



removal of perfusion lag structure before ICA-FIX cleaning boosts reproducibility in fcMRI



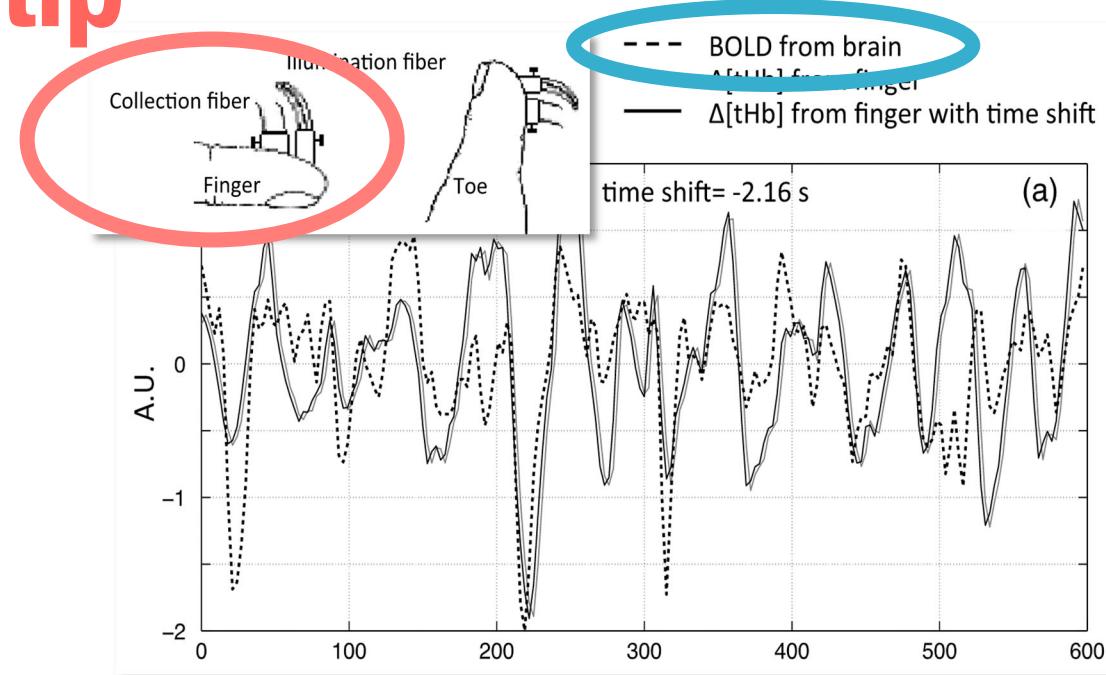
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Brain Connectomics Imaging Lab.
RIKEN Center for Biosystems Dynamics Research,
Kobe, Japan

Bad news: fcMRI signal contains “slow noise”

NIRS signal from
the fingertip

sLFO
Spontaneous
low-frequency
oscillation

≈ fMRI global mean signal



Systemic circulation

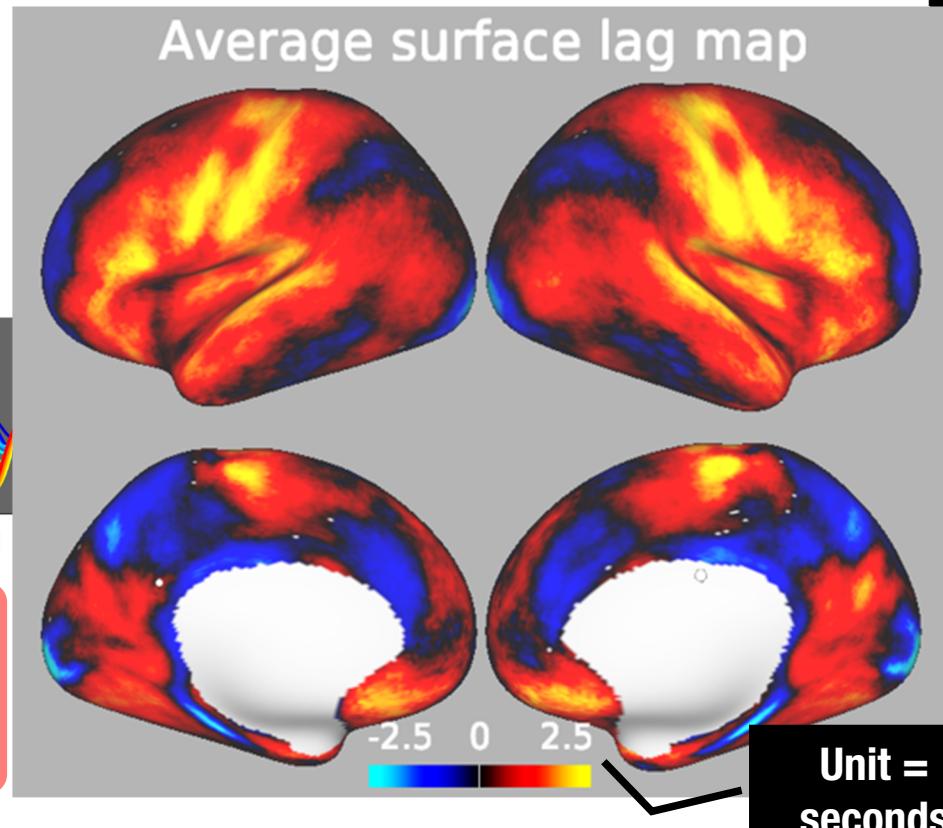
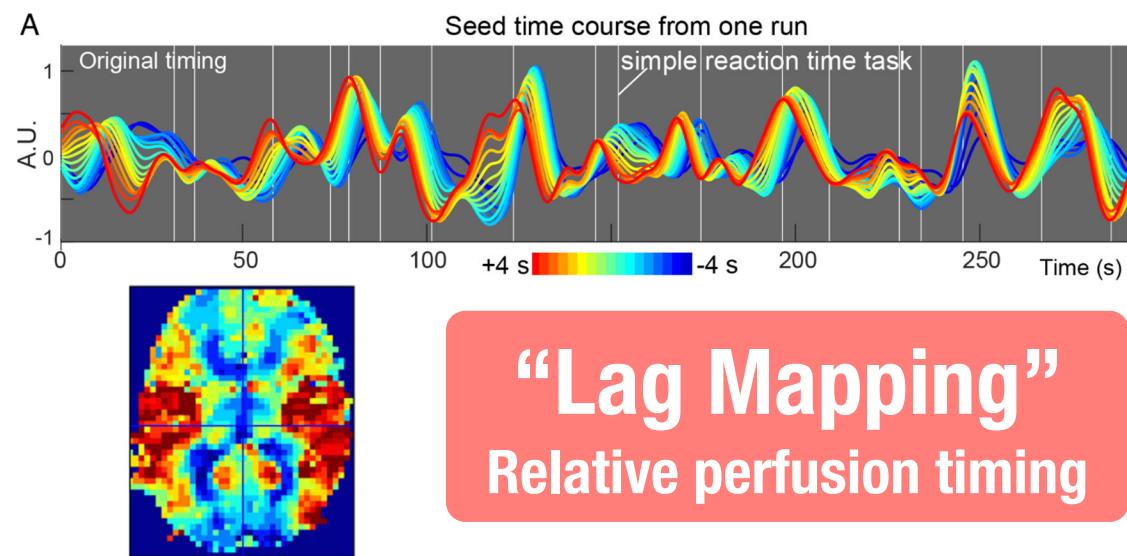
↓
Not from local
neuro-vascular
coupling

