# Optimizing MR Scan Design for Parameter Estimation (with Application to $T_1$ , $T_2$ Relaxometry)

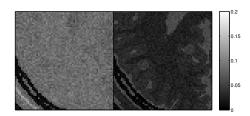
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Student SPEECS Seminar December 11, 2015

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  - Motivation
  - Problem Formulation
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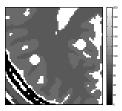
### Why Quantitative MRI?



(a) Anatomical Image



(b) Latent  $T_1$  Map



(c) Latent  $T_2$  Map

### Anatomical MRI: seek to reconstruct qualitative images

- √ Linearly related via Fourier transform to raw k-space data
- Same anatomy + varied acquisitions = varied image contrasts!
- Confounds nuisance contrast mechanisms with those of interes

#### **Quantitative** MRI: seek to estimate *intrinsic* parameters of interes

- Parameter maps are physical and have direct medical relevance
- √ Tissue alterations detectable with high sensitivity
- Many studies suggest (potential) clinical applications
  - Brain: multiple sclerosis, epilepsy, Parkinson's, ...
  - Other: cartilage degeneration, cardiac edema/infarction, ...
- $\nearrow$  In general, *nonlinearly* related to **k**-space data  $\rightarrow$  challenging recon
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### How to popularize QMRI clinically?

#### Multidisciplinary approaches:

- Health sciences: find specific applications for which QMRI outperforms as a diagnostic or prognostic tool (Cheng et al., 2012)
- Hardware engineering: improve MR hardware to produce better data (higher SNR, better field uniformity, etc.) (Roemer et al., 1990)
- Image reconstruction: for a given dataset, estimate latent parameters of interest rapidly and "reliably" (Nataraj et al., 2014)
- Data acquisition: prescribe a fast scan profile, or a combination of scan parameters from one or more pulse sequences, that enables "good" parameter estimation
  - Prior work: measured with CNR variations (Deoni et al., 2003, 2004)
  - This talk: measured with estimator **precision** (Nataraj et al., 2015)

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### **Problem Statement**

We seek a systematic method to guide **robust scan design** to enable **precise** latent object parameter estimation.

Scan design consists of two subproblems:

- Scan profile selection Given a collection of candidate pulse sequences, how to best assemble a scan profile?
- Scan parameter optimization For a fixed time constraint, how to optimize a given scan profile's acquisition parameters for latent object parameter estimation?

Robust means unbiased estimators maintain high precision across a wide range of object parameters



### General signal model

Many MR pulse sequences yield images (at position  $\mathbf{r}$ ) described as:

$$y_d(\mathbf{r}) = f_d(\mathbf{x}(\mathbf{r}); \mathbf{v}(\mathbf{r}), \mathbf{p}_d) + \epsilon_d(\mathbf{r}), d = 1, \dots, D$$
 (1)

#### Notation:

- $\mathbf{x}(\mathbf{r}) \in \mathbb{C}^L$  collects L latent object parameters at  $\mathbf{r}$
- $\mathbf{v}(\mathbf{r}) \in \mathbb{C}^K$  collects K known object parameters at  $\mathbf{r}$
- $\mathbf{p}_d \in \mathbb{R}^P$  denotes set of P scan parameters for dth dataset
- $\epsilon_d(\mathbf{r}) \sim \mathbb{C}\mathcal{N}(0, \sigma_d^2)$  modeled as independent, complex Gaussian noise<sup>1</sup>

## Scan profile model

A candidate scan profile collects *D* datasets from a combination of (possibly different) pulse sequences:

$$\mathbf{y}(\mathbf{r}) = \mathbf{f}(\mathbf{x}(\mathbf{r}); \mathbf{v}(\mathbf{r}), \mathbf{P}) + \boldsymbol{\epsilon}(\mathbf{r})$$
 (2)

