

# Quantifying exchange using optimized small-tip fast recovery (STFR) sequences

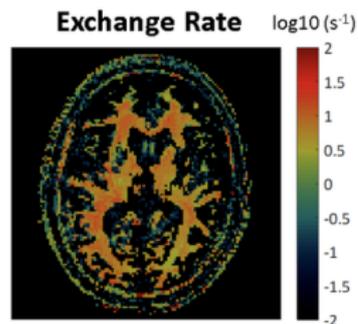
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University of Michigan

MRI Workshop on Acquisition and Reconstruction

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# Motivation: Mapping exchange



- ▶ Exchange captures inter-compartmental interactions (exchange of water + electrolytes)

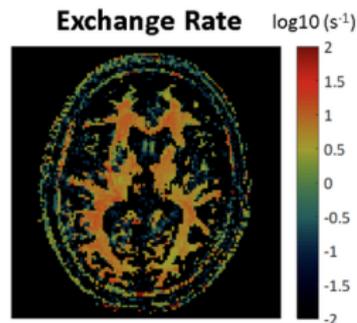
**Figure:** 2-D exchange map from MRF-X<sup>1</sup>

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<sup>1</sup>Hamilton, Jesse, et al. "MR fingerprinting with chemical exchange (MRF-X) for in vivo multi-compartment relaxation and exchange rate mapping." ISMRM 2016.

<sup>2</sup>Harkins, Kevin D., et al. "Effect of intercompartmental water exchange on the apparent myelin water fraction in multiexponential T2 measurements of rat spinal cord." MRM 67.3 (2012): 793-800.

# Motivation: Mapping exchange



- ▶ Exchange captures inter-compartmental interactions (exchange of water + electrolytes)
- ▶ Exchange can be a useful clinical biomarker<sup>2</sup>

**Figure:** 2-D exchange map from MRF-X<sup>1</sup>

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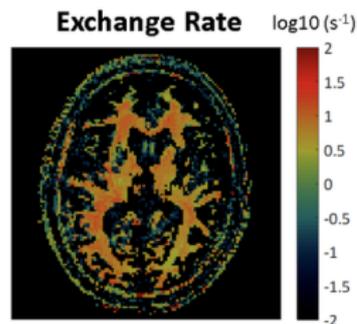


Figure: 2-D exchange map from MRF-X<sup>1</sup>

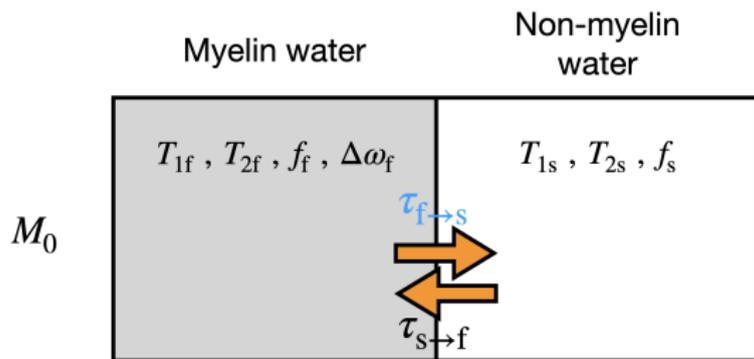
- ▶ Exchange captures inter-compartmental interactions (exchange of water + electrolytes)
- ▶ Exchange can be a useful clinical biomarker<sup>2</sup>
- ▶ Hard to estimate exchange *in vivo* currently

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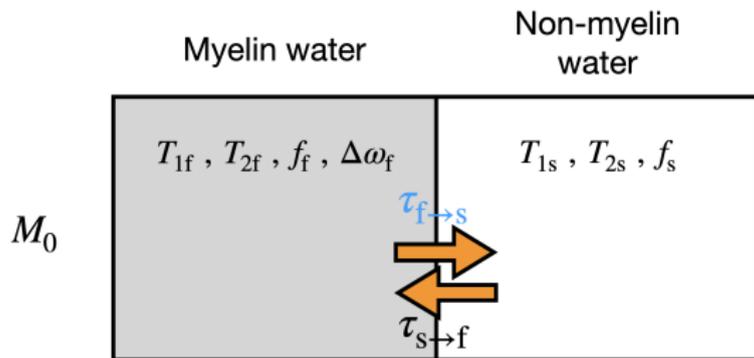
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# Two pool exchanging model



