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TITLE: Magnetic Resonance Arterial Spin Tagging for Noninvasive Pharmacokinetic Analysis of Breast Cancer

PRINCIPAL INVESTIGATOR: Michael H. Buonocore, M.D.

CONTRACTING ORGANIZATION: University of California
Davis, California 95616-8671

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<td>This project is designed to develop and evaluate arterial spin tagging techniques for the non-invasive measurement of breast tissue perfusion. The long term goal is to show that arterial spin tagging can provide comparable tissue information as that currently obtained using dynamic first-pass contrast-enhanced imaging. To date, the research effort has focused on the pulse sequence, protocol and image processing software needed for analysis and visualization of the anatomic, functional, and dynamic breast images. Specific tests and analyses of the sequence have been performed to understand and correct for driven equilibrium effects, and for inversion and excitation slice profile mismatch. Software development includes methods for estimating, at each pixel, the $T_1$, the perfusion (as defined by $f/A$), and standard errors, and from these compute a “suspicion index” that the tissue is abnormal. Algorithms for image registration and image display have been developed to allow direct comparison of high resolution $T_1$, fast spin echo $T_2$ and proton density, perfusion, and contrast enhanced images. The research effort provides crucial software for acquisition and definitive review of the images, and automation for detailed and thorough statistical analysis of the very large data sets that are obtained.</td>
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(5) INTRODUCTION

Subject: This study concerns the development of MRI arterial spin tagging to non-invasively measure breast tissue perfusion. The project will test whether the tagging technique is a low cost and more reliable alternative to dynamic first-pass contrast-enhanced imaging, giving possibilities for use in specific clinical settings, as well as in screening. MR dynamic first-pass contrast-enhanced imaging has shown that malignant and benign breast lesions can be distinguished. However, it may have limited importance in clinical breast diagnosis due to significant false-negatives and false-positives. Also, dynamic contrast enhancement cannot be used in screening, even within defined high-risk sub-populations such as patients with dense breasts, due to the high cost of contrast material and administration. The arterial spin tagging technique was developed to measure tissue perfusion parameters without the use of contrast, and has been successfully demonstrated in brain and kidney. The spin tagging technique may provide more sensitive and specific assessment of these parameters, at higher spatial resolutions, by virtue of its ability to acquire repeated measurement of flow sensitive data to improve signal to noise ratio (SNR). Dynamic first-pass contrast-enhanced imaging has a limited data acquisition time and thus a limited resolution and SNR.

Purpose: The purpose (long-term goal) of the project is to develop MRI arterial spin tagging as a low cost and more reliable alternative to first-pass contrast-enhanced imaging. We will test the hypothesis that arterial spin tagging provides accurate and precise discrimination between normal tissue, benign and malignant lesions, when differences in perfusion and $T_1$ exist. Lesions will have been previously detected by clinically accepted diagnostic imaging procedures, and by biopsy. Statistical analysis will be performed to access the correspondence between arterial spin-tagging and biopsy, and to establish the relative value of spin tagging compared to first-pass contrast-enhanced MRI. We hope to establish that, relative to using first-pass contrast-enhanced imaging, false positives and negatives are reduced using arterial spin tagging by virtue of increased image signal-to-noise ratio (SNR), higher spatial resolution, and the unique ability to obtain estimates of macromolecular bound fluid fractions.

Scope: The effort on the project is mainly limited to the technical aspects of development of a new methodology. However, the project also includes a rigorous performance comparison with the current gold-standard methodology. The specific aims are to (1) refine arterial spin tagging pulse sequences and imaging protocols, (2) develop automated data analysis software for measurement of breast tissue parameters, and (3) compare the technique to first-pass contrast-enhanced MRI and biopsy. The technique will be evaluated in sixty patients, with roughly equal numbers of benign and malignant lesions. Multi-component differential equation models of magnetization distribution within breast tissue will be used to estimate perfusion and $T_1$, from which regions suspicious for malignant disease will be identified. While sub-population screening is a critical potential application, it is premature to propose a study to evaluate the technique for specific sub-population screening, before it’s capability for detection and discrimination of previously detected lesions is established.

Background: Studies have shown that dynamic first-pass contrast-enhanced magnetic MRI discriminates malignant from many types of benign breast disease. This technique detects malignant disease by measuring tissue fluid distribution parameters, which are derived from the rate of rise of MRI signal intensity. The rate of decrease of signal intensity, after the peak signal is reached, is also believed to be related to malignancy, based on the idea that the vasculature of
malignant tissues possesses a high degree of arteriovenous shunting causing more rapid elimination of contrast agent. Unfortunately, this technique may have limited importance due to significant false-negatives and false-positives. Further development and refinement is required, but major improvements are unlikely given the limited data acquisition time inherent to first-pass techniques. In particular, the first-pass techniques cannot provide, with high temporal resolution, high spatial resolution required to reveal morphological details that can reduce false positives. Arterial spin tagging has been developed for measuring tissue parameters without the use of contrast, and has been successfully demonstrated in brain and kidney. Spin tagging measures absolute blood perfusion (cc/min/g tissue), longitudinal relaxation time ($T_1$) of the tissue water, and possibly macromolecular bound fluid fractions. Spin tagging can provide more sensitive and specific assessment of tissue parameters, at higher spatial resolution, using the ability to build up signal to noise over reasonable scan times to investigate tissue parameters. It is not limited to data collection during a first pass.