#### Notation:

- $\mathbf{y}(\mathbf{r}) := [y_1(\mathbf{r}), \dots, y_D(\mathbf{r})]^\mathsf{T} \in \mathbb{C}^D$  collects noisy signals
- $\mathbf{f}: \mathbb{C}^L \times \mathbb{C}^K \times \mathbb{R}^{P \times D} \mapsto \mathbb{C}^D$  naturally extends scalar function f
- $\mathbf{P} := [\mathbf{p}_1, \ldots, \mathbf{p}_D] \in \mathbb{R}^{P \times D}$  gathers all scan parameters
- $oldsymbol{\epsilon} \in \mathbb{C}^D$  denotes Gaussian noise with diagonal covariance matrix  $oldsymbol{\Sigma}$

#### The Cramér-Rao Bound

Log-likelihood function (to within a constant c):

$$\ln L(\mathbf{x}(\mathbf{r})) = -\frac{1}{2} \|\mathbf{y}(\mathbf{r}) - \mathbf{f}(\mathbf{x}(\mathbf{r}); \boldsymbol{v}(\mathbf{r}), \mathbf{P})\|_{\mathbf{\Sigma}^{-1/2}}^2 + c$$
 (3)

Fisher information matrix: useful for characterizing estimator precision:

$$\mathbf{F}(\mathbf{x}(\mathbf{r}); \boldsymbol{\nu}(\mathbf{r}), \mathbf{P}) := \mathbb{E}\left(\left[\nabla_{\mathbf{x}} \ln L(\mathbf{x}(\mathbf{r}))\right]^{\dagger} \left[\nabla_{\mathbf{x}} \ln L(\mathbf{x}(\mathbf{r}))\right]\right)$$

$$= \left[\nabla_{\mathbf{x}} \mathbf{f}(\mathbf{x}(\mathbf{r}); \boldsymbol{\nu}(\mathbf{r}), \mathbf{P})\right]^{\dagger} \mathbf{\Sigma}^{-1} \left[\nabla_{\mathbf{x}} \mathbf{f}(\mathbf{x}(\mathbf{r}); \boldsymbol{\nu}(\mathbf{r}), \mathbf{P})\right]$$
(4)

(Matrix) Cramér-Rao Bound on covariance of unbiased estimates:

$$\operatorname{cov}(\widehat{\mathbf{x}}(\mathbf{r}); \boldsymbol{\nu}(\mathbf{r}), \mathbf{P}) \ge \mathbf{F}^{-1}(\mathbf{x}(\mathbf{r}); \boldsymbol{\nu}(\mathbf{r}), \mathbf{P}) \tag{5}$$

### Towards an Objective Function

#### Desirable to choose P such that precision matrix $F^{-1}$ "small"

Statisticians have considered minimizing various summary statistics:

$$ullet$$
 G-optimality: max diag  $\left( \mathbf{F}^{-1} \right)$  (Smith, 1918)

• *D*-optimality: 
$$det(\mathbf{F}^{-1})$$
 (Wald, 1945)

• A-optimality: 
$$\operatorname{tr}\left(\mathbf{F}^{-1}\right)$$
 (Chernoff, 1953)

. . . .

We consider a weighted variation of A-optimality

$$\Psi(\mathbf{x}(\mathbf{r}); \boldsymbol{\nu}(\mathbf{r}), \mathbf{P}) = \operatorname{tr}\left(\mathbf{W}\mathbf{F}^{-1}(\mathbf{x}(\mathbf{r}); \boldsymbol{\nu}(\mathbf{r}), \mathbf{P})\mathbf{W}^{\top}\right)$$
(6)

• Diagonal weight matrix  $\mathbf{W} \in \mathbb{R}^{L \times L}$  controls relative importance of precisely estimating L latent object parameters



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### Min-max Optimization

Cannot minimize  $\Psi(x; \nu, P)$  directly due to spatial variation of  $x(\cdot)$  and  $\nu(\cdot)$ 

Instead, seek candidate scan parameters  $\breve{\mathbf{P}}$  that minimize the max cost  $\widetilde{\Psi}^t$ :

$$\check{\mathbf{P}} \in \underset{\mathbf{P} \in \mathcal{P}}{\arg \min} \widetilde{\Psi}^{\mathsf{t}}(\mathbf{P}), \qquad \text{where}$$
(7)

$$\widetilde{\Psi}^{t}(\mathbf{P}) = \max_{\substack{\mathbf{x} \in \mathcal{X}_{t} \\ \boldsymbol{\nu} \in \mathcal{N}_{t}}} \Psi(\mathbf{x}; \boldsymbol{\nu}, \mathbf{P}). \tag{8}$$