Good News: It is detectable!

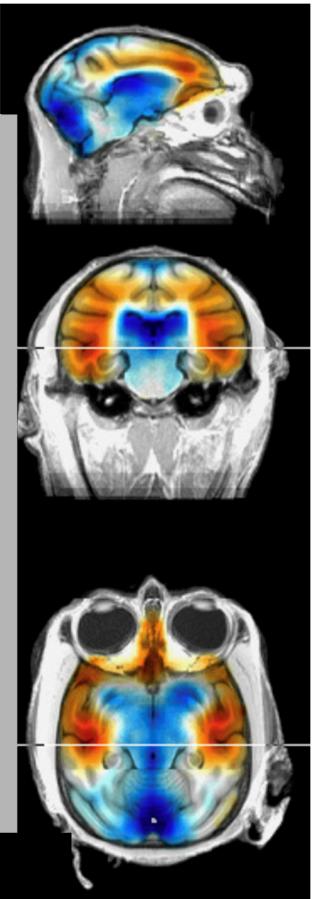
Phase delay (lag)
observed between
fingertip and brain
→ How about
within the brain?



Tong & Frederick
(2010) *NeuroImage*

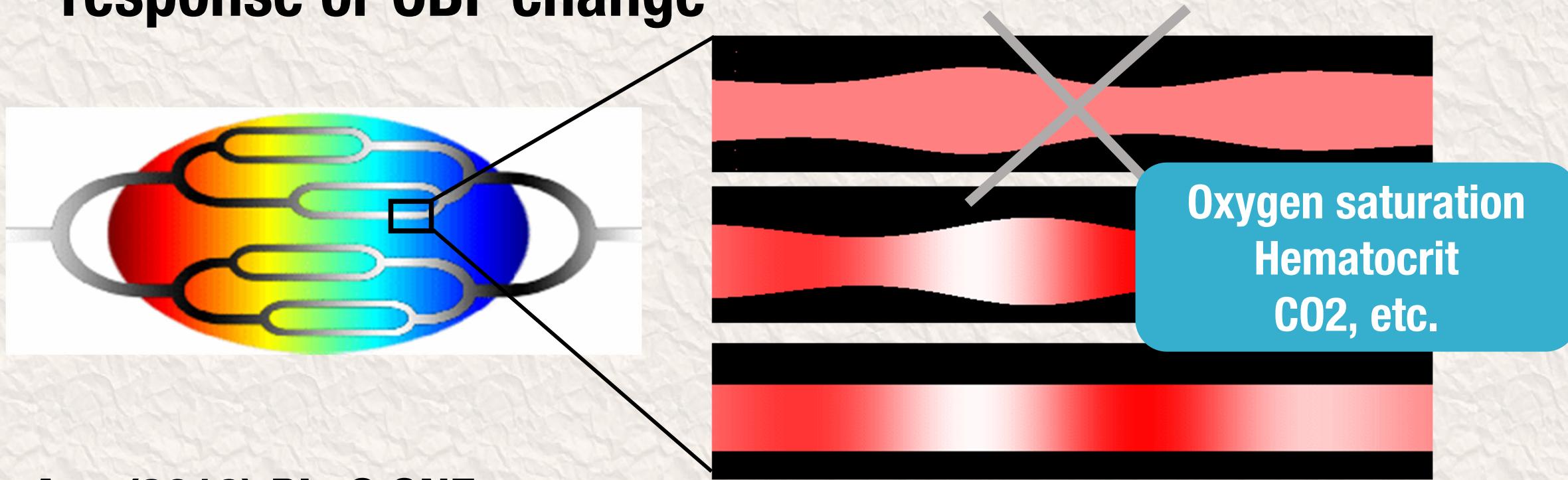


Aso, Hayashi
HBM2020



sLFO = variation of deoxy-Hgb density

- Working hypothesis: sLFO is independent of vascular response or CBF change



Aso (2019) PLoS ONE

OK, so what's the problem?