Previous studies have confirmed that first-pass enhancement of carcinomas by Gd-DTPA are often (but not always) more intense and faster compared to normal parenchyma or benign breast disease [1,2,3,4,5,6,7,8]. Enhancement is attributed to tumor-induced angiogenesis resulting in increased vascularity within the malignant tumor [9,10,11] which often also develops anastomoses and shunts. Due to greater tissue uptake, carcinomas enhance maximally within 1.5 to 2 minutes of contrast injection and then, due to accumulation in extracellular fluid, tend to maintain the enhancement for several minutes. Kaiser [12] postulated that false positive benign lesions may be distinguished by the behavior of signal enhancement after the peak occurs, i.e., during the elimination phase [13]. Malignant lesions have greater degrees of A/V shunting, and hence more rapid elimination of contrast and more rapid return to non-enhanced signal levels. Due to the importance of the time course of signal enhancement in these criteria, pharmacokinetic analysis [8,14,15] has been clinically important. Motion correction [16] and image subtraction [17] have been important for reliable temporal processing. However, the importance of morphology is regarded by many researchers as equally or more important than temporal profiles, and some have emphasized high spatial resolution, not temporal resolution, in discrimination of tumors [18,19 page 595]. Overall, inconsistent reports and general overlap of benign and malignant temporal profiles precludes the role of MRI as an “imaging biopsy” [19,20].

Techniques for non-invasive, spatially resolved MRI perfusion measurement have been developed based on magnetic labeling of arterial water [21,22] (i.e. arterial spin-tagging), and tested in both animals and humans [23,24,25,26]. Mathematical models of the magnetization distribution are used to relate changing signal intensity to tissue perfusion (flow of magnetization into tissue), magnetization of free fluid, magnetization of fluid bound to macromolecules, and fluid exchange rates between these compartments [23,27]. Tagged spins will not accumulate significantly in tissues as does contrast agent, and therefore do not obtain precisely the same measurements as contrast-enhanced scans. However, arterial spin tagging will measure macromolecular bound fluid fraction, that is affected by extracellular fluid fraction, and can be adapted to measurement of fluid elimination.
(6) BODY

Overview

This section details the experimental methods, results and discussion in relation to the Statement of Work outlined in the proposal. The first subsection provides brief paragraphs of the accomplishments in relation to the Statement of Work. Detailed discussion of these accomplishments is provided in the second subsection. There are two main contributions of the work. The first is the discovery of a multitude of unrecognized sources of $T_1$ estimate errors, the recognition of the profound sensitivity of perfusion measurements to $T_1$ estimate errors, and the development of robust pulse sequences and analysis methods to derive accurate $T_1$ estimates with inherently noisy and confounded signal data. The second is the development of the software program, BrView, to allow rapid review, quantitative analysis, and assessment of the multitude of different breast images and timeseries data that is obtained in each patient study.

Summary of accomplishments in relation to Statement of Work

Technical objective 1

Task 1: Months 1-6: Implementation and testing of magnetization transfer pulses for both arterial tagging and first-pass contrast enhanced sequences.

Magnetization transfer pulses have not been implemented yet. Mainly, this delay was due to the fact that the MRI system at UC Davis Medical Center is scheduled to be upgraded to the new Horizon LX system soon, and we opted to defer this pulse sequence development until we were using this new LX software platform. There have been multiple delays, but this upgrade is scheduled to occur by January 1999. The existing pulse sequence will be converted from the 5.4 OS platform to the LX2 platform (a substantial change), and the magnetization transfer pulses will be implemented also.

Because of concerns regarding the ultimate utility of the magnetization transfer technique, we have decided that implementation and testing of arterial spin-tagging sequence based on echo planar imaging (EPI) data acquisition is a higher priority. EPI based spin-tagging technique can theoretically improve the accuracy of the $T_1$ estimation, which we have studied and worked with extensively (see below). The major problem with magnetization transfer is that it represents a perturbation on an already small signal. Therefore, whether we will observe a magnetization transfer effect is questionable. Nevertheless, it will be implemented as part of the project.

Task 2: Months 1-9: Implementation and testing of interleaved high-resolution imaging technique, for both arterial tagging and first-pass contrast enhanced sequences.

An interleaved arterial spin-tagging technique, that allows high resolution (e.g. 256 x 256 or more) has been implemented and tested. We also implemented a rectangular data acquisition technique, which acquires data sets that have different numbers of points along the frequency and phase encode direction. The 256 x 240 interleaved acquisition provides the best spatial versus temporal resolution tradeoff, and has been used for all of our recent studies.

Task 3: Months 1-9: Analysis of spin tagging sequence to understand causes of existing baseline offsets, effects of inversion slice transition profiles, and effects of RF flip angle profile on the measurements, with implementation and testing pulse sequence modifications to minimize these imperfections.
Matching of inversion and excitation slice profiles was greatly improved by using customized RF pulses designed with the Shinnar-Le Roux (SLR) algorithm. The new data processing technique (based on a semi-log linear regression of inversion time ($T_I$) dependent signal) significantly reduced the previously reported problem in measuring the longitudinal relaxation time ($T_1$) caused by baseline offsets, and by the uncertainty in the effective inversion time. The optimal RF flip angle for spin excitation during spatial encoding was found to be 10 degrees, based on experiments over a range of flip angles.

**Technical objective 2**

**Task 4: Months 3-15:** Software for automatic estimation and error analysis of perfusion, tissue water longitudinal relaxation time, and extracellular fluid volume fraction from mathematical models and user-defined ROIs from spin tagging timeseries. Implementation of pharmacokinetic model calculations and error analysis for first-pass contrast enhanced imaging.

A software program (BrView) has been written in C, X Window System, and Motif, and implemented on an SGI O2 computer (paid for by grant funds) for analyzing and visualizing the MR images and timeseries. Much effort has been expended on developing the capability to easily review all of the images taken in each study, and cross-reference pixel locations. Much more effort than anticipated has been expended in the development of a robust technique for identifying so-called "suspicious regions" based on the $T_1$ and perfusion measurements. The measurement of $T_1$, upon which the perfusion measurement entirely depends, is based on a semi-log linear regression technique developed by the investigators. The $T_1$, perfusion ($f/A$) and standard errors of these quantities are estimated automatically using this robust technique. The so-called feature images display these quantities in color and grayscale. These quantities are used to calculate a “suspicion index” for each pixel, and thereby identify regions of breast tissue suspicious for malignancy.

