# Reconstruction of Undersampled Cardiac Cine MRI data Using Compressive Sensing Principles

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Abstract-Reduction of MRI data acquisition time is an important goal in the MRI field. Undersampling k-t space is a solution to reduce acquisition time. MR images may have sparse or compressible presentations in appropriate transform domains, such as Wavelet. According to the Compressive Sensing (CS) theory, they can be recovered from randomly undersampled k-t space data that guarantees incoherency between sparsifying transform and sampling operator. However, pure random k-space sampling can be more time-consuming than full k-space sampling because of MRI principles. In this paper, we propose a new method based on Hidden Markov Models (HMM) to undersample k-space along the phase-encoding direction. To this end, we cluster extracted features of each k-space line by fuzzy c-means (FCM) method and consider the resulting class labels as the states of a Markov chain. Then we train a HMM and find the related transition matrix to each k-space line. We choose the lines having more non-diagonal transition matrices to sample data along them. We reconstruct the image by minimizing the L1 norm subject to data consistency using conjugate-gradient method and use simulation to set the parameters of the proposed method (e.g., iteration number). We apply our method to reconstruct undersampled Cardiac Cine MRI data with and without sparsifying transform, successfully.

#### I. Introduction

It seems necessary to find some solutions to increase MRI speed without losing the quality of images because of developing of some time-consuming applications of MRI such as functional and spectroscopic MRI, imaging some moving organs (such as heart and joints) and need to improve spatial resolution at the same imaging time [1]. On the other hand, the hardware solution to reduce MRI data acquisition time, using more powerful gradient amplifiers, is limited by technical and biological considerations. Therefore, the researchers have searched some methods to use intrinsic redundancy and correlation of MRI data in k-t space and reconstruct images without complete sampling of k-t space. We can divide these approaches into the following three categories [2]:

- 1. Exploiting correlations in k-space, such as parallel imaging [3], [4],
- 2. Exploiting temporal correlations, such as UNFOLD [5] and 3. Exploiting correlations in both k-space and time domain, such as k-t BLAST [2].

We should emphasize that in the mentioned methods, the

images signal to noise ratio (SNR) decrease by the increase in speed.

In the next section we describe the used concepts in this article such as Compressive Sensing (CS) [6], [7] and Hidden Markov Models (HMMs). In the third part we attempt to detail the proposed method. Throughout the fuzzy clustering, will study the k-space lines by use of HMMs and then select the ones having more variations in time. Afterwards, we reconstruct the desired images by CS concepts. In the fourth section of article, we present some results of successful applying the proposed method to reconstruct undersampled Cardiac Cine MRI data. Finally, we present the strengths and drawbacks of our method and give some suggestions to improve it.

#### II. THEORY

In this section, we introduce the theory of compressive sensing [8] then review two major attempts to speed up MRI using CS principles. Finally we learn about training HMMs based on some observations.

# A. Compressive Sensing

Let  $x \in \mathbb{R}^N$  and the matrix  $\Psi = [\psi_1, \psi_2, ..., \psi_N]$  be a basis for  $\mathbb{R}^N$ . We say that x is K-sparese if we have:

$$x = \sum_{i=1}^{K} \theta_i \Psi_i \tag{1}$$

and  $K \ll N^2$ . Consider also an  $M \times N$  measurement  $matrix \Phi, M \ll N$  where the rows of  $\Phi$  are incoherent with the columns of  $\Psi$ . Incoherenency between rows of measurement matrix and basis vectors means that we need all of the vectors in second set to expand each of the vectors in first set and vice versa. Compressive Sensing theory, states that only cK incoherent  $measurements: y = \Phi x$  are required to reconstruct signal x, with high probability. It is important to know that the c ratio is always bigger than 1 and in the most of scientific applications is a number between 3 and 4. Incoherency is an essential condition for this theory to be successful. For example, if the rows of the measurement matrix be i.i.d. Gaussians random vectors, such a matrix is incoherent with any other fixed matrix, with high probability.

The problem of using CS in MRI is exactly lying here, because random sampling of k-t space can be more time-consuming than complete sampling. Now, we want to review the proposed methods dealing with CS in MRI.

#### B. Compressive Sensing in MRI

In this part we consider the effect of incoherency in reconstruction of undersampled MR images.