- The lag structure has its own correlation structure



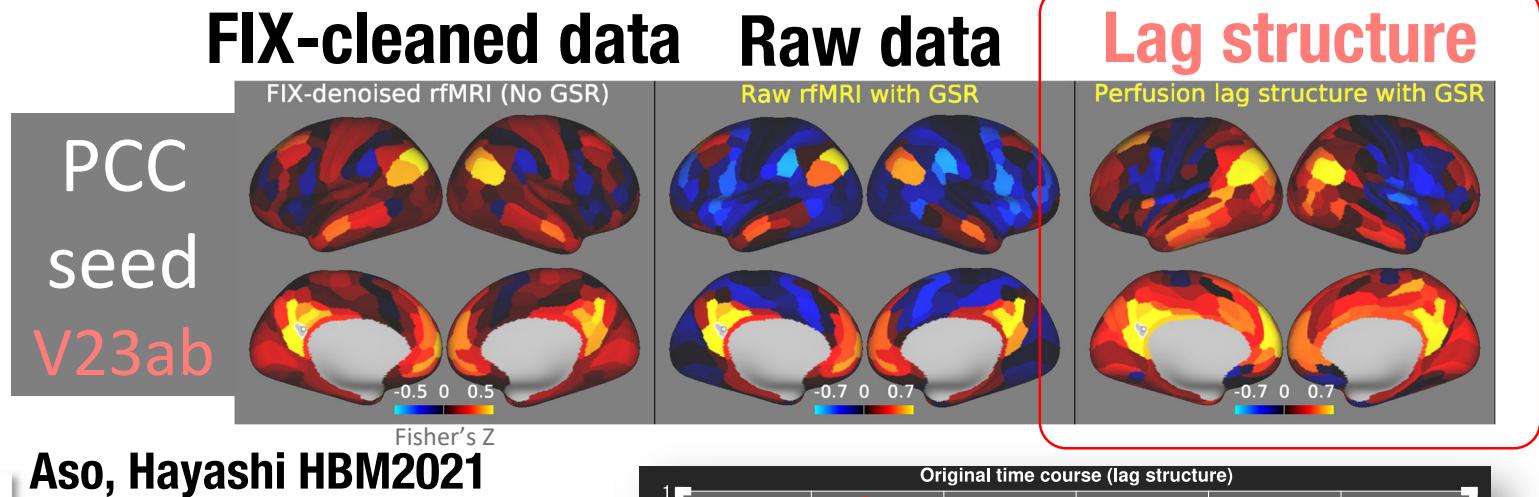
Chen (2020) NeuroImage; Tong (2015) FNS
Resting-state “physiological networks”

Jingyuan E. Chen ^{a,b,*}, Laura D. Lewis ^c, Catie Chang ^d, Qiyuan Tian ^{a,b}, Nina E. Fultz ^a,
Ned A. Ohringer ^a, Bruce R. Rosen ^{a,b,e}, Jonathan R. Polimeni ^{a,b,e}

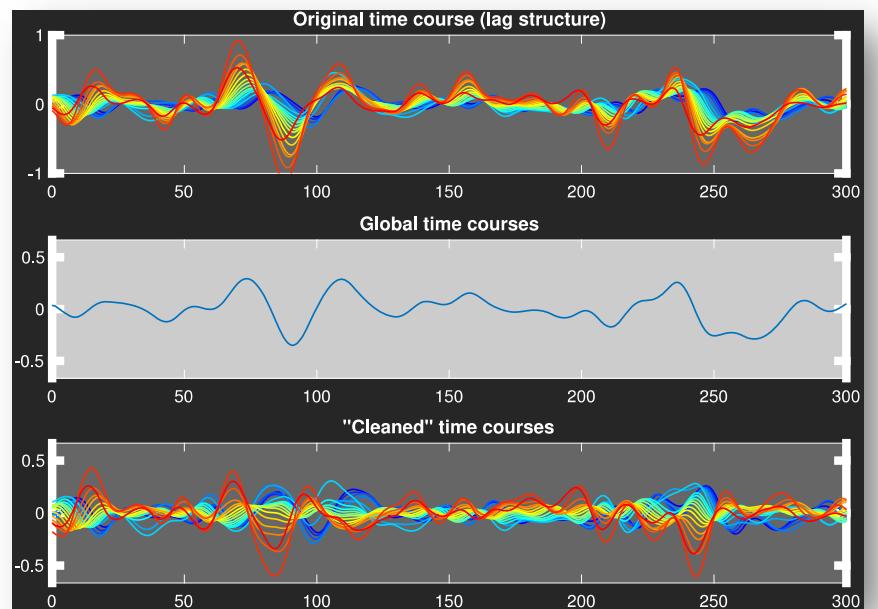
- GSR doesn't do!!

- The lag structure is a major part of global signal, but GSR leaves the correlation structure
- Tailored regression of sLFO for each voxel is needed

Erdogán, 2016 “Dynamic GSR”, Aso 2019



Aso, Hayashi HBM2021



• The lag structure has its own “connectivity”, or correlations

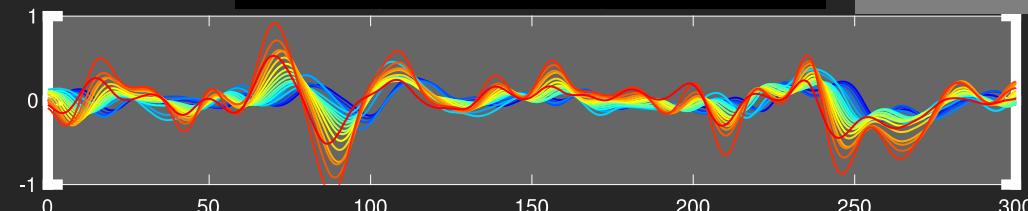
- Tong (2015) FNS, Chen (2020) NeuroImage