The implementation of the pharmacokinetic model calculations and error analysis for first-pass contrast enhanced imaging has not been completed. Thus far, the software allows the user to click on the reference image of the dynamic study, to display the time profile of the signal at that pixel. The implementation of the pharmacokinetic model is currently a major focus of the current effort.

**Task 5: Months 3-15:** Software for registration (including implementation and testing of motion correction and physiological noise reduction algorithms) and overlay of images from high resolution $T_1$ weighted, spin-tagging, and contrast enhanced studies.

Using the BrView software program, image pixels containing high values of perfusion ($f/A$), and moderate to high values of $T_1$, and low standard errors, are identified as suspicious, and assigned a “suspicion index” based on finding similar abnormal values at spatially adjacent pixels. Suspicious pixels can be overlaid on the high-resolution $T_1$ and $T_2$ clinical images by color mapping, and simultaneously presented with the first-pass contrast enhanced timeseries at these pixels. This provides the user with a comprehensive anatomical and functional view of the suspicious regions, and facilitates making a decision regarding the malignant nature of the lesion. Finally, a motion artifact estimation and correction algorithm has been developed and implemented in a separate software program.
Technical objective 3

Task 6: Months 9-24: 60 patients with malignant and benign breast lesions will be imaged using $T_1$ weighted, first-pass contrast enhanced, and arterial spin tagging MRI pulse sequences.

We have recruited only two subject studies for both first-pass contrast enhanced and arterial spin tagging MRI pulse sequences. Now that the software development is substantially completed, we are trying to improve the rate at which patients are recruited into the study. It is likely that the patient studies will extend into the third year of the proposal. We are currently discussing with physicians at a local Kaiser Hospital for recruitment of additional patients. Appropriate documentation will be submitted for approvals upon reaching a firm agreement.

Detailed description of the accomplishments

Background to theory of arterial spin tagging

This background material is taken from the literature, and represents the starting point for the project. In the so-called “on-slice” spin tagging technique that we implemented, spins in a slab of tissue are inverted first using a RF pulse. The thickness of the slab can be controlled by the strength of the slab selective gradient ( normally a Z gradient). Then the levels of recovery to equilibrium are measured at different times (Figure 1). The rate of recovery (i.e. the effective $T_1$) can be estimated from the recovery curve. In the research protocol, 3mm slices are imaged

![Figure 1](image1.png)

Figure 1. After the Magnetization is inverted, it recovers towards thermal equilibrium position according to characteristic relaxation time $T_1$. The signal intensity along a typical magnetization recovery curve is shown. The image displays the magnitude of the magnetization. The rate of recovery can be determined from images acquired at different times of the recovery curve represented by $T_1$. $T_1$ is a point in time at which an zero k-space line is acquired.

Figure 2. A voxel within the imaged slice is identified. Arterioles and capillaries pass through the voxel resulting in net blood flow (indicated by long arrows) and replacement of tissue water. Blood water (indicated by short arrows) exchanges readily with the extracellular fluid.

sequentially (one at a time). The goal is to measure capillary blood perfusion, which is defined as the net amount of arterial blood per second entering a gram of breast tissue. This definition can be extended to a per unit volume (voxel) measurement, i.e., the net amount of arterial blood per second entering a voxel of breast
tissue. Therefore, perfusion represents the directed microscopic flow within a voxel. In breast tissue, blood vessels, which are basically composed of capillaries, are very small. Blood flow is very slow in these vessels. Water molecules can move easily between the vessels and their surroundings (Figure 2). A two-component model is used to approximate the activity of the water molecules. In this model, water molecules in blood vessels entering a slice of breast tissue mix thoroughly with water molecules outside the vessels before they leave the slice. For this two-component model, the Bloch Equation for the breast tissue water magnetization at the slice of tissue investigated can be written as [23]

\[
\frac{dM(t)}{dt} = \frac{M_0 - M(t)}{T_1} - k_{for} M_m(t) + k_{rev} M_m(t) + f M_a(t) - f \left( \frac{M(t)}{\lambda} \right) \quad (1.1)
\]

where, \( M(t) \) = the longitudinal water magnetization of a voxel of breast tissue at time \( t \), \( M_0 \) = the value of \( M(t) \) under fully relaxed conditions, \( T_1 \) = the spin-lattice relaxation time constant of the breast tissue, \( M_m(t) \) = the longitudinal magnetization of macromolecules in a voxel of tissue, \( k_{for} \), \( k_{rev} \) = the magnetization transfer rate constants from water to macromolecules and vice versa, \( M_a \) = the longitudinal water magnetization per ml of arterial blood entering a voxel from outside of the slice investigated, \( f \) = perfusion in ml of arterial blood per second in a voxel, and \( \lambda \) = the ratio of water content between breast tissue overall and the arterial blood within it.

