## MR micro- and macrostructural biomarkers inked to nonpathological brain aging in

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## Introcuction

## Using a robust Magnetic Resonance Imaging (MRI) microscopy battery composed of • Diffusion MRI (dMRI) • Relaxometry based MRI (rMRI) Goal: To quantify changes in brain structure related to normal aging.



• Cohort Size: 12 female non-human primates (bonnet macaques) brains provided by the Barnes lab. Ages ranging from 10 to 25 years old (30 to 75 human equivalent years).

Ex-vivo MRI battery is developed to examine brain specimens at high resolution (200-600 micron scale) for quantitative biomarkers that indicate age-related brain dysfunction.



## <u>Multiple Spin Echo (MSE) – rMRI Technique:</u> Myelin Water Fraction (MWF) – Quantifies water trapped between myelin bilayers (a marker for myelin content – quantifies myelin found in each voxel).

## Selective Inversion Recovery (SIR) - rMRI Technique: **Bound Pool Fraction (BPF)** – Measures myelin content and is sensitive to molecular size.

**Diffusion Tensor Imaging (dMRI):** Fractional Anisotropy (FA) – Reports that there is a preferred direction of water diffusion in each voxel. **Trace (TR)** – Reports the total water diffusion regardless of direction.

## Background

## **Acquired Multiple Spin Echoes**



N = 1 - 32 (only showing 27 for space reasons) TE (Echo Times) = 6.2 - 198.4 ms





## Summary of Methods

## **Sample Preparation**

Post-mortem Bonnet Macaque Brain

Medical Gauze



## Map Generation

Different pipelines avenues depending on scan type.



MWF





Vacuum Chamber





BPF



DEC



TORTOISE Unix/Bash

## REMMI MATLAB





## Summary of Methods

## **Image Registration** Using Advanced Normalization Tools (ANTs) to

## **Voxel-wise Analysis**

Using fsl randomise tool threshold free cluster enhancement and assessing negative and positive correlations with age.









## **ROI Analysis** Masked high resolution anatomical (HRA) and conducted a Pearson correlation test across cohort.





## **Results – Voxel-wise Analysis**

## **Fractional Anisotropy**



## Trace







## **Myelin Water Fraction**

**Threshold Free Cluster** Enhancement corrected p value 5000 permutations (conservative)

0.05

p value

p value

0

0

0.05

## Results - ROI Analysis

## • Found strong correlations when assessing interactions between different MR metrics to one another.













## White Matter: MWF vs. BPF

## Concusions

In Voxel-wise and ROI analysis with respect to aging: No significant positive or negative correlations were found between our MR metrics and age using voxel-wise analysis. No significant positive or negative correlations were found in the hippocampus or white matter with respect to age using ROI analysis. Both analyses produced consistent results with regards to aging.

In ROI analysis with respect to MR metric correlation: Strong positive correlation with FA and MWF measures in white matter were observed.

 Strong positive correlation with BPF and MWF measures in the hippocampus. • This indicates that these 3 measures, depending on location in the brain, contain similar information.

• The combination of MR metrics may be better at identifying anatomical information than a single metric on its own.

## Euture Drectons

 Quantitative Susceptibility Maps (QSM) which is sensitive to iron and calcium accumulation. Voxel-wise and ROI analysis with behavioral data and age. • Use principal component analysis to perform dimensionality reduction to test if the combination multiple MRI metrics (features) can characterize normal aging. • ROI analysis of subfields of the hippocampus.





# TORTOISE (https://tortoise.nibib.nih.gov/)

ANTs (https://github.com/ANTsX/ANTs) Lowekamp BC, Chen DT, Ibáñez L and Blezek D (2013) The Design of SimpleITK. Front. Neuroinform. 7:45. doi: 10.3389/fninf.2013.00045 Z. Yaniv, B. C. Lowekamp, H. J. Johnson, R. Beare, "SimpleITK Image-Analysis Notebooks: a Collaborative Environment for Education and Reproducible Research", J Digit Imaging., https://doi.org/10.1007/s10278-017-0037-8, 2017. Winkler AM, Ridgway GR, Webster MA, Smith SM, Nichols TE. Permutation inference for the general linear model. NeuroImage, 2014;92:381-397.

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## References

REMMI-matlab (https://github.com/remmi-toolbox/remmi-matlab)