Towards Computationally Efficient Autocalibration for Accelerated MRI using Compressed Sensing Parallel Imaging

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Introduction:
A number of highly accelerated data acquisition methods for 3D volumetric MRI using both Compressed Sensing and parallel imaging have been developed. Among them L1-ESPIRiT[1], and an efficient version of it called ESPIRiT[2] have been proposed in the recent past. ESPIRiT addresses the computational challenges with the use of an image-domain Parallel imaging (PI) operator based on sensitivity maps obtained from a SPIRiT [4] convolution kernel, in place of performing computationally expensive convolution with the kernel in k-space in L1-ESPIRiT. But the computation of the SPIRiT kernel weights itself is still computationally expensive, especially for high channel count reconstructions. The weights computed by obtaining a least squares fit for predicting target points in the calibration region using a set of source points in their neighborhood. The computational complexity for generating the kernel weights is dependent on the number of source points and the number of target points. In the past, some approaches for optimal kernel support were explored in the context of GRAPPA [5]. This work is aimed at developing an optimally shaped neighborhood that reduces the number of source points without sacrificing image quality, but has much improved computational performance.

Theory:
The autocalibration procedure for ESPIRiT generates reconstruction weights \( w \) by fitting a source matrix \( A \) to a target matrix \( B \) using generalized matrix inversion. Number of rows \( m \) of \( A \), are the number of points in the calibration region. Conventional ESPIRiT calibration uses all source points in a cube-shaped k-space kernel. For a \( N \)-channel dataset with a kernel size of \( k_1 \times k_2 \times k_3 \), the number of columns \( n \) of \( k_1 \times k_2 \times k_3 \times N \). The weights for each coil are obtained by solving Moore-Penrose pseudo inverse [2] i.e., solve for \( C=A^T \hat{A}^{-1} \hat{D} = A^T \hat{B} \) Coefficient weights diminish rapidly, as k-space distance increases as shown in Figure 1(a). Based on this observation, the significance of source points formed by \( k_1 \times k_2 \times k_3 \) neighborhood around each target pixel was studied by experimenting with different neighborhoods defined by the Lp-norm of the target point. If \( X_i=(x_1,y_1,z_1) \), \( X_j=(x_2,y_2,z_2) \) the \( L_p \) distance \( d_p(X_i,X_j) \) between \( X_i \) and \( X_j \) is defined as \((x_1-x_2)^p + (y_1-y_2)^p + (z_1-z_2)^p \) \( L_p \)-neighborhood of a point \( X \) is defined as \( Y \) \( L_p \), \( Y_{L_p}(X) \). Different neighborhoods for \( p=1, 2, \infty \) would take different shapes as shown in figure 1(a). For a typical kernel of radius \( r=3 \), an \( L_\infty \) kernel has 343 neighbors. The number of columns \( n \) of matrix \( A \) goes down by a factor of 5.4 for an \( L_1 \)-neighborhood and by 2.78 for an \( L_2 \)-neighborhood. Since the solver is of the order of \( n^3 \), this gives an overall speed-up of 150x for an \( L_1 \) kernel and 21x for \( L_2 \) kernel over a typical box shaped \( L_\infty \) kernel. The correlation computation time for computing the correlation matrix \( C \) scales down linearly.

Methods:
Based on the above theory, the calibration step mainly consists of the following operations: 1) Forming the \( A \) matrix, based on the neighborhood shape. 2) Finding the correlation matrices \( C=A^T \hat{A}, D=A^T \hat{B} \) 3) Solving for the kernel weight \( C=A^T \hat{B} \) 4) Updating \( D \). We exploit the redundancy and directly compute the correlation matrix \( \hat{C} \) as proposed in [3]. We experimented the method described above for \( L_p \) neighborhoods, with \( p=0.1 \) to 1 in steps of 0.1, for \( p=2 \) and \( p=\infty \) on 4 patient datasets (T2-weighted brain MRI using 8-channels and 32 channels, Proton-density-weighted Knee MRI, noncontrast-enhanced renal MRA).

Results:
While we observed some difference in the image quality for \( p=1 \), we found that for \( p=1, p=2 \), \( p=\infty \) there was no visible difference in image quality. The ESPIRiT reconstruction outputs using by L1,L2,L_\infty norms for calibration weights computation are shown in figure 1(d). This is confirmed by the fit we observe between source and target points for these neighborhoods and the coil-maps generated as shown in figure 1(c). We implemented this approach on a dual Intel Xeon 55xx. With our multi-threaded implementation using 8-threads, for a 32-channel dataset, and a calibration region of size \( 27 \times 27 \times 128 \), the compute time for the solver is 720 seconds for \( L_\infty \) kernel, 5 seconds for \( L_1 \) kernel, and 35 seconds for \( L_2 \) kernel.

Discussion:
Using an optimal neighborhood for calculating kernel weights for calibration in ESPIRiT, we can significantly improve the computational efficiency. This study demonstrates that we can achieve this, ensuring that we are not compromising on the fit between the source and target points from the calibration region and more importantly the perceptual image quality.

References:

Figure 1: T2-weighted 32 channel Brain MRI (a) Plot of weights for a typical 7x7 neighborhood (b) Neighborhood shapes (c) Coil Maps (d) Final Output (e) difference Images with respect to fully sampled for L1 L2 and \( L_\infty \) norms

Figure 2: PDW- 8-channel Knee MRI: Final Output for L1,L2 and \( L_\infty \) norms