To simplify Eq.(1.1), we assume that the magnetization transfer contributions are negligible, specifically,

\[ (k_{rev} - k_{for}) M_m(t) \ll \frac{M_0 - M(t)}{T_1} + f M_a(t) - f \left( \frac{M(t)}{\lambda} \right) \]

Then Eq. (1.1) can be simplified to

\[
\frac{dM(t)}{dt} = \frac{M_0 - M(t)}{T_1} + f \frac{M_a(t)}{\lambda} - f \frac{M(t)}{\lambda} \quad (1.2)
\]

Eq. (1.1) will be utilized without this approximation when magnetization transfer is implemented. Parameters \( T_1 \) and \( f / \lambda \) in Eq. (1.2) can be estimated from two spin tagging conditions. In the so-called "Selective Tagging Condition", all spins within the slice of breast tissue being imaged are inverted, and the signal from the recovering magnetization is acquired at a specified later time (defined as the inversion time TI). Images are acquired at several TI values in order to generate a magnetization recovery curve at each pixel. Since the arterial spins entering the slice are not influenced by these selective inversion pulses, they enter the slice with full longitudinal magnetization (i.e. they are in thermal equilibrium). Then,

\[ M_a(t) = M_a^0 \]

where \( M_a^0 \) = the value of \( M_a(t) \) at thermal equilibrium. In the two-component model, we assume that the arterial blood spins entering the slice mix thoroughly with spins in the surrounding breast tissue before leaving the slice. Thus,

\[ f M_a^0 = f \frac{M_0}{\lambda} \]
and Eq. (1.2) becomes
\[
\frac{dM(t)}{dt} = M_0 \left( \frac{1}{T_1} + \frac{f}{\lambda} \right) - M(t) \left( \frac{1}{T_1} + \frac{f}{\lambda} \right)
\] (1.3)

A MR system measures the apparent \( T_1 \), here represented as \( T_{1s} \), as if all spins are stationary (not flowing):
\[
\frac{dM(t)}{dt} = \frac{M_0 - M(t)}{T_{1s}}
\]
where,
\[
\frac{1}{T_{1s}} = b_s = \frac{1}{T_1} + \frac{f}{\lambda}
\] (1.4)

and the subscript \( s \) denotes that the parameter was measured using the selective RF pulse.

Images are also acquired in a so-called "Non-selective Tagging Condition". Here, spins are inverted everywhere in the sensitive volume of the body coil, but signals are acquired only from the slice being investigated, as in the Selective Tagging Condition. The arterial spins flowing into the slice thus experience the same inversion pulses as those within the slice, and the magnetization of entering arterial blood can be written
\[
f M_a(t) = f \frac{M(t)}{\lambda}
\]
Eq. (1.2) reduces to the simple expression
\[
\frac{dM(t)}{dt} = \frac{M_0 - M(t)}{T_1}
\]
where \( T_1 \) in this case represents the true \( T_1 \) of the tissue without the confounding influence of blood flow. This \( T_1 \) and corresponding rate \( b_n \) is denoted
\[
T_{1n} = T_1
\] (1.5)
and
\[
b_n = \frac{1}{T_{1n}}
\] (1.6)

where the subscript \( n \) is used to distinguish these parameters from those found in Selective Tagging Condition. Based on the \( b_s \) and \( b_n \) values obtained from the two tagging conditions, we can determine the perfusion parameter \( f/\lambda \):
\[
\frac{f}{\lambda} = b_s - b_n
\] (1.7)
This $f/\lambda$ is the rate of magnetization replacement in a voxel. The importance of $f/\lambda$ is that, compared to normal breast tissue and benign breast lesions, a higher $f/\lambda$ is expected in malignant tumors, due to higher blood flow rates into the malignant tumor.

Contributions to pulse sequence design

The University of California Davis Medical Center operates a Signa Advantage 1.5 T GE MR system (Milwaukee, Wisconsin) (with 1.0 g/cm peak gradient strength, 500 microsecond rise time, and version 5.4 OS) for MR clinical and basic science research. An arterial spin tagging pulse sequence has been developed for this MR system based on the original source code of the commercial GE 2D fast Spoiled Gradient Recalled Echo (SPGR) pulse sequence. In this sequence, after the inversion RF pulse, raw data is acquired for an entire 2D image using the fast SPGR k-space acquisition (Figure 3). Several innovations have been implemented in this custom pulse sequence: 1. The original 2D fast SPGR data acquisition scheme has been modified to support a symmetric centric $k$ space data acquisition. Data acquisition starts at the zero $ky$ line of the $k$ space, and alternates below and above this $ky$ line, end at the highest frequency $ky$ lines. This allows the low spatial frequency components, which have the dominant effect on image contrast, to be acquired at the beginning. The effective time of inversion, defined by the time from spin inversion to the acquisition of the center $k$ space line, can be adjusted most flexibly with this acquisition. 2. Implementation of an optimized RF inversion pulse and an optimized RF acquisition pulse, both derived from the Shinnar Le-Roux (SLR) algorithm [28]. These optimized RF pulses allow slice profiles with sharp edges so that, in the selective condition, the alignment between the inversion slice and the image slice is as good a match as physically possible. Sections of inverted magnetization (slabs) are created by applying SLR inversion pulses in the presence of slice selection gradients. The inversion slab thickness is controlled by

\[2D\ fast\ SPGR\ data\ acquisition\ scheme\]

\[TR\]

Figure 3. Spin-tagging sequence. $TR$ is fixed at 2.7 sec. In each TR interval, an inversion RF pulse is applied first, followed by phase encoding steps starting at zero $k$ space. The time $TI_{br}$ is changed by changing the pulse sequence control variable $t_{befiv}$. The RF pulses are magnified, and $t_{befiv}$ and $TI_{br}$ are not to scale (actually longer than shown).
changing the gradient strength. By applying the SLR pulse without an associated slice selection gradient, a non-selective slab containing inverted spins in the entire sensitive volume of the transmit RF coil is created. By applying the pulse with the gradient, a selective slab of specific thickness is created. 3. The so-called odd-number interleaved k-space acquisition [29,30] has been implemented to allow high spatial resolution in the phase encoding direction. With this acquisition, temporal resolution can be traded for image spatial resolution. Overall, the time required to acquire an arterial spin tagging image at a single slice is 2 minutes 30 seconds. This longer time than originally proposed is due to the fact that images at a multitude of TI times were needed to accurately estimate the contribution of blood flow the apparent longitudinal relaxation time, and thus to estimate perfusion.

