- **Model-based and model-free denoising strategy** in combination such as FIX +WM/CSF leads to **stable clustering of the hippocampus**
- This approach successfully **eliminated structured noise** rendering the seed time-series' less inter-correlated and **enhanced the distinctiveness** of the connectivity profiles
- **FIX+WM/CSF** also yielded **most reproducible** clusterings across samples and modalities
- **Conclusion: higher biological validity of RSFC-CBP** after FIX +WM/CSF denoising, revealing a hippocampal organization confirmed by tracing, lesion and genetic studies across species [4,5,6].