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# 1 Purpose

This is a rough draft of the theory and methodology behind the Toeplitz subspace approach. This should help in understanding the code.

# 2 Theory

#### 2.1 Toeplitz Embedding

We first consider the non-Cartesian SENSE [1] forward model

$$\mathbf{F} \begin{bmatrix} \mathbf{S}_1 \\ \vdots \\ \mathbf{S}_C \end{bmatrix} \mathbf{x} = \mathbf{b} \tag{1}$$

where  $\mathbf{F}$  is the Non-Uniform Fourier Transform (NUFFT) operator,  $\mathbf{S}_i$  form a set of C sensitivity maps,  $\mathbf{x}$  is the volume of interest, and  $\mathbf{b}$  is the acquired data from the scanner. After acquiring the scanner data  $\mathbf{b}$ , the volume  $\mathbf{x}$  is usually reconstructed by solving the regularized inverse problem:

$$\underset{\mathbf{x}}{\operatorname{argmin}} ||\mathbf{A}\mathbf{x} - \mathbf{b}||_{2}^{2} + R(\mathbf{x}) \tag{2}$$

Due to the problem scale of MRI, it is not computationally tractable to analytically solve (2). Thus, iterative methods [2] are used to solve for the reconstructed volume. These iterative methods will almost always involve the repeated application of the Gram matrix  $\mathbf{A}^H \mathbf{A}$ . The Gram matrix is costly to compute due to the NUFFT term. An NUFFT consists of a Fast Fourier Transform (FFT) followed by a Kaiser-Bessel interpolation operation, which can become computationally expensive. Hence, it is desirable to eliminate the need for an NUFFT, and only use FFTs.

Previous works [3], [4] on Toeplitz embedding show that the operator  $\mathbf{F}^H\mathbf{F}$  is a shift-invariant operator and therefore has a block-Toeplitz structure that can be exploited:

$$\mathbf{F}^{H}\mathbf{F} = \tilde{R}^{H}\tilde{\mathcal{F}}^{H}\mathbf{diag}(\tilde{\mathbf{T}})\tilde{\mathcal{F}}\tilde{R}$$
(3)

Here, we use the  $\tilde{}$  notation to denote an operation with a twice oversampled representation.  $\tilde{\mathcal{F}}$  is a twice oversampled FFT,  $\tilde{R}$  zero pads the input volume by  $2\times$  in each spatial dimension,  $\tilde{\mathbf{T}}$  is called the Toeplitz

kernel, and the operator  $\tilde{R}^H$  crops a  $2\times$  padded volume down to the original dimensions of  $\mathbf{x}$ . According to [3], the Toeplitz kernel can be computed as

$$\tilde{\mathbf{T}} = \tilde{\mathcal{F}}\tilde{\mathbf{F}}^H\tilde{\mathbf{F}}\delta \tag{4}$$

where  $\tilde{\mathbf{F}}$  is a NUFFT on the zero-padded input volume, and  $\delta$  is the Dirac delta function. Avoiding Kaiser-Bessel interpolation by replacing  $\mathbf{F}^H\mathbf{F}$  with  $\tilde{R}^H\tilde{\mathcal{F}}^H\mathbf{diag}(\tilde{\mathbf{T}})\tilde{\mathcal{F}}\tilde{R}$  as in (3) leads to significant speedups, but comes at the expense of the memory required to store  $\tilde{\mathbf{T}}$ .

#### 2.2 Subspace Reconstruction with Toeplitz Embedding

Many time-resolved MRI acquisitions [5], [6], [7] reconstruct a time series of T volumes  $\mathbf{x} = (\mathbf{x}_1, \dots, \mathbf{x}_T)$ . We will assume that each time-point t with corresponding volume  $\mathbf{x}_t$  has a unique sampling trajectory, and hence will have a corresponding unique NUFFT  $\mathbf{F}_t$  with Toeplitz embedding  $\mathbf{diag}(\mathbf{T}_t)$ . The idea of subspace reconstructions [5], [6], [7] is to leverage a temporal subspace to reduce the dimensionality of the problem:

$$\begin{bmatrix} \mathbf{x}_1 \\ \vdots \\ \mathbf{x}_T \end{bmatrix} = \Phi \begin{bmatrix} \alpha_1 \\ \vdots \\ \alpha_K \end{bmatrix}$$
 (5)