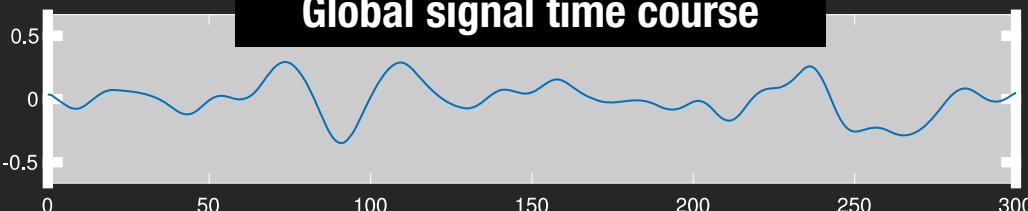
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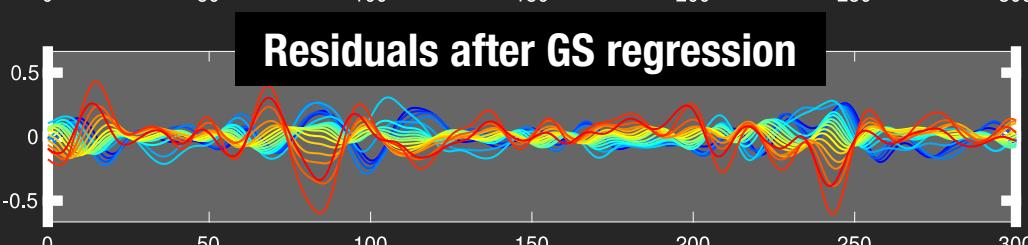
Lag structure time courses



Global signal time course

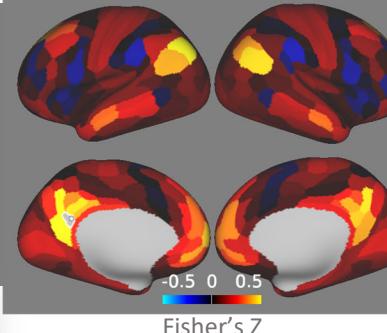


Residuals after GS regression

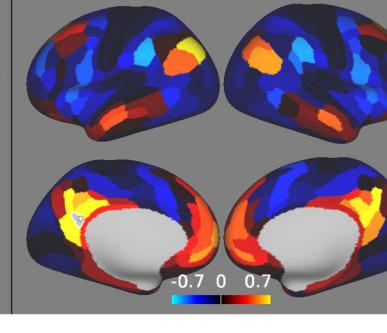


FIX-cleaned data Raw data

FIX-denoised rfMRI (No GSR)



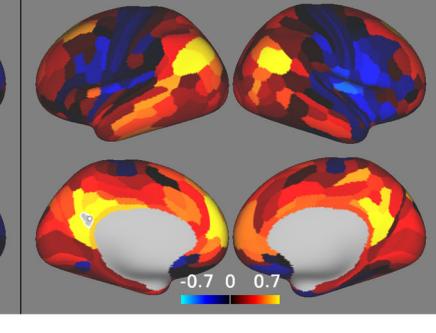
Raw rfMRI with GSR



Aso, Hayashi HBM2021

Lag structure

Perfusion lag structure with GSR



• GSR doesn't do!!

- The lag structure is a major part of global signal, but GSR leaves the correlation structure
- Tailored regression of sLFO for each voxel is needed

Erdogán (2016) FNS “Dynamic GSR”, Aso (2019) PLoS One

Method: Let's get rid of it

HCP test-retest dataset N=40

<https://github.com/RIKEN-BCIL/HCPstyle-BOLDLagMappingAndCleaning>



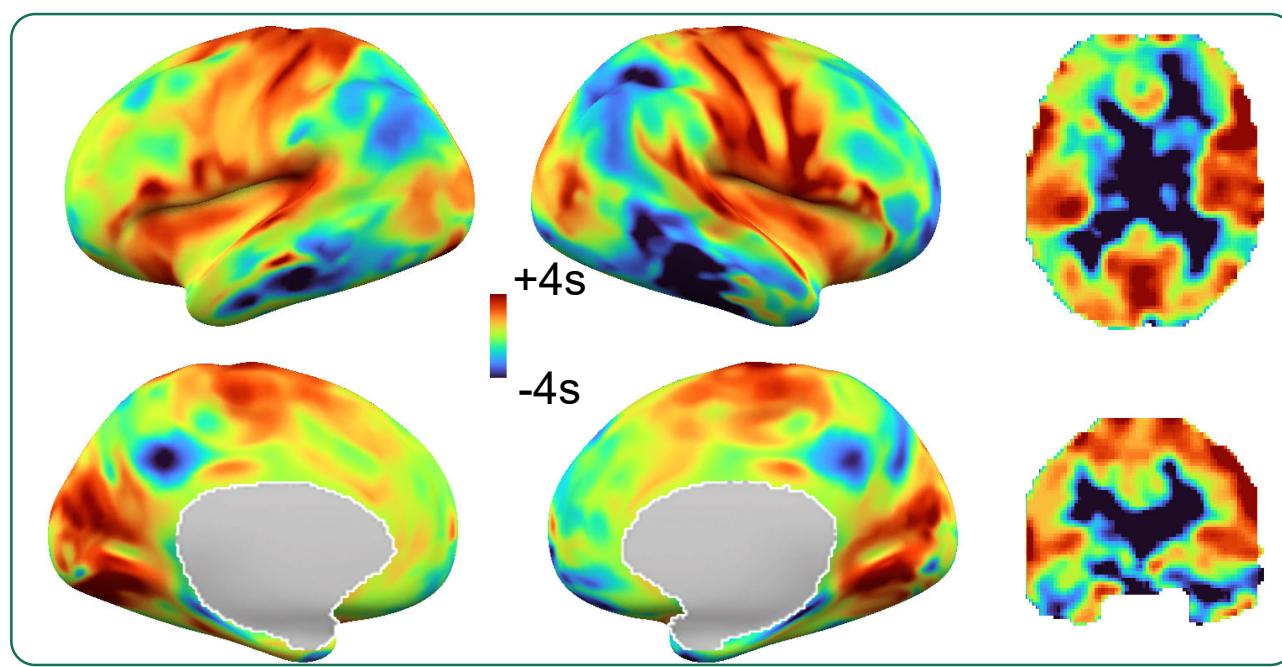
“De-perfusioning”
affects only slow components

