

Mark Nishimura 1,* , Daniel Abraham 1,* , and Kawin Setsompop 1,2

¹Stanford EE ²Stanford Radiology *Equal contribution

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1 Purpose

This is a rough draft of the theory and methodology behind the Toeplitz subspace approach. We provide this document to assist 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:

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

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

where $\tilde{\mathbf{F}}$ is an NUFFT on the zero-padded input volume, and δ 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 a subspace reconstruction [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 Φ is the linear subspace with dimension [T, K]. The linear subspace represented by Φ is typically determined by taking the first K principal components of a matrix containing all possible signal evolutions. We can incorporate this into the forward model as

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

$$\tag{6}$$

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

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

The relationship between the symmetric operator \mathbf{M} and the time-series NUFFTs 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 Φ

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

 \triangleright The terms with are zero-padded by $2\times$

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

 \triangleright Initialize to zeros

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

 \triangleright Set the j^{th} coefficient to a twice-over sampled 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^{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 T Toeplitz kernels, we instead use a PSF approach similar to [3]. The details of this algorithm are shown in Algorithm 1. Note that we use the $\tilde{}$ 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 α . 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 α .

References

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