#### More notation:

- ullet Search space  ${\mathcal P}$  can incorporate scan time constraints
- Tight latent object parameter set  $X_t$  chosen based on **application**
- $\bullet$  Tight known object parameter set  $\mathcal{N}_t$  chosen using prior knowledge

### **Incorporating Robustness**

Generally,  $\Psi(\mathbf{x}; \boldsymbol{v}, \mathbf{P})$  is non-convex, and may have multiple global minimizers and/or near-global minimizers. Collect these candidates as

$$\mathcal{S} := \left\{ \mathbf{P} : \widetilde{\Psi}^t(\mathbf{P}) - \widetilde{\Psi}^t(\check{\mathbf{P}}) \le \delta \widetilde{\Psi}^t(\check{\mathbf{P}}) \right\}, \qquad \text{where } \delta \ll 1. \tag{9}$$

**Robustness problem** – select *one* scan parameter  $P^*$  that degrades least when worst-case cost viewed over *broadened* sets  $\mathcal{X}_b$  and  $\mathcal{N}_b$ :

$$\mathbf{P}^* = \underset{\mathbf{P} \in \mathcal{S}}{\text{arg min }} \widetilde{\Psi}^b(\mathbf{P}), \qquad \text{where}$$
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### Summary: Robust, Application-specific Scan Design

## Recall: We sought a systematic method to guide robust scan parameter optimization and scan profile selection

- $\checkmark$  Candidate scan parameters  $\mathcal S$  found via min-max problem (7)
- $\checkmark$  Robust parameter  $\mathbf{P}^*$  chosen from  $\mathcal{S}$  via robustness problem (10)
- Scan profile selection?...
  - Given some candidate pulse sequences, construct all possible scan profiles that satisfy constraints, *e.g.*, acquisition time
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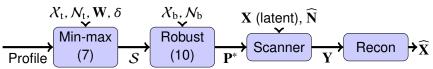
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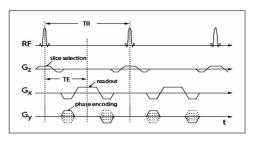
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#### MR Parameters of Interest



#### Prescribed scan parameters, p

- T<sub>R</sub>: repetition time between RF excitations
- a<sub>0</sub>: nominal flip angle by which spins are tipped

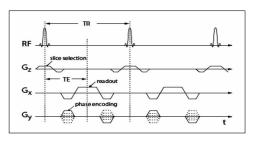
**Latent** object parameters,  $\mathbf{x}(\mathbf{r})$ 

- $T_1(\mathbf{r}), T_2(\mathbf{r})$ : longitudinal and transverse relaxation times (of interest)
- M<sub>E</sub>(r): spin density (nuisance)

Known object parameters, v(r)

•  $\kappa(\mathbf{r})$ : spatial variation in flip angle (true flip is  $a_0 \kappa(\mathbf{r})$ ),  $a_0 \kappa(\mathbf{r})$ 

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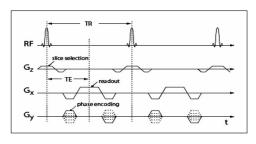
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### **Detailed Application**

#### **Example Problem**: scan design for joint $T_1$ , $T_2$ estimation in brain

- Candidate (fast) pulse sequences
  - Spoiled Gradient-Recalled Echo (SPGR): sensitive to T<sub>1</sub>
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- Candidate scan profiles
  - Profile consisting of  $C_{\rm SPGR}$  SPGR and  $C_{\rm DESS}$  DESS scans yields  $D = C_{\rm SPGR} + 2C_{\rm DESS}$  datasets
  - Can write SPGR, DESS signal models to group L = 3 latent object parameters  $\mathbf{x}(\mathbf{r}) := [M_{\mathrm{E}}(\mathbf{r}), T_1(\mathbf{r}), T_2(\mathbf{r})]^{\mathsf{T}}$  together
  - Prior works have considered T<sub>1</sub> and T<sub>2</sub> estimation from as few as
     2 SPGR (Deoni et al., 2003) or 1 DESS (Welsch et al., 2009) scan(s)
  - Examine scan profiles no longer than  $(C_{SPGR}, C_{DESS}) = (2, 1)$  profile
  - Ensuring  $D \ge L = 3$ , only other possibilities: (1, 1) and (0, 2)
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  - Two scan parameters  $\mathbf{p} := [a_0, T_R]$  available to optimize for each scan