**Our protocol using the pulse sequences**

The protocol has been modified from that described in the proposal, to account for the longer scan time required for the spin tagging sequence. For each subject, 10 or more equally spaced slices in the breasts are analyzed. Breasts are scanned bilaterally. A dual phased array breast imaging RF coil (Medical Advance, Inc., consigned specifically for this project) is used for the breast studies. Spin tagging measurement is based on three conditions during image acquisition: 1. selective inversion slab on (Selective Condition), 2. non-inversion slab on (Non-selective Condition), and 3. no inversion slab (Regular Condition). At each slice, a time series of 61 images are acquired with a fixed time of repetition (TR) of 2.7 seconds, with a total scan time of about 3 minutes. At each time series, Conditions 1, 2 and 3 are applied in sequence. For either the Selective or Non-selective condition, there are seven TI cycles with seven different TI's (interval between the spin inversion and the effective center of k space of each image acquisition). In each TI cycle, four images are acquired. For the Regular Condition, five images are acquired (Figure 4). Repeated image acquisition with the identical parameters is done to improve the signal-to-noise ratio (SNR). The total scan time for the whole scan of 12 slice locations is about 30 minutes.

**Contributions to data analysis**

A data analysis technique described below has been developed from scratch (no prior description in the literature) to properly interpret the timeseries of signal changes obtained at each pixel. It has been developed taking into account the characteristics of 2D fast SPGR data acquisition. The analysis rests on the assumption that the 2D fast SPGR data acquisition does not disturb the recovery curve of the inverted spins or the equilibrium states of non-inverted spins. Although not strictly true, the deviation from reality does not significantly affect the outcome of the
measurements. Using images obtained under the “Regular” and the Selective or Non-selective Tagging Conditions, $T_{ls}$ or $T_{ln}$ is calculated at each pixel using a semi-log linear regression (Figure 5),

$$\frac{1}{T_{ls}} = -\frac{\ln (S_{reg} - S_{sel}(i)) - \ln (S_{reg} - S_{sel}(j))}{TI_{br}(i) - TI_{br}(j)}$$

(1.8)

and

$$\frac{1}{T_{ln}} = -\frac{\ln (S_{reg} - S_{non}(i)) - \ln (S_{reg} - S_{non}(j))}{TI_{br}(i) - TI_{br}(j)}$$

(1.9)

where $S_{reg}$ = the average steady-state signal intensity at a pixel under the Regular Condition, $TI_{br}(i)$ = the ith effective inversion time (TI) (used for the ith image), $S_{sel}(i)$ = the steady-state signal intensity at a pixel under the Selective Condition using the ith TI, and $S_{non}(i)$ = the steady-state signal intensity at a pixel under the Non-selective Condition using the ith TI. According to the theory (reviewed above), the value of $f/\lambda$ is the difference between the inverse of the $T_1$ values as expressed in Eq.(1.7). A derivation of the above linear regression equations is given in the Appendix.

As mentioned previously, four images are acquired for each TI under the Selective Condition, four images are required for each TI under the Non-selective Condition, and five images are acquired under the Regular Condition. Generally, we have observed that only the last three repeated images are at the steady state for their particular condition. The first two images the signal intensity is approaching the steady state. Therefore, only these last three are used for the above linear regression. A total of 21 data points are available for the calculation of $T_{ls}$ or $T_{ln}$, however, the three low signal data points near the zero-crossing line are unreliable and are usually eliminated (based on an outlier analysis). Finally, there are 18 remaining points for the calculation of the $T_{ls}$. Because the $T_1$ and $f/\lambda$ calculations based on this pulse sequence are very sensitive to structured and random noise, in particular the structured noise due to motion (pixel displacements), two noise-reduction techniques have been implemented. In the first technique, the resolution of each image is reduced from 256 x 256 to a lower resolution, for example, 64 x 64, by pixel averaging. This improves the signal-to-noise per pixel, and reduces the effect of small tissue motions occurring during the 61 images being analyzed. This reduction is due to the simple fact that any particular
motion results in a lower fractional change in the pixel volume. The second technique uses a data elimination strategy to reduce the mean-square-error of the semi-log regression used for determination of the $T_1$s. This technique permits the elimination of up to five noisy data points (out of 18) to reduce the $T_1$ regression error.

Contributions to image processing

A computer program, called BrView, has been developed for display, analysis, and interpretation of the images obtained in the breast imaging protocol, starting from previous programs developed by the PI for neuro-functional MRI research (as described in the proposal). The program is written in C, the X Window System, and Motif. For all of the analysis methods described above, the values of $T_{1s}$, $T_{1n}$ and $f/\lambda$ at different pixel locations on each slice are calculated. Images, which we call feature images, are formed based on these values. The three most useful feature images are those showing $T_{1n}$ values, the perfusion image showing $f/\lambda$ values, and the $f/\lambda$ error image based on the standard deviations of $f/\lambda$ values. These feature images contain pixels with a wide range of values. However, values of $T_1$, $f/\lambda$ and $f/\lambda$ errors of normal breast tissues are within narrow ranges, while pixels displacing large $f/\lambda$, $T_1$ above 500 ms, but small $f/\lambda$ error are regarded as suspicious of malignant disease. Pixels containing fat display very short $T_1$ and the perfusion values at these pixels tend to be inconsistent. To display these feature images, an appropriate positive shift, scaling and window size is applied [31].

