Simulating single-coil MRI from the responses of multiple coils

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Abstract
We convert the information-rich measurements of parallel and phased-array MRI into noisier data that a corresponding single-coil scanner could have taken. Specifically, we replace the responses from multiple receivers with a linear combination that emulates the response from only a single, aggregate receiver, replete with the low signal-to-noise ratio and phase problems of any single one of the original receivers (combining several receivers is necessary, however, since the original receivers usually have limited spatial sensitivity). This enables experimentation in the simpler context of a single-coil scanner prior to development of algorithms for the full complexity of multiple receiver coils.

1 Introduction
Typical modern MRI scanners (such as all those represented in the fastMRI data of [13]) include multiple receivers — so-called “coils” of conductors — for making measurements. In this brief yet self-contained technical note, we seek to conveniently simulate the response of a single-coil machine, complete with errors that are reasonably representative of the measurement errors which a real scanner would make, given only the responses from a multi-coil, “parallel-imaging” machine. The fastMRI competition issues two challenges — single-coil and multi-coil — under the expectation that the single-coil track could provide a simplified stepping-stone toward the full multi-coil challenge.

The aim of fastMRI is compressed sensing — the accelerated acquisition of image reconstructions by taking fewer measurements than classical signal processing would need to attain the same resolution; compressed sensing yields full-resolution or “superresolved” reconstructions based on so-called “undersampled” measurements; measurements acquired at lower than the Nyquist rate required to reconstruct any arbitrary signal for a given bandwidth. Two approaches to compressed sensing are especially popular: (1) optimization for individual images and (2) machine learning over collections of images. Combinations of the two approaches are also possible. Techniques representative of the first approach (via optimization) include Fourier reconstruction with total-variation regularizers (or related regularizers, such as the sum of the absolute values of wavelet coefficients); representative of the second approach (via machine learning) is statistical modeling with deep neural networks. The terminology, “compressed sensing,” “compressive sensing,” and “compressive sampling,” is due to [1] and [2]; they took the first approach (via optimization). The second approach (machine learning) is the focus of [13].

Complicating the development of compressed sensing for MRI is the longstanding focus on leveraging multiple receiver coils in the design of MRI scanners, as reviewed by [7]. All scanners of [13] are multi-coil by default, designed and optimized for multi-coil measurements. However, compressed sensing is more complex with multi-coil data. Distilling the multi-coil challenge into a single-coil stepping-stone could lower barriers to entry into research on compressed sensing for MRI and accelerate the development of accelerated MRI acquisitions, by teasing apart the fundamental mathematical issues from the multi-coil complications.

Thus, we need to simulate a single-coil scanner with sufficient fidelity to actual errors in measurement, using only multi-coil data. As hoped (and illustrated with numerical examples in Section 3 below), our single-coil emulation does indeed yield phase similar to the notorious phase problems tackled by [5] and others. Encouragingly, the method of [9] is similar to ours and yielded highly successful results in a related setting (though, admittedly, our mathematical motivation, practical application, optimization procedure and targets, and detailed objective differ). Section 2 now introduces our simulation of a single-coil scanner.

∗ “MRI” abbreviates “magnetic resonance imaging”
Figure 1: The wide solid lines represent coils of conductors — 16 small ones and 1 large one. The arrows indicate circulation of current through the conductors (with solid lines for the small coils and dashed lines for the large coil). The currents in the small coils would sum to induce the indicated current in the large coil (flux through the small coils would be equivalent to flux through the large coil), neglecting mutual inductances — and actual designs of coil arrays tend to minimize mutual inductances (as per [11]).

2 Emulated single-coil

We propose using a linear combination of the responses from multiple coils for the emulated single-coil (ESC) response. We least-squares fit the complex-valued coefficients in the linear combination to the “ground-truth” reconstruction, estimating the ground truth using the canonical full multi-coil reconstruction, the root-sum-square (RSS) of [8]. Most notably, linearly combining the raw coil responses sums complex-valued fluxes directly, rather than summing nonnegative energies as in the RSS. This simulates a single-coil scanner for the reasons discussed next.