$\Delta\omega_f$  : Myelin-specific frequency shift

# Two pool exchanging model

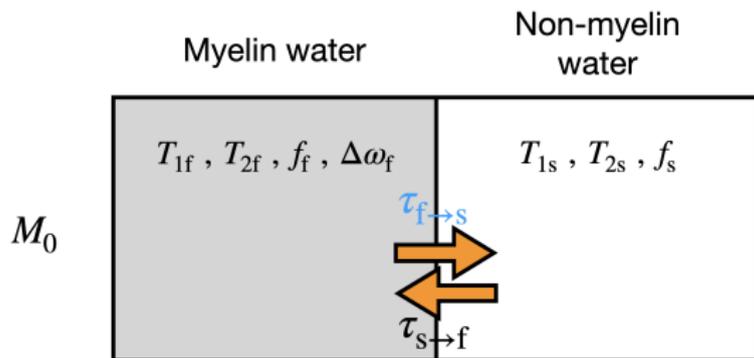


## Objectives

- ▶ Obtain a scan design that is able to estimate exchange with high precision
- ▶ Quantify exchange rates using an appropriate estimator

$\Delta\omega_f$  : Myelin-specific frequency shift

# Two pool exchanging model



## Setup

- ▶  $f_f + f_s = 1$
- ▶ Well-mixed:  $f_f/f_s = \tau_{f \rightarrow s}/\tau_{s \rightarrow f}$
- ▶ 8 unknowns:  $(M_0, f_f, T_{1f}, T_{1s}, T_{2f}, T_{2s}, \tau_{f \rightarrow s}, \Delta\omega_f)$

$\Delta\omega_f$  : Myelin-specific frequency shift

## Scan design

- ▶ Cramer-Rao Bound (CRB) based optimization (weighted wrt exchange)
- ▶ Pulse sequence: Small-tip Fast Recovery (STFR) sequence<sup>3</sup>

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<sup>3</sup>Sun, Hao, et al. "Steady-state functional MRI using spoiled small-tip fast recovery imaging." *Magnetic resonance in medicine* 73.2 (2015): 536-543.

# High-level overview

## Scan design

- ▶ Cramer-Rao Bound (CRB) based optimization (weighted wrt exchange)
- ▶ Pulse sequence: Small-tip Fast Recovery (STFR) sequence<sup>3</sup>

## Estimation

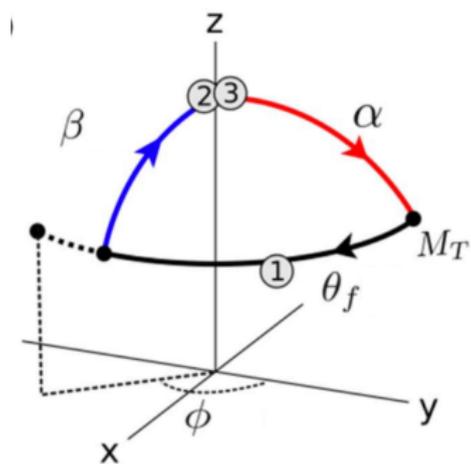
- ▶ Parameter Estimation via Regression with Kernels (PERK)<sup>4</sup>
- ▶ Fast dictionary-free parameter estimation method
- ▶ Lifts measurements to a higher-dimensional space followed by kernel regression

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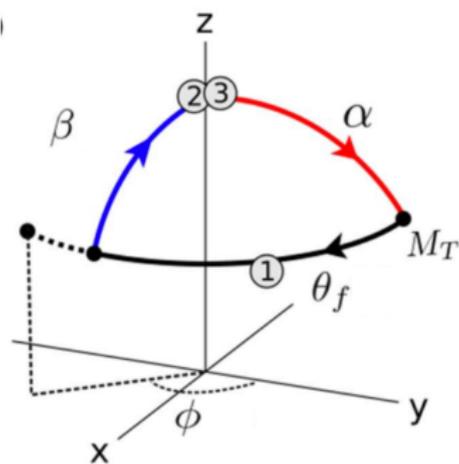
<sup>4</sup>Nataraj, Gopal, et al. "Dictionary-free MRI PERK: Parameter estimation via regression with kernels." *IEEE transactions on medical imaging* 37.9 (2018): 2103-2114.