This BrView program is used to display and interpret the feature images for identification of suspicious regions. Since a sizable breast lesion covers several pixels within the same slice and possibly neighboring slices, these pixels should have similar tissue characteristics. It is assumed that pixels with a breast lesion will have approximately equal $T_1$ values, and that these $T_1$ values will be relatively high compared to fat, and also show high perfusion rates with low perfusion standard errors. When a pixel is found to have a moderately long $T_1$, and a high $f/\lambda$, and the $f/\lambda$ error is low, surrounding pixels within the same slice and across slices are also checked to see if they have similar characteristics. The “suspicion index” for this pixel is defined as the percentage of surrounding pixels also classified as suspicious. By applying this method to all pixels in all slices, “suspicion index maps” are created. If the pixel has a suspicion index exceeding a threshold, this pixel would be highlighted on the display. Clearly, the level of suspicion here is qualitative. Neither a rigorous statistical analysis of the procedure, nor a statistical comparison and analysis of suspicion regions with the results of dynamic contrast enhanced MRI, has not been carried out yet. Nevertheless, the goal is to develop a quantitative tool for breast cancer detection. The suspicion index threshold for identification of malignant disease, as well as independent thresholds for $f/\lambda$, $T_1$ and the standard deviation of $f/\lambda$, should ultimately be based on the results of a large set of patient studies.

Based on the small set of studies that have been done, the thresholds used to identify suspicious pixels are as following:

$$T_1 > 0.5 \text{ sec}$$
$$f/\lambda > 0.1 \text{ sec}^{-1}$$
$$\text{STD of } f/\lambda < 0.1 \text{ sec}^{-1}.$$  

and the suspicion index threshold, used for asserting a “high” probability that the tissue is abnormal, is set at 20%. A quantitative analysis of the false positive and negative error rates
associated with this threshold has not been done. The BrView program allows clinician researchers to change the thresholds and observe the changes in the suspicion index maps. This visualization program can simultaneously display feature images and standard clinical MR images. Parameters $T_1, f/\lambda$, and the standard deviation of $f/\lambda$ at any pixel location of the breast objects can be found by a click of the mouse on that location. If a dynamic contrast enhancement study has been done, a click of the mouse reveals a plot of the change of signal intensity as a function of time at this location. The suspicious regions can also be mapped to high-resolution traditional clinical images (256 x 256), which are always acquired for comparison during each subject study. This mapping provides a radiologist the traditional view of the breast images with the suspicious regions identified. In summary, BrView is designed to allow pixel cross-registration and comparison between results from different MR imaging techniques. It provides an efficient working environment for clinicians making diagnostic decisions.

Other image processing contributions are methods to reduce tissue misregistration that cause inaccuracy in the $T_1$ estimation. Breast images are expected to precisely align in all 61 images acquired in the timeseries at each slice location. Unfortunately, the patient may move during the two minute 30 second scan, resulting in a change in breast position and orientation. The motion estimation used here is based on binary segmentation, following by statistical best-fit alignment of defined regions across the collection of 61 images. Geometric constraints of patient motion and breasts, within the scanner and dedicated RF coil, suggest that only lateral displacements in the X and Y directions should be considered. The studies performed suggest that typical displacements are not large, within 1 cm. Such motions are correctable with segmentation and region alignment (see Figure 6).

**Results of in-vitro and in-vivo testing**

The success of the arterial spin tagging pulse sequence in measuring perfusion was first proven by the result of flow phantom studies. All studies showed that there was a linear relationship...
between \(f/\lambda\) and actual flow. All evaluated flows were within the range of tissue perfusion (Figure 7). Furthermore, the \(T_{1n}\) values obtained in the non-selective condition, at four different flow rates, were as expected, specifically 1.232 +/- 0.034 sec. A brain imaging study was performed to further evaluate the arterial spin tagging pulse sequence. In 11 different regions of interest (ROIs), defined at several different slice locations, white matter had \(T_{1n}\) equal to 0.495 +/- 0.021 sec, \(T_{1s}\) equal to 0.487 +/- 0.022 sec, and \(f/\lambda\) equal to 0.030 +/- 0.075 sec\(^{-1}\), ranging from 0 to 0.131 sec\(^{-1}\). Gray matter in this study had \(T_{1n}\) equal to 0.665 +/- 0.065 sec, \(T_{1s}\) equal to 0.629 +/- 0.035 sec, and \(f/\lambda\) equal to 0.081 +/- 0.054 sec\(^{-1}\), ranging from 0.037 to 0.170 sec\(^{-1}\). In two normal breast studies, at 14 different ROIs defined at several different locations, parenchyma (glandular tissue) had \(T_{1n}\) equal to 0.772 +/- 0.088 sec, \(T_{1s}\) equal to 0.709 +/- 0.062 sec, and \(f/\lambda\) equal to 0.109 +/- 0.053 sec\(^{-1}\), ranging from 0.043 to 0.190 sec\(^{-1}\). Normal fat tissue had \(T_{1n}\) equal to 0.377 +/- 0.069 sec, \(T_{1s}\) equal to 0.363 +/- 0.056 sec, and \(f/\lambda\) equal to 0.086 +/- 0.073 sec\(^{-1}\), ranging from 0.003 to 0.177 sec\(^{-1}\). The values resulting from the head and normal breast studies were within expected ranges.

Results of patient studies

Introduction
For this and all cases summarized below, suspicious pixels were identified based on the following criteria:

\[ T_1: > 0.5 \text{ sec} \]
\[ f/\lambda: > 0.1 \text{ sec}^{-1} \]
\[ \text{STD of } f/\lambda: < 0.1 \text{ sec}^{-1}. \]

and the suspicion index threshold was set at 20.2 %. Analysis was done using the program BrView.