2.1 Motivation

As seen in Figure 1, the flux through a single coil is physically the sum of the flux through multiple coils forming a disjoint partition of the single coil that loops around the multiple coils (provided that the multiple coils are all in phase and have the same gain, while neglecting the mutual inductance that coil design typically tries to minimize, as per [11]). This is much like in the proof of Stokes’ Theorem (which also uses Figure 1). Since we do not know the correct phase offset and gain calibrations, which amount to multiplying a whole coil’s measurements by the same complex number, we can fit the complex number so as to match the ground-truth RSS reconstruction as well as possible.

As described, for example, by [11], many MRI machines use “bird-cage” or other volumetric coils rather than surface coils in a flat plane. A phased “single-coil” array is often a cylindrical array of coils with capacitative (or inductive) coupling between the coils. This coupling effectively introduces phase offsets between the multiple conductors forming the “single-coil” bird-cage. Our linear combination accounts for phase offsets and is physically realizable and similar to bird cages used in practice. The use of coils selecting for circular polarization (including the quadrature pairs, butterflies, figure eights, etc. described by [11]) is an additional complication, implicitly resolved by how the least-squares fit of the linear combination to the ground truth decides to combine the coils. The following subsection formalizes the mathematical details.
2.2 Mathematical formulation

Given data $A$ from multiple coils, we define $k$ to be the number of coils, $l$ to be the number of slices in a volume of cross-sectional images, $m$ to be the height of each slice, and $n$ to be the width of each slice: $A$ then consists of an $(nm 	imes k)$ matrix with complex-valued entries. More precisely, the images in $A$ are cropped inverse Fourier transforms of the original two-dimensional Fourier domain (“k-space”) measurements, cropping to the center $m \times n$ block of pixels. We define $b$ to be the $(nm 	imes 1)$ column vector containing “ground-truth” reconstructions from the full multi-coil data, such as the RSS of [8]; specifically, each entry of $b$ is the Euclidean norm of the corresponding $1 \times k$ row in $A$. We then calculate the $k \times 1$ column vector $x$ with complex-valued entries minimizing

$$\| \sqrt{|Ax|} - \sqrt{|b|} \|_2^2,$$

where $\| \cdot \|_2$ is the Euclidean norm, $\sqrt{\cdot}$ takes the square root entrywise, and $| \cdot |$ takes the absolute value entrywise, so that $|b|$ is in fact equal to $b$, as the entries of $b$ are nonnegative. The distance $\| \sqrt{|Ax|} - \sqrt{|b|} \|_2$ is known as the Hellinger metric of [3]; the square roots amplify entries of small magnitude and attenuate entries of large magnitude. Furthermore, $\sqrt{|Ax|} - \sqrt{|b|} = (|Ax| - |b|)/(\sqrt{|Ax|} + \sqrt{|b|})$, so the Hellinger distance is more like a measure of relative errors than the conventional Euclidean distance $\| |Ax| - |b| \|_2$. Our estimate of $|b|$ is $|Ax|$, where $x$ minimizes (1). Minimizing (1) is far from the only possibility; minimizing (1) performed well in our numerical experiments and facilitates the nonlinear optimization required.

To solve the nonlinear least-squares problem minimizing (1), we tried three different methods. The three methods (all reviewed, for example, by [6]) are {1} gradient descent, {2} the Levenberg-Marquardt variant of Gauss-Newton iterations, and {3} the LBFGS (limited-memory Broyden-Fletcher-Goldfarb-Shanno) version of quasi-Newton iterations. For all, we started the iterations with the solution to the linear least-squares problem minimizing

$$\| Ax - b \|_2^2;$$

the objective function in (1) is the same as in (2) aside from taking the square roots of the absolute values of the entries of $Ax$ and of $b$.

Gradient descent was the slowest, and its best step size varied dramatically. Levenberg-Marquardt and LBFGS both worked well, but LBFGS was an order of magnitude faster on the wall clock. Levenberg-Marquardt requires forming a large system matrix explicitly, whereas LBFGS uses only matrix-vector multiplications and so can be “matrix-free,” not requiring the explicit formation of the system matrix. The results reported below come from LBFGS. We used the implementations in SciPy of [4] for all experiments.