In [9], some of the k-t space lines along the phase coding have been skipped to decrease data acquisition time. To this end, the authors have been suggested using constrained random sampling pattern, as follows:

Sampling points on a regular undersampling pattern are randomly shifted along the phase encoding direction by 1, 0 or +1 with same probability.

Their proposed method has been yielded to better results in comparison with pure random selecting of some k-t space lines along the phase-encoding direction and particularly proposed for dynamic MRI. As we know, dynamic MR images are naturally sparse in temporal frequency domain. Thus we don't need to any sparsifying transform if we reconstruct desired images in temporal frequency domain. Need to high SNR data and time consuming calculations are the drawbacks of their method.

In [10], at first, a relation has been introduced to measure the coherency between sampling pattern and the basis of sparsifying transform. Then the incoherency of the common sampling patterns has been evaluated by this relation. The author has shown that the pure random sampling has the best result and random selecting of k-space lines has the worst record among the others. After that a general relation, has been proposed for non-linear reconstruction of undersampled MRI data based on CS concepts. We describe this relation further, in the third section of the paper.

#### C. Hidden Markov Models

Markov chains [11] are mathematical descriptions of Markov models with a discrete set of states. Markov chains are characterized by:

- A set of states  $\{1, 2, ..., M\}$ ,
- An  $M \times M$  transition matrix (T) whose (i,j) entry is the probability of a transition from state i to state j,
- A set of possible outputs or emissions,  $\{s_1, s_2, ..., s_N\}$ ,
- An  $M \times N$  emission matrix (E) whose (i,k) entry gives the probability of emitting symbol  $s_k$  given that the model is in state i.

A hidden Markov model is one in which we observe a sequence of emissions, but do not know the sequence of states the model went through to generate the emissions.

In this paper, we use the Baum-Welch algorithm [11] to estimate the HMM parameters (T and E) for time variations of k-space lines, after fuzzy clustering.

### III. PROPOSED METHOD

It was mentioned that speed up MRI by undersampling k-t space, contributes to decrease in SNR of reconstructed

images. On the other hand, the reconstruction step in CS is robust to noise. Further more, MR images may have sparse or compressible presentations in appropriate transform domains, such as Wavelet. Also, use of CS in MRI does not need any special hardware changes. Therefore, in this research we focus on increasing MRI speed by the idea of CS. We limit the proposed method, to Cartesian sampling pattern in order to simplicity and applicability. Different steps of the proposed method, are described below:

- 1. Preprocessing: Detaching of under study tissue from available images and making k-t space corresponding to each array coil channels.
- 2. Feature extraction of each k-t space lines along the phase-encoding direction. These features are: average, standard deviation, median and maximum.
- 3. Clustering the time samples of each k-space lines (using the features described above and for each channel of array coil) by fuzzy c-means algorithm. The number of clusters is supposed  $\frac{1}{4}$  of time samples.
- 4. Training of Hidden Markov model for each k-space lines using Baum-Welch algorithm. The results of above clustering (for each channel) are considered as training sequences for the HMM describing time behavior of each k-space line.
- 5. Using the transition matrix came from above, to decide about selecting or rejecting a k-space line to sample. We suppose that the lines which have more non-diagonal transition matrix have more information in them. Thus they are more adequate to be sampled.
- 6. According to [10], image reconstruction by the following constrained L1 norm minimization:

$$\begin{cases}
min. \|\Psi m\|_1 + \lambda TV(m) \\
s.t. \|F_u m - y\|_2 \le \epsilon
\end{cases}$$
(2)

It is done by the means of conjugate gradient and use of available codes in (www.stanford.edu/~mlustig). It should be noted that in (2), m denotes the desired image,  $\Psi$  is the sparsifying transform,  $F_u$  is the undersampled Fourier operator, y is the collected samples of k-space and  $\epsilon$  shows the strength of noise. A total variation term has been added to the object of minimization to yield better result. The first row in (2) shows the sparsity of the desired image and second row guarantees the consistency of collected data and final image. In the following section we present the results of above method.