ICA-FIX

ICA-FIX

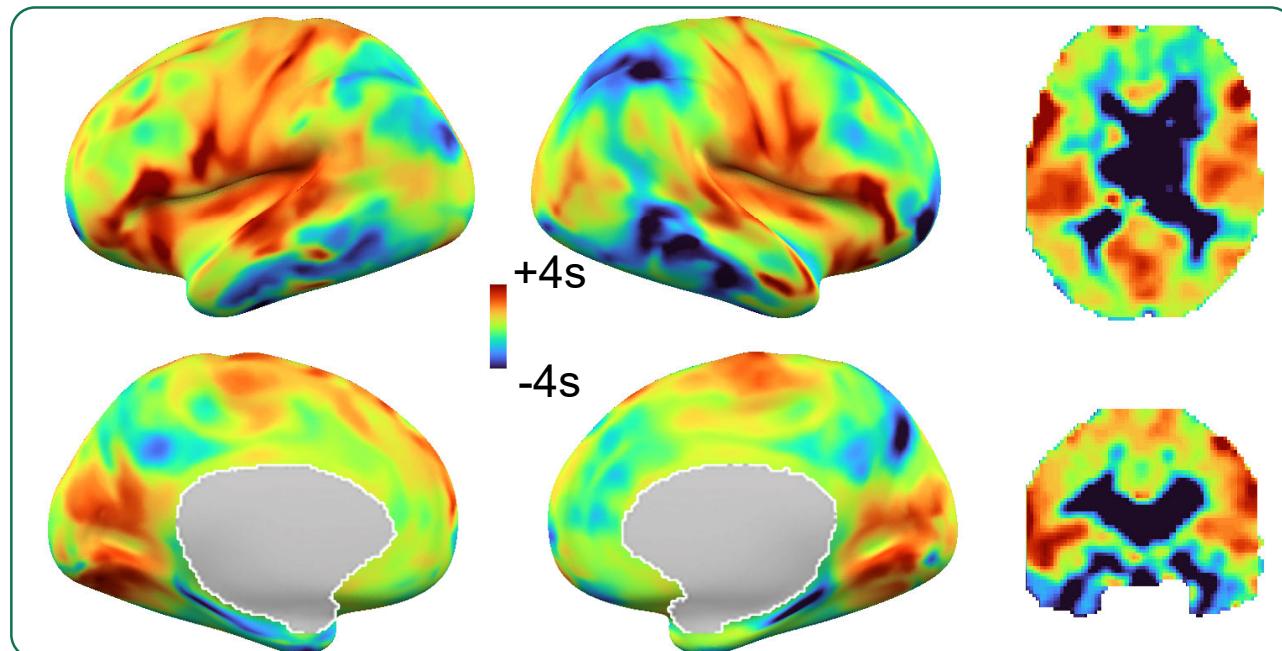
Compare Day-1 and Day-2

**Subject
#599671**
Lag map
day-1



**What was removed
by deperfusion:
Consistent across
visits,
because it's
predominantly
vascular anatomy**

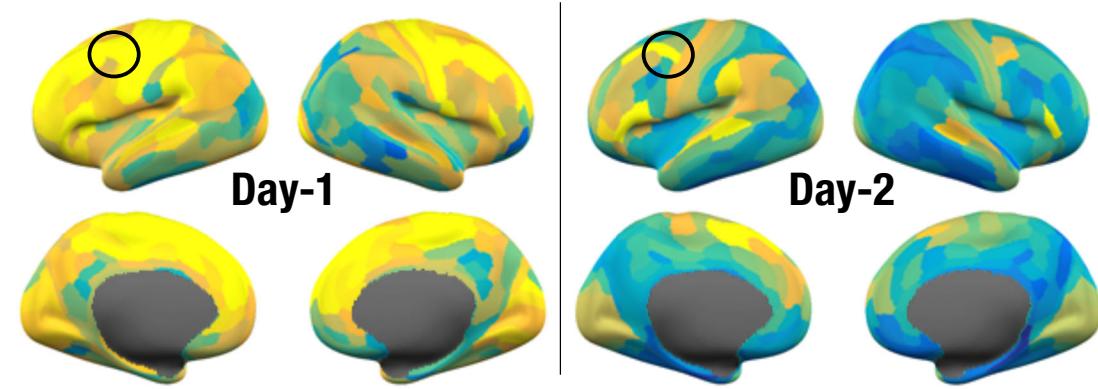