where  $\Phi$  is the linear subspace with dimension [T, K]. We can incorporate this into the forward model as

$$\begin{bmatrix} \mathbf{F}_1 \\ \vdots \\ \mathbf{F}_T \end{bmatrix} \begin{bmatrix} \mathbf{S}_1 \\ \vdots \\ \mathbf{S}_C \end{bmatrix} \boldsymbol{\Phi} \boldsymbol{\alpha} = \underbrace{\mathbf{F}_s \mathbf{S} \boldsymbol{\Phi}}_{\mathbf{A}_s} \mathbf{b}$$
 (6)

where  $\mathbf{A}_s$  is the subspace forward operator, and  $\mathbf{F}_s$  contains the  $N_T$  NUFFTs. It can be shown that the subspace Gram matrix is

$$\mathbf{A}_{s}^{H}\mathbf{A}_{s}\alpha = \sum_{i=1}^{C} \mathbf{S}_{i}^{H} \tilde{R}^{H} \tilde{\mathcal{F}}^{H} \underbrace{\begin{bmatrix} \mathbf{M}_{11} & \cdots & \mathbf{M}_{1K} \\ \vdots & & \vdots \\ \mathbf{M}_{K1} & \cdots & \mathbf{M}_{KK} \end{bmatrix}}_{\mathbf{M}} \tilde{\mathcal{F}} \tilde{R} \mathbf{S}_{i} \begin{bmatrix} \alpha_{1} \\ \vdots \\ \alpha_{K} \end{bmatrix}$$
(7)

And the symmetric operator **M** is a grid structure of  $K^2$  Toeplitz kernels:

$$\mathbf{M}_{ij} = \sum_{t=1}^{T} \bar{\mathbf{\Phi}}_{ti} \mathbf{diag}(\tilde{\mathbf{T}}_t) \mathbf{\Phi}_{tj}$$
 (8)

Equations (7), (8), show that there exists a much more compact subspace Toeplitz matrix. This can be efficiently computed without the need to compute Toeplitz embeddings for each time point, as shown in Algorithm 1.

# 3 Methods

#### Algorithm 1 Computing Toeplitz Embedding Matrix M

Input: Over-sampled Time-Series NUFFTs  $\tilde{F}_s$ , Temporal Subspace  $\Phi$ 

1: **for**  $j \in \{1, \dots, K \text{ do } \}$ 

▶ The terms with ~are zero-padded by 2X

2:  $\tilde{\alpha} \leftarrow \tilde{0}$ 

▷ Initialize to zeros

3:  $\tilde{\alpha_i} \leftarrow \tilde{\delta}$ 

 $\triangleright$  Set the  $j^{\text{th}}$  coefficient to a twice-oversampled delta

4:  $\{M_{1j}\cdots M_{Kj}\} \leftarrow \tilde{\mathcal{F}}\Phi^H \tilde{F_s}^H \tilde{F_s}\Phi \tilde{\alpha}$ 

 $\triangleright$  Compute  $j^{\text{th}}$  column of M

5: end for

6: return M

From equations (4), (7), (8), we can formulate a simple, memory efficient algorithm for calculating  $\mathbf{M}$ . Instead of computing  $N_T$  Toeplitz embeddings, one can instead use a PSF approach similar to [3]. The details of this algorithm are shown in Algorithm 1. Note that we use the notation to denote image space zero-padding by a factor of 2. After completing Algorithm 1, one must store the Toeplitz embedding matrix  $\mathbf{M}$  in memory. Since  $\mathbf{M}$  is symmetric, we will need to store K(K+1)/2 embeddings. Each embedding has a twice oversampled volume size, so the total memory requirement will be  $\frac{K(K+1)}{2}(2N)^d$ , where d is the dimension of the problem (usually d=2 or d=3).

During image reconstruction, we will need to repeatedly apply  $\mathbf{M}$  to  $\alpha$ . In order to combat GPU memory constraints, we store the relevant terms of  $\mathbf{M}$  in CPU memory, and only move the rows of  $\mathbf{M}$  on and off the GPU to perform the dot product with  $\alpha$ .

#### References

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