Remark 1. Anuroop Sriram points out that the mean of the noise on the “ground-truth” RSS will be strictly positive in regions of the image where the actual object being imaged is absent, as the RSS is always nonnegative. Thus, when we fit the linear combination of the coil responses, we should try to match the RSS only where the RSS is substantially above a threshold that is significantly above the noise level. For the current data from fastMRI of [13], essentially the entire cropped image of interest meets this criterion, so we try to match the full cropped image.

3 Numerical examples

We compare the emulated single-coil (ESC) method of Section 2 with ESPiRiT of [10] and the leading singular vector of the singular value decomposition (SVD) of $A$ from (1), as in the “eigencoil” or “coil compression” reviewed by [7]. ESC could be regarded as a special kind of the “virtual-coil” methods discussed by [7]. We also display the root-sum-square (RSS) of [8] for reference as our best estimate of the ground truth.

For our data — scans of 1594 knees from [13] — the parameters from Section 2 take the values $k = 15$ receiver coils, $m = 320$ rows and $n = 320$ columns in a cross-sectional image, with Figure 2 specifying $l$, the number of cross-sectional slices. We used a memory of 10 vectors in the limited-memory BFGS quasi-Newton iterations to minimize (1) starting from the linear least-squares solution minimizing (2).

In Figures 3-8, the first row shows (the magnitude of) the reconstructions. The second row shows the difference between the RSS ground truth and the magnitude of each method’s reconstruction. The third row shows the phase of the reconstructions (estimates of the RSS ground truth discard the phase). Figures 9–14 display the entries of the solution $x$ minimizing (1), where the ESC reconstruction is $Ax$. 
The method proposed in the present paper is heavily data-driven. If computational simulations or excellent empirical estimates of the “physical coil sensitivity maps” defined in [7] could be calculated with high fidelity to the physical reality represented in the fastMRI data of [13], then less data-intensive methods might be feasible. Unfortunately, actual MRI scanners are exceptionally complex and deviate significantly from textbook idealizations; interactions of the objects being imaged with the scanners further distort the sensitivities of receiver coils, as detailed, for example, by [5]. While autocalibration procedures such as those from ESPiRiT of [10] work well for multi-coil reconstruction, their estimates of sensitivity maps are specifically tailored for multi-coil reconstruction, not for constructing single-coil measurements from multi-coil.
Figure 3: first randomly chosen example; “abs” refers to the absolute value of a complex number and “angle” refers to the phase.
Figure 4: second randomly chosen example; “abs” refers to the absolute value of a complex number and “angle” refers to the phase
Figure 5: third randomly chosen example; “abs” refers to the absolute value of a complex number and “angle” refers to the phase.
Figure 6: fourth randomly chosen example; “abs” refers to the absolute value of a complex number and “angle” refers to the phase.
Figure 7: fifth randomly chosen example; “abs” refers to the absolute value of a complex number and “angle” refers to the phase
Figure 8: sixth randomly chosen example; “abs” refers to the absolute value of a complex number and “angle” refers to the phase
Figure 9: entries of $x$ minimizing (1) corresponding to the data in Figure 3

Figure 10: entries of $x$ minimizing (1) corresponding to the data in Figure 4

Figure 11: entries of $x$ minimizing (1) corresponding to the data in Figure 5
Figure 12: entries of $x$ minimizing (1) corresponding to the data in Figure 6

Figure 13: entries of $x$ minimizing (1) corresponding to the data in Figure 7

Figure 14: entries of $x$ minimizing (1) corresponding to the data in Figure 8
4 Conclusion

Although our simulation of a single-coil scanner might appear perverse and pointless at first glance — after all, we throw away the information necessary to realize the gains in signal-to-noise ratios that imaging with multiple receivers provides — maintaining fidelity to a physically realizable simple MRI machine in fact yields a convenient test case, a reliable stepping-stone toward the complexity of full parallel-MRI reconstruction. Less than 0.5% of the resulting emulated single-coil (ESC) images exhibit artifacts such as those displayed in Figures 15–17; Figures 3–8 display typical examples. The obvious alternative to our scheme would be to take measurements on an actually manufactured single-coil MRI scanner; however, MRI is extremely expensive, especially in the medical domain considered for the fastMRI data set of [13].
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