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### Scan Profile Comparisons: Visualization

$$(C_{SPGR}, C_{DESS}) = (2, 1) (C_{SPGR}, C_{DESS}) = (1, 1) (C_{SPGR}, C_{DESS}) = (0, 2)$$

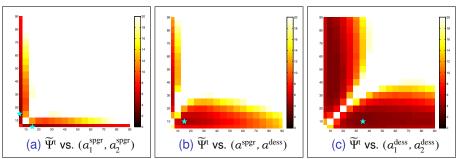


Figure 1: Appears that  $\widetilde{\Psi}^t$  at minimizers are similar, but the optimized (0,2) profile appears most robust to flip angle variation. All values in milliseconds.

### Scan Profile Comparisons: Performance Summary

Scan	$\widehat{a}_0^{ ext{spgr}}$	$\widehat{a}_0^{ ext{dess}}$	$\widehat{T}_R^{ ext{spgr}}$	$\widehat{T}_R^{\mathrm{dess}}$	$\widetilde{\Psi}^t(\pmb{P}^*)$	$\widetilde{\Psi}^b(\pmb{P}^*)$
(2, 1)	(15, 5)°	$30^{\circ}$	(12.2, 12.2)	17.5	4.0	17.7
(1, 1)	15°	10°	13.9	28.0	4.9	17.9
(0, 2)	_	(35, 10)°	_	(24.4, 17.5)	3.5	12.2

Table 1: Reflecting Fig. 1,  $\widetilde{\Psi}^b$  recommends (0,2) more emphatically than  $\widetilde{\Psi}^t$ . Flip angles are in degrees; all other values are in milliseconds.

#### New findings

- DESS sequences alone can be useful for precise T<sub>1</sub> mapping
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### Simple simulation study

Neglect to model several effects to simplify study of estimator statistics:

- No transmit field inhomogeneity
- No receive coil sensitivity variation
- No partial volume effects: deterministic knowledge of WM/GM ROIs
- ...

Max-likelihood (ML)  $T_1$ ,  $T_2$  estimation...

- ...using precomputed dictionary of signal vectors
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#### Estimator statistics

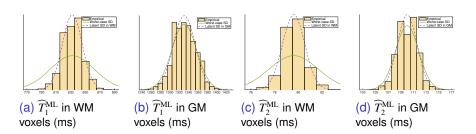


Figure 2: At realistic noise levels, ML estimates exhibit negligible bias and appear nearly Gaussian-distributed. Thus, CRB reliably approximates  $\widehat{T}_1^{\text{ML}}, \widehat{T}_2^{\text{ML}}$  errors.

### (Selected) Acquisition/Reconstruction Details

#### Fast steady-state acquisitions

- Combinations of (2, 1), (1, 1), and (0, 2) SPGR and DESS scans
- ullet Prescribe flip angles  $\widehat{m{a}}$  and repetition times  $\widehat{m{T}}_R$  in Table 1
- $256 \times 256 \times 6$  matrix over  $240 \times 240 \times 30$ mm FOV
- Effective scan time: 10.73s per slice

#### Slow reference acquisitions

- ullet Optimized combination of 2 IR scans for reference  $\widehat{T}_1$  map
- Optimized combination of 2 SE scans for reference  $\widehat{T}_2$  map
- 256 × 256 matrix over 24 × 24 × 5mm FOV
- Effective total scan time: 51m12s per slice

#### Reconstruction overview

- Regularized Least Squares (RLS) optimization using ML initialization, followed by alternating minimization
- Flip angle variation  $\widehat{\kappa}(\mathbf{r})$  separately estimated from pair of Bloch-Siegert (BS) shifted SPGR scans