# Small-tip Fast Recovery (STFR)



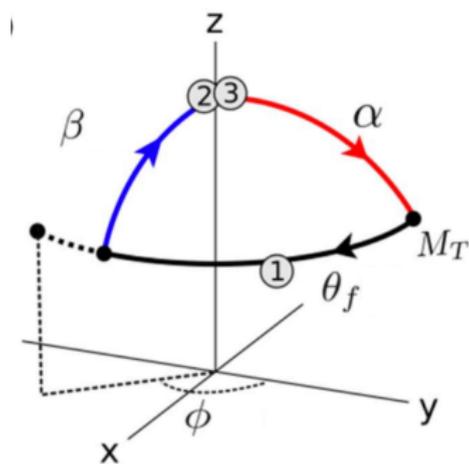
- Goal: Design a set of  $D$  STFR scans.

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- ▶ Scan parameters:  
 $(T_{\text{free}}, T_g, \alpha, \beta, \phi, T_E)_d$ ,  
where  $d = 1, 2, \dots, D$ .

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## Hypothesis

Optimized STFR sequences are sensitive to  $\Delta\omega_f$ , which might lead to more precise exchange estimates.

$\Delta\omega_f$  : Myelin-specific frequency shift

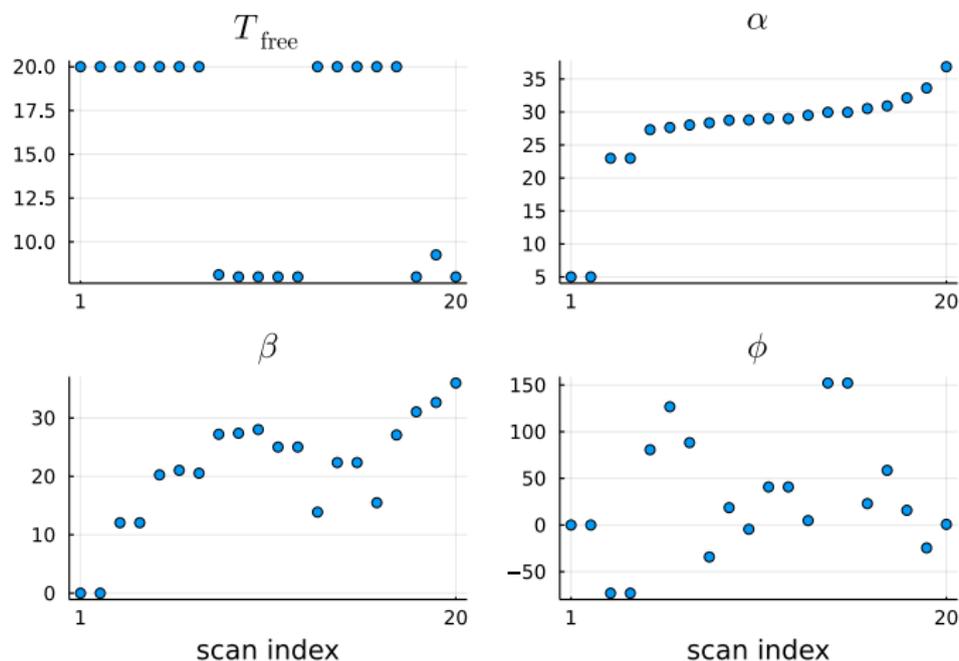
## Setup:

- ▶  $D = 20$  scans
- ▶ max FA =  $40^\circ$ , range of  $T_{\text{free}} = 8\text{-}20\text{ms}$

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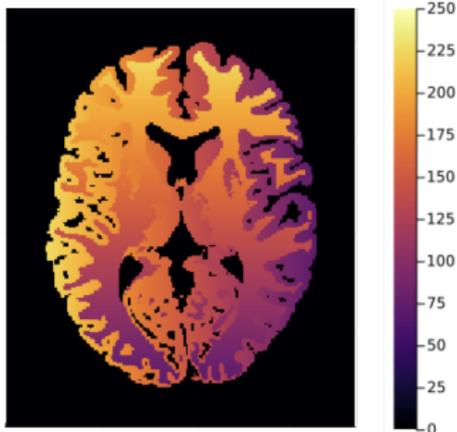
- ▶  $D = 20$  scans
- ▶ max FA =  $40^\circ$ , range of  $T_{\text{free}} = 8\text{-}20\text{ms}$
- ▶ B0 and B1+ maps are computed through separate SPGR and Bloch-Siegert scans
- ▶ Total scan time:  $\sim 21$ mins for 10 slices

# Optimized STFR design



- ▶ Coeff. of variation is **20%** for exchange estimation in WM (vs. 93% for initial design).