**Lag map
day-2**



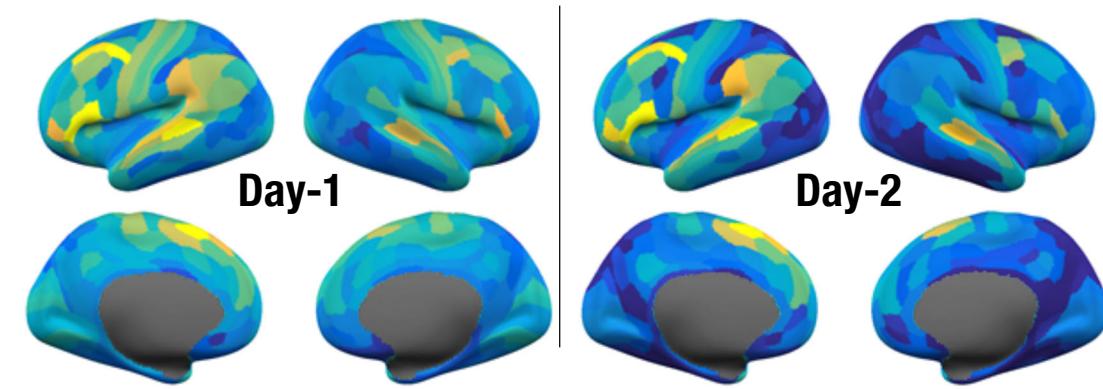
Subject #599671, Superior Language Network/Area 55b seed

Glasser (2016) Nature

Raw data



Deperfusing



Day-1

Day-2

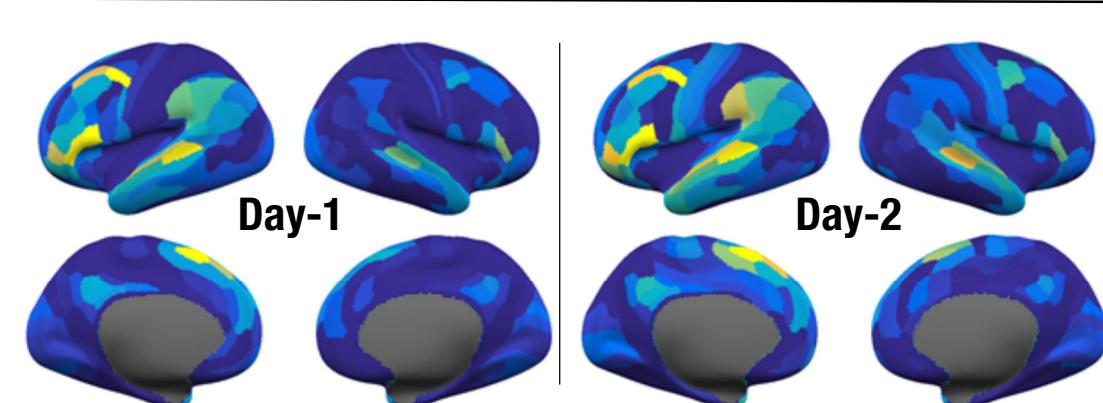
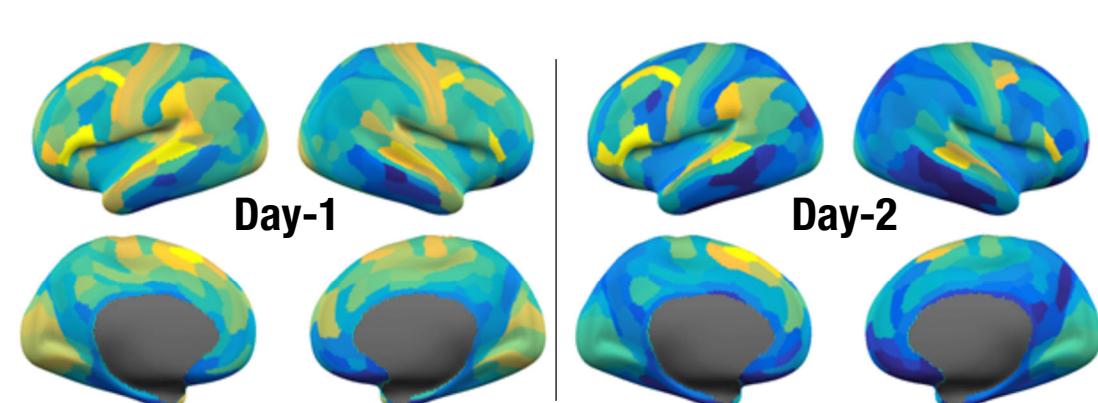
Day-1