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#### Slow reference acquisitions

- Optimized combination of 2 IR scans for reference  $\widehat{T}_1$  map
- Optimized combination of 2 SE scans for reference T<sub>2</sub> map
- 256 × 256 matrix over 24 × 24 × 5mm FOV
- Effective total scan time: 51m12s per slice

#### Reconstruction overview

- Regularized Least Squares (RLS) optimization using ML initialization, followed by alternating minimization
- Flip angle variation  $\widehat{\kappa}(\mathbf{r})$  separately estimated from pair of Bloch-Siegert (BS) shifted SPGR scans

### Phantom Results: T<sub>1</sub>

#### Coronal scans of **NIST MR system phantom**

(Russek et al., 2012)

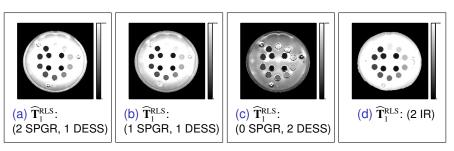


Figure 3:  $T_1$  RLS phantom estimates. Colorbar range is [0, 2000]ms.

### Phantom Results: T<sub>2</sub>

#### Coronal scans of **NIST MR system phantom**

(Russek et al., 2012)



Figure 4: T<sub>2</sub> RLS phantom estimates. Colorbar range is [0, 500]ms.

### Phantom Results: Comparisons

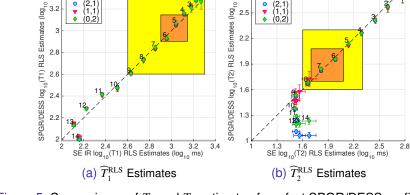


Figure 5: Comparisons of  $T_1$  and  $T_2$  estimates from fast SPGR/DESS profiles versus slow IR and SE profiles, respectively. Within tight and broad ranges of interest, estimates in good agreement.



### (Jeff's) Brain Results: T<sub>1</sub>

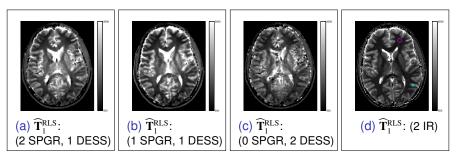


Figure 6:  $T_1$  RLS brain estimates. Colorbar range is [500, 2000]ms.

Scan	(2, 1)	(1, 2)	(0, 2)	(2 IR)
$\overline{WM}\widehat{T}_1^{\mathrm{RLS}}$	$773 \pm 51$	$711 \pm 53$	$721 \pm 38$	660. ± 13
$GM \; \widehat{T}_1^{RLS}$	$1110 \pm 160$	$1110\pm180$	$990 \pm 110$	$1029 \pm 39$

### (Jeff's) Brain Results: T2

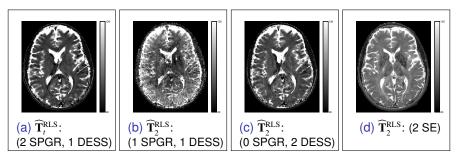


Figure 7:  $T_2$  RLS brain estimates. Colorbar range is [20, 120]ms.

Scan	(2, 1)	( ' /	· · /	, ,
$\overline{WM}\widehat{T}_2^{RLS}$	$42.3 \pm 3.3$		$45.5 \pm 3.6$	
$GM \; \widehat{T}_2^{RLS}$	$54 \pm 11$	$71 \pm 11$	$54.7 \pm 8.4$	$68.7 \pm 5.0$

### Summary and Future Directions

#### Summary

- Introduced a CRB-inspired min-max approach to aid robust, application-specific MR scan design
- Practical application: optimized (SPGR, DESS) combinations for  $T_1$ ,  $T_2$  relaxometry in WM/GM regions of the brain
- Numerical simulations + phantom and brain experiments

#### Ongoing and Future Work

- Scan design for est. flip angle scaling  $\kappa(\mathbf{r})$  also (Nataraj et al., 2014)
- Scan design when analytical signal model unknown

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Figure 8: http://collaborate.nist.gov/mriphantoms

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