#### IV. RESULTS

The mentioned approach was applied to two Cardiac Cine MRI data sets, came from 8-channel coil. The results of fuzzy clustering and training of HMMs for 4 k-space lines are presented in Fig.1 and Fig.2 respectively. It must be noted that the k-space lines which have been finally selected to sample, have more non-diagonal transition matrices, as shown in Fig.2. The original (desired) cardiac image, result of zero padding and inverse Fourier transform, reconstructed image by randomly selected k-space lines and reconstructed image by proposed sampling mask, are shown in Fig.3. The image

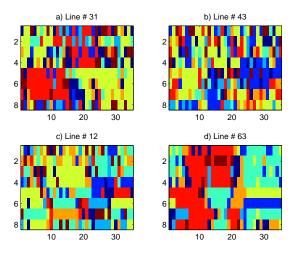


Fig. 1. Clustering Results for different k-space lines. The vertical axis refers to 8 coil channels and the horizontal one shows 35 time samples, in each image.

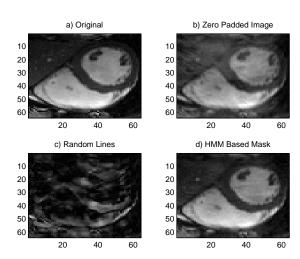


Fig. 3. a) desired image, b) result of zero-padding and inverse Fourier transform, c) result image by random selecting some k-space lines, d) result image by selecting some k-space lines according the proposed method.

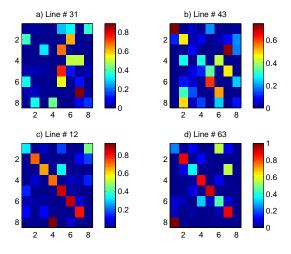


Fig. 2. The result transition matrices for different k-space lines, by Baum-Welch algorithm. a) and b) refer to finally selected lines to sample and c) and d) refer to non-selected ones.

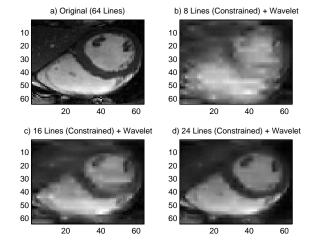


Fig. 4. a) desired image, b), C) and d) result images for different reduction factors, using wavelet as the sparsifying transform and selecting some k-space lines according the proposed method (constrained).

related to random mask has poor quality in comparison with the result of proposed method.

The reduction factor for these images is 2 and we have not used any sparsifying transform. For higher undersampling (speed up) factors, we have to use sparsifying transform.

Fig.4 shows original image and some undersampled Cardiac images. They have been reconstructed by Wavelet (Daubechies-4) as the sparsifying transform and sampling pattern according to proposed method. We have added selecting the central k-space lines along the phase encoding direction to our method, as a constraint.

## V. DISCUSSION AND CONCLUSION

In this article, we increased Cardiac Cine MRI speed by undersampling k-t space. To this goal, we studied time behavior of k-space line along phase-encoding direction by HMM and selected which have more time variations. We used CS principles to reconstruct desired images. We successfully applied our method to reconstruct half sampled Cardiac Cine MRI, without any sparsifying transform. It is possible to earn higher speed up factor with using sparsifying transform and adding the central k-space lines to the selected lines by our method.

Use of fuzzy clustering as an intermediate tool to study complicated phenomena by HMM, applicability to non-dynamic MRI data and simplicity can be accounted as the specifications of our method.

It must be noted that we supposed to have access to all of the k-t space data to develop our algorithm. But it is not possible in the reality and we want to know the most important part of

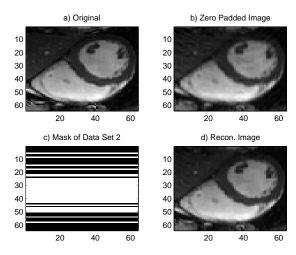


Fig. 5. a) desired image, b) result of zero-padding and inverse Fourier Transform, c) Extracted mask of data set 2 by the proposed method, d) reconstructed image by selecting some k-space lines according the shown mask.

k-t space data to reconstruct acceptable images. On the other hand, we can use the result pattern for a person or group of persons for the others by assuming similarity of MRI data of an organ in different reasons. Fig.5 shows the reconstructed image using the k-space of data set 1 and sampling pattern extracted from data set 2.

In the proposed method, we didn't use the correlation between 2 adjacent k-space lines to decide about selecting them or not. But it may possible to use Distributed Compressive Sensing (DCS) [12] to enter this correlation and get better results.

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