# Simulations



(a) Exchange map - ground truth (in ms) <sup>5</sup>

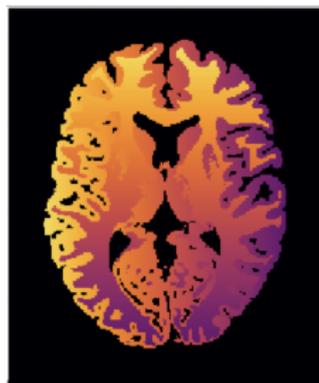
Parameter	White Matter	Gray Matter
$M_0$	0.77	0.86
$f_f$	0.15	0.03
$T_{1,f}$ (ms)	400	500
$T_{1,s}$ (ms)	832	1331
$T_{2,f}$ (ms)	20	20
$T_{2,s}$ (ms)	80	80
$\Delta\omega_f$ (Hz)	15	5
$\Delta\omega$ (Hz)	Varies	Varies
$\kappa$	Varies	Varies

(b) Nominal values for WM and GM at 3T <sup>6</sup>

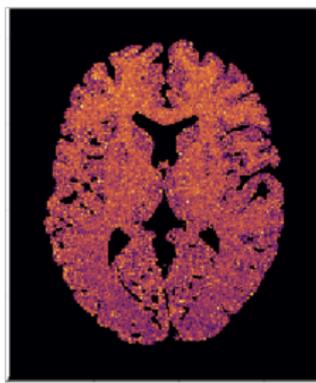
<sup>6</sup>Collins, D. Louis, et al. "Design and construction of a realistic digital brain phantom." *IEEE TMI* 1998.

<sup>7</sup>Whitaker, Steven T., et al. "Myelin water fraction estimation using small-tip fast recovery MRI." *MRM* 2020.

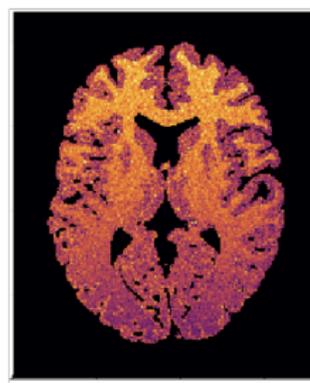
# Results



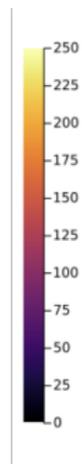
Ground truth



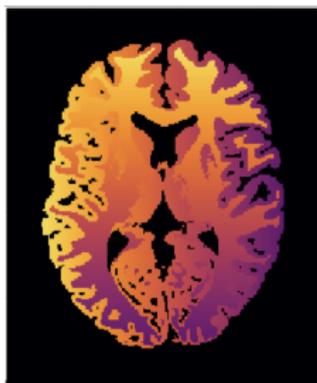
Initial rand design  
(RMSE in WM: 38ms)



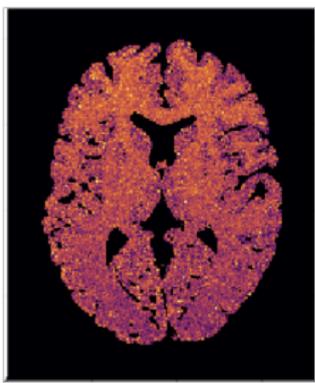
Opt STFR design  
(RMSE in WM: **23ms**)



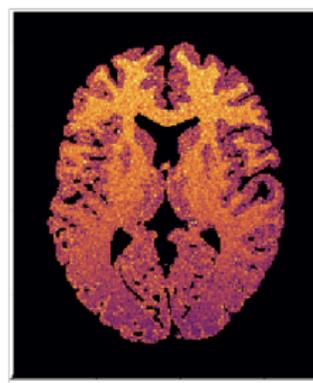
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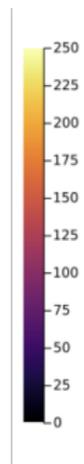
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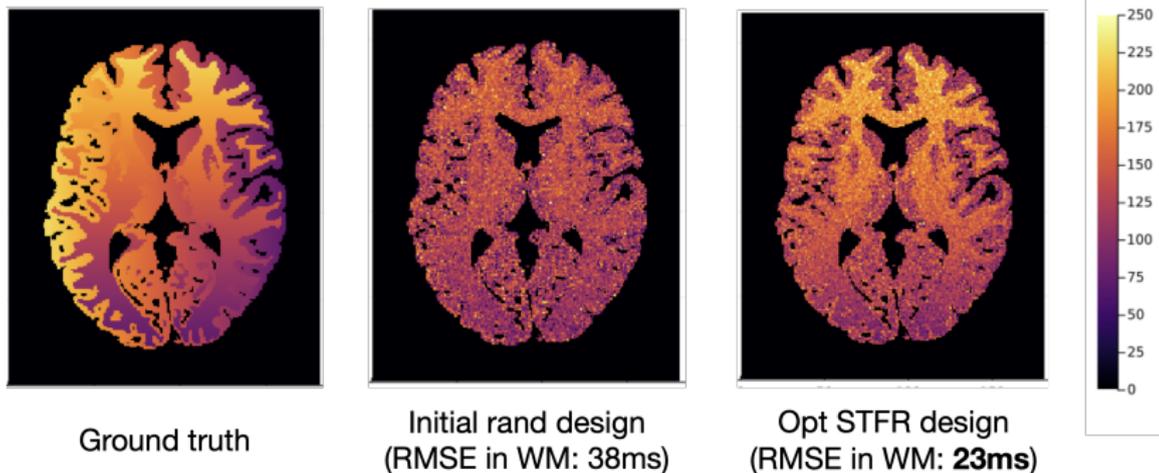