Case 1
Case Study #1: palpable mass
Date of study: Dec 10, 1997
Level of suspicion based on \(T_{1n}\) and perfusion: Moderately High
Clinical MR images: Negative (lesion not visualized)
Assessment based on spin tagging: Positive for malignancy
Biopsy result: benign (spin tagging INCONSISTENT with biopsy)

Discussion: Figure 8 shows the three feature images, and an image acquired under the “Regular Tagging Condition” of the arterial spin tagging sequence. All three feature images are 64 x 64. The \(T_1\) image has excellent SNR and clearly shows the distribution of \(T_1\) values. In particular, bright areas at the centers of the breasts represent glandular tissue with long \(T_1\) values. The corresponding perfusion image has relatively low SNR. Some pixels at the edge of the breasts have high intensity due to high calculation error, not high \(f/\lambda\). The calculation at edge pixels is extremely sensitive to breast motion, as well as to the presence of Gibbs ringing artifacts common in low resolution images. Pixels with high \(T_1\) values generally have less perfusion error. Because the perfusion estimates are more sensitive to errors in \(T_1\) when \(T_1\) is low. Fortunately, this property of the error should not limit the identification of malignant tissue very much, because malignant tissue will typically have relatively high \(T_1\) values (> 500 ms). Figure 9 shows the overlay of the suspicious regions onto the high-resolution clinical images. Highlighted
(pure white) pixels on the left breast (right side of image) on slices 4, 5, 6 and 7 may be identifying one contiguous mass. Physical examination of this patient revealed a mass at the nine o'clock position in the left breast, consistent with the suspicious cancer region identified by spin tagging.

**Case 2**
Case Study #2: palpable mass
Date of study: Nov 16, 1997
Level of suspicion based on $T_1$ and perfusion: Low
Clinical MR images: Positive (lesion visualized, consistent with palpation)
Assessment based on spin tagging: Negative for malignancy
Biopsy result: benign (spin tagging CONSISTENT with biopsy)

Discussion: The low suspicion index, based on $T_1$ relaxation time and perfusion, at and around the palpated lesion location was strong indication that the lesion was benign. Only one suspicious pixel was found at that location at the 20.2% threshold. It was judged that the breasts did not contain malignant tissue. This case is shown in Figures 10(a)-(d).

**Case 3**
Case Study #3: Palpable mass
Date of study: Aug 6, 1997
Level of suspicion based on $T_1$ and perfusion: High
Clinical MR images: Negative (mass not clearly visualized, prior cyst visualized)
Result based on spin tagging: Positive for malignancy
Biopsy result: Positive (spin tagging CONSISTENT with biopsy)

Discussion: The high suspicion index, based on $T_1$ relaxation time and perfusion, at and around the lesion detected by palpation, was strong indication that the lesion was malignant. This case is shown in Figures 11(a)-(d).

**Case 4**
Case Study #4: Palpable mass
Date of study: Aug 7, 1998
Level of suspicion based on $T_1$ and perfusion: High
Clinical MR images: Positive (lesion visualized)
Assessment based on spin tagging: Positive for malignancy
Assessment based on dynamic contrast-enhanced study: Positive for malignancy
Biopsy result: Negative (spin tagging NOT CONSISTENT with biopsy)

Discussion: High suspicion indices, based on relaxation time and perfusion, were found in several breast regions. The visible lesion location on clinical MRI was consistent with that of the palpable mass. It overlapped but was considerably smaller than that found by the suspicion indices. First-pass contrast enhancement study results correlated strongly with the high perfusion regions detected by spin tagging. Nevertheless, biopsy showed the mass to be benign. This case is shown in Figures 12(a)-(d).

In summary, there was only partial agreement (2 of 4) between spin tagging and biopsy assessments of malignant disease. More patient studies (as planned in project), as well as follow-up of patients, is needed. Spin tagging may be detecting abnormal regions not evaluated by the
biopsies. Arterial spin tagging results were, subjectively, more convincing if the suspicious regions overlapped abnormal regions seen on standard clinical $T_1$, $T_2$ and proton-density weighted images. Unfortunately, standard clinical images often did not reveal any abnormal tissue, concurring with the view that standard clinical MRI is not definitive in the assessment of breast lesions. Dynamic first-pass contrast-enhanced imaging is definitely needed to help validate the spin tagging technique, and this imaging will be done on all subsequent patients in this project. In addition, results from X-ray mammography and ultrasonography will be compared, as has been done for prior studies (see page 40).

**Future Implementations**

**Arterial spin tagging with an echo planar type data acquisition**

To derive the semi-log regression analysis, it was assumed that SPGR data acquisition would not disrupt the recovery of longitudinal magnetization of inverted spins, nor the equilibrium of the non-inverted spins. Unfortunately, this assumption is a cause of error in the method, and our simulations have shown that the $T_1$ calculated based on the assumption was slightly lower than the true $T_1$. Using the simulation results, a conversion table was developed to convert the estimate derived based on the assumption to a better $T_1$ estimate. Nevertheless, a better approach would be to replace the 2D fast SPGR data acquisition scheme with an echo planar data acquisition scheme. Echo planar imaging (EPI), requires only a single RF pulse for acquisition of the entire data set for one image. The disturbance due to the repeated RF pulses to the recovery curves of the inverted spins or the equilibrium states of the non-inverted spins would be much more predictable. Also, EPI is faster than 2D fast SPGR, so the $TI$ cycles could be redesigned, and a multi-slice technique introduced, to shorten the scan time by a factor of 4-8. This implementation would make possible high resolution scanning of the entire breast tissue in under 10 minutes. Starting September 1998, the UC Davis Medical Center will operate a GE Signa Horizon Echospeed MR system. This system will be capable of echo planar imaging (128 x 128 single shot is typical, but up to 256 x 256 single shot can be done). Conversion of the arterial spin tagging sequence to incorporate echo planar acquisition will be carried out. In addition, conversion will also be done for the existing spin tagging sequence that is based on 2D fast SPGR.