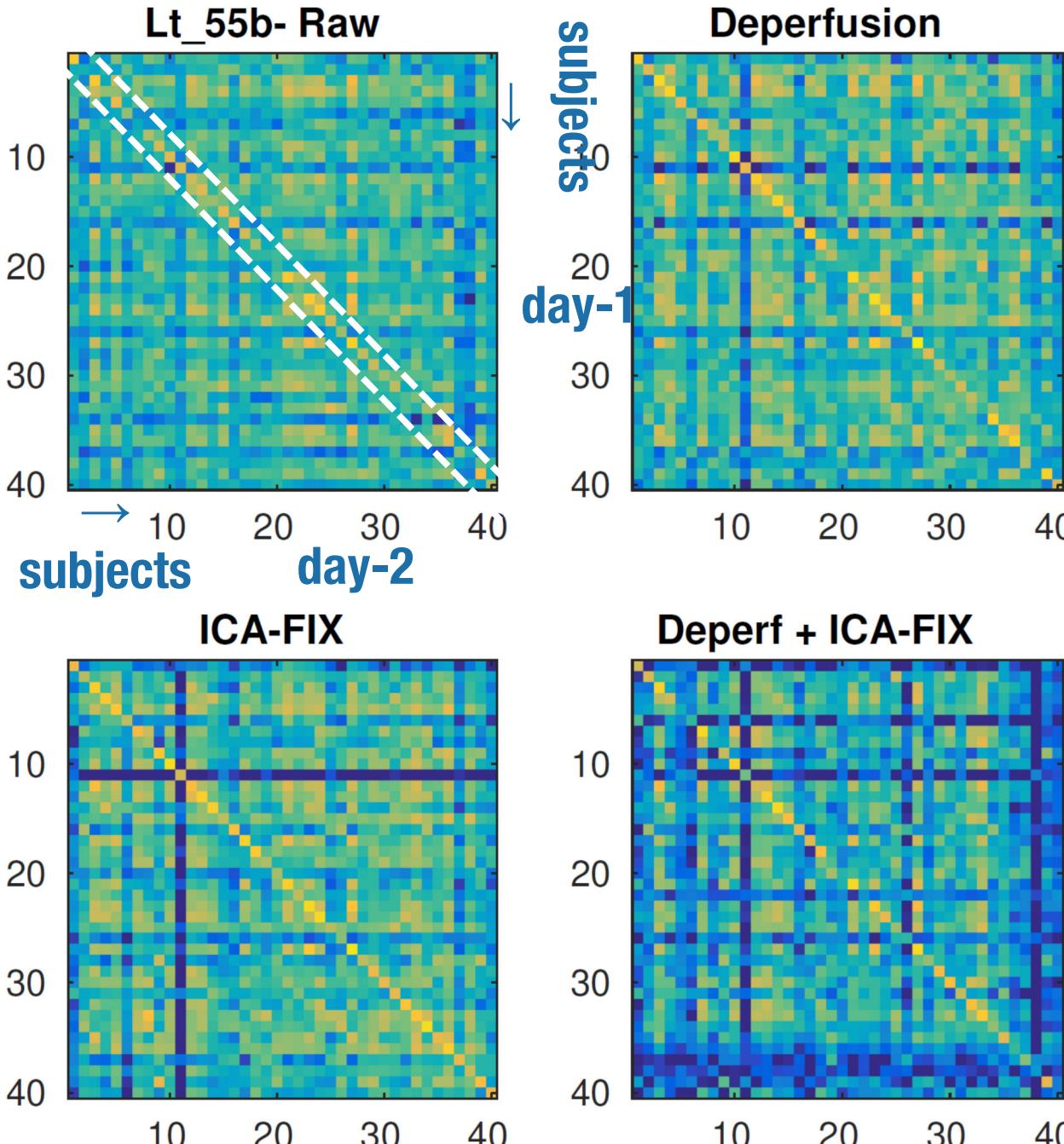
Day-2

ICA-FIX



ICA-FIX after Deperf

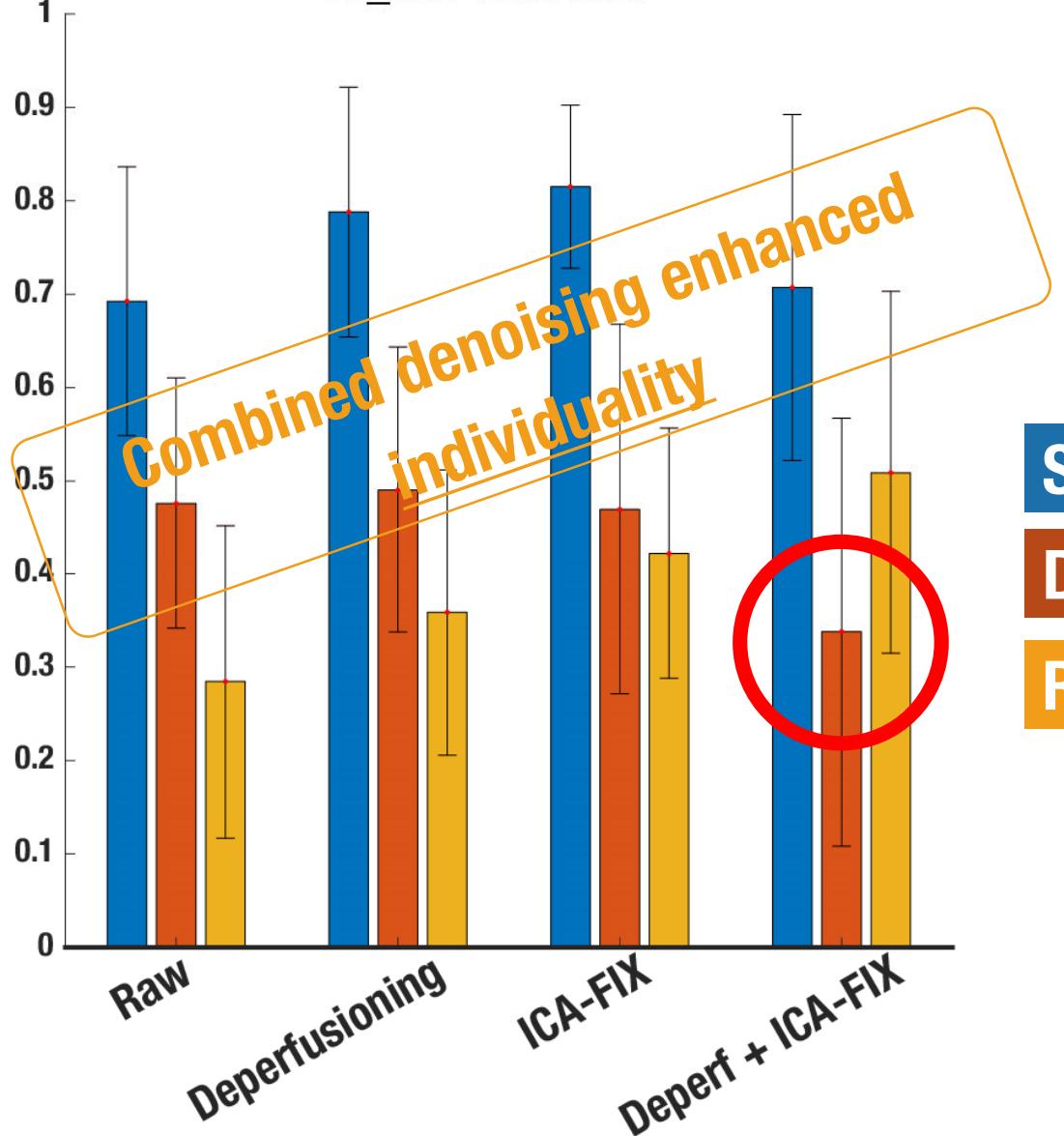




- # Network similarity as a reproducibility measure
- Every combination b/w day-1 and 2 (Spearman's rho)
 - Diagonal elements are within-subject, i.e., reproducibility

Denoising makes them stand out by diminished between-subject similarity!