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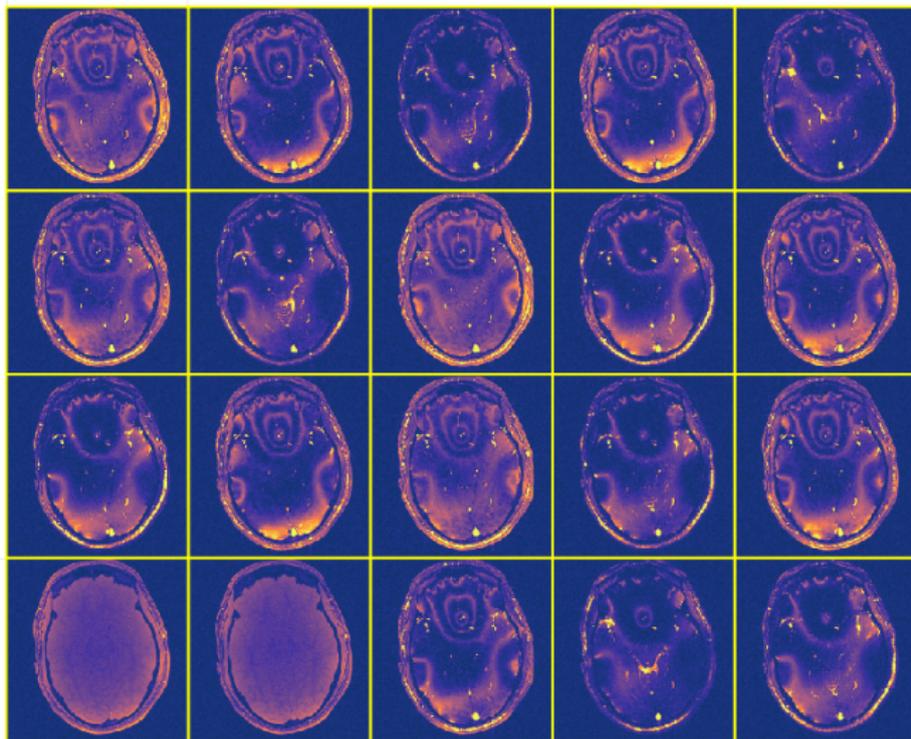
- ▶ Opt STFR design has an RMSE of **15%** in WM (vs. 25% for initial design).

# Results



- ▶ Opt STFR design has an RMSE of **15%** in WM (vs. 25% for initial design).
- ▶ Performs as worse as a mean estimator in GM (caveat: very little myelin content in GM)

# Invivo data (in progress)



Acknowledgements for data: Melissa Haskell

## Conclusions:

- ▶ Obtained an optimized STFR design that has  $\sim 20\%$  coefficient of variation for estimating exchange in WM.
- ▶ Optimized design performs better in WM than a random design (by  $\sim 10\%$ ); does as worse as a mean estimator in GM.

## Conclusions:

- ▶ Obtained an optimized STFR design that has  $\sim 20\%$  coefficient of variation for estimating exchange in WM.
- ▶ Optimized design performs better in WM than a random design (by  $\sim 10\%$ ); does as worse as a mean estimator in GM.

## Next steps:

- ▶ Investigate the performance of the estimator in the presence of off-resonance.
- ▶ Investigate how the opt design does for invivo data; compare with other state-of-the-art methods.