Lt_55b-Area 55b



Individuality:
relative within-subject reproducibility

Same subject

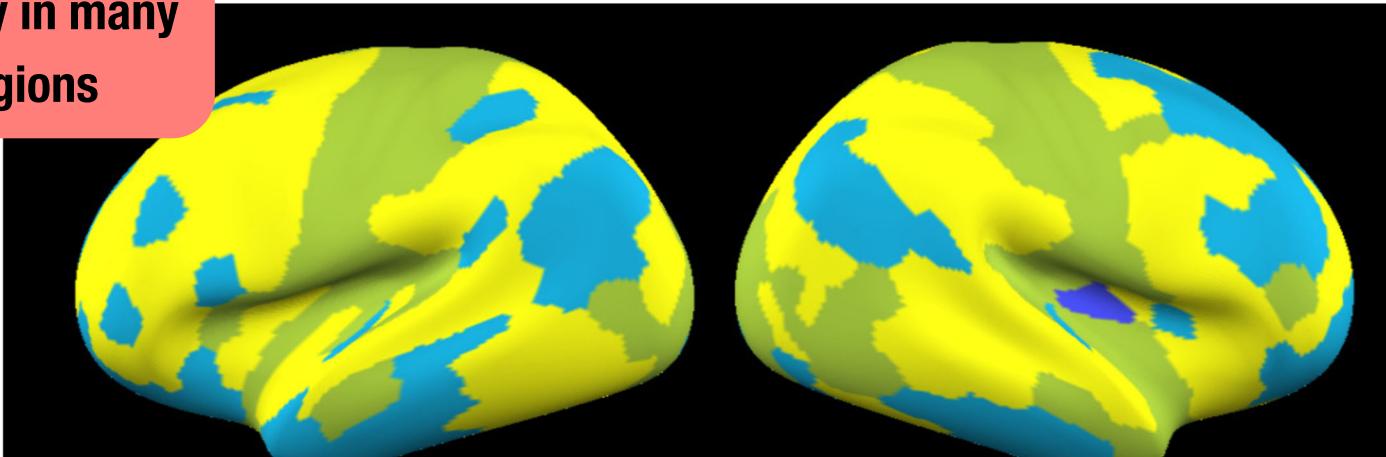
Different subject

Relative reproducibility (ratio)

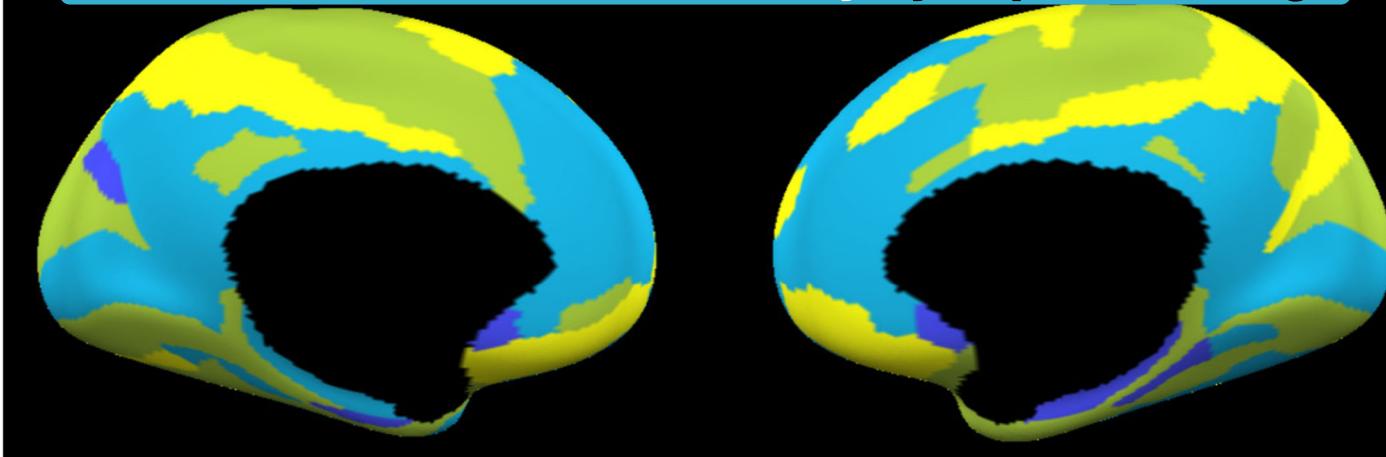
How about the
other seeds?

Best denoising (for individuality) for each seed region

Combined method
enhanced
individuality in many
seed regions



DMN seems to lose individuality by deperfusioning



Dep+ICAFIX

Combined method

Deperfusing

ICA-FIX

Raw

Summary

- Test-retest reproducibility of FC generally improved by the two denoising procedures, suggesting both approaches are basically valid.
- The deperfusioning+ICA-FIX combined method achieved highest relative reproducibility (=individuality) by suppressing between-subject similarity.
- Denoising effects were not uniform across networks, mirroring the previous works on physiology-driven spurious FC.
- The combined method is recommended.

“...low reproducibility does not necessarily imply low accuracy”
(Harrison 2015 *NeuroImage*)