**Magnetization transfer implementation**

The second major assumption, used in the derivation of the perfusion formula, is that the contribution of the magnetization transfer term in Eq. (1.1) is negligible. To allow inclusion of this term, the spin tagging pulse sequence must include a binomial or high order RF pulse to prepare the spins prior to data acquisition. Differential magnetization transfer effects have been detected in head and neck neoplasms [32]. Development of this pulse sequence modification to estimate the size of the magnetization transfer contribution will be completed.
Figure 8. Dec 10, 1997. Example of feature images derived at single slice. (a) Image obtained during spin tagging sequence during Regular Tagging Condition. (b) $T_1$ image derived, at each pixel, from best-fit of signal intensity from sequence of images obtained at different inversion times ($TI$). (c) Perfusion ($j/\lambda$) image derived from comparison of $T_1$ images during Non-selective and Selective Tagging Conditions. (d) Perfusion error image, measured as standard error based on $T_1$ standard errors.
Figure 9. Dec 10, 1997 study. The suspicious pixels (threshold 20.2%), showing moderate $T_1$, high $f/\lambda$, low $f/\lambda$ error, are identified on six (a-f) contiguous slices, displayed on fat suppressed $T_2$ weighted fast spin echo images.
Figure 10(a). Nov 16, 1997 study. Suspicion index map is created using the “Create Detection Image” button after setting thresholds.
Figure 10(b). Nov 16, 1997 study. Feature images (top row), suspicion index map (bottom left, threshold 0.5%), and mapping (pure white pixels) of the map on proton density and T2 images (bottom row). Also shown: Axial T1.
Figure 10(c). Nov. 16, 1997 study. Feature images (top row), suspicion map with the suspicion index threshold at 20.2% (revealing single pixel) and mapping (pure white pixel) onto proton density and $T_2$ images. Clicking on pixels displays numerical $T_1$, $f/\lambda$, standard error of $f/\lambda$, and suspicion index. Also shown: Axial $T_1$. 
Figure 10(d). Nov. 16, 1997 study. The $T_1$ and $f/\lambda$ values of pixels representing the lesion identified by proton density and $T_2$ images, are superimposed on the histogram of the $T_1$ and $f/\lambda$ values from the entire breast. Shows that criteria for suspicious pixels is based on moderate $T_1$ and high $f/\lambda$. See text for complete criteria.
Figure 11(a). Aug 6, 1997 study. Suspicion index map is created using the “Create Detection Image” button after setting thresholds.
Figure 11(c). Aug 6, 1997 study, slice adjacent to that shown in Figure 11(b). Feature images (top row), suspicion index map with threshold at 20.2%, and mapping (pure white pixels) onto proton density and T2 images. Also shown: Axial $T_1$. 
Figure 11(d). Aug 6, 1997 study. The $T_1$ and $f/\lambda$ values of pixels representing the lesion identified by proton density and T2 images in Figure 11(c), are superimposed on the histogram of the $T_1$ and $f/\lambda$ values from the entire breast. Shows that criteria for suspicion is based on moderate $T_1$ and high $f/\lambda$. See text for complete criteria.
Figure 12(a). Aug 7, 1998 study. Suspicion index map is created using the “Create Detection Image” button after setting thresholds.
Proton density, T2, and T1 images are shown without suspicion index map (bottom row). Also shown: Axial T1.
Figure 12(c). Aug 7, 1998 study. Feature images (top row), suspicion index map with threshold at 20.2% (revealing one cluster of pixels) and mapping (pure white pixels) onto proton density and T2 images. Also shown: Axial T1.
Figure 12(d). Aug 7, 1998. Timeseries of signal changes at a rapidly enhancing pixel during dynamic contrast enhancement. Also shown: $T_1$ “feature” image (1st from left), pre-contrast 3D image (2nd from left), and post-contrast 3D image, last image of dynamic study, $T_2$ image, and $T_1$ axial image.
(7) CONCLUSIONS

In this year of the project, we have developed the necessary pulse sequences for anatomical and functional imaging of breast tissue, and the necessary software for the analysis, display and interpretation of this data. These developments will allow us to evaluate complete breast imaging studies that are composed of anatomical scans, dynamic first pass contrast enhanced scans, and arterial spin tagging scans. The integrated nature of the software allows easy and consistent interpretation of both spatial dependencies as well as temporal dependencies of signal changes indicative of disease. The in-vitro and in-vivo studies done thus far indicate that first and foremost, arterial spin tagging is a sensitive and reliable method for measuring $T_1$ and parameters related to tissue perfusion. Second, they indicate that arterial spin tagging is a viable alternative to dynamic first-pass contrast-enhanced imaging. The fact that arterial spin tagging can be easily added to any standard breast imaging protocol, without requiring special nursing or MR technologist expertise, means that it is especially attractive for routine clinical use. It could be used to supplement contrast-enhanced studies in the initial study of the patient, and used exclusively in follow-up studies. Since 10/1/98, six subjects have been studied without the dynamic first-pass contrast-enhanced scan, and two have been studied with this scan. Comparison of contrast-enhanced images versus arterial spin-tagged images has not yet been made. In particular, no statistical comparison of the dynamic study results (considered to be the gold standard for non-invasive assessment of malignant disease) and the arterial spin tagging results have been performed. However, there is considerable spatial overlap of suspicious regions obtained by these methods noted on visual inspection. Collecting data on human subjects, and statistical confirmation of this agreement, will be major focuses of next year’s work.

(8) REFERENCES


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(9) APPENDIX

Contributions to Data Analysis

The regression method derived in this project resolves a key confound that is present in all arterial spin tagging sequences, that of the mismatch of the slice profiles of the inversion and excitation RF pulses. Our sequence is longer than many arterial spin sequences reported in the literature, but we are quite sure that these other methods fail to account for the mismatch, and result in dubious measurements of perfusion. It is interesting to note that the mismatch results in signal contamination whose spatial dependence resembles an inverted $T_1$ weighted image, thus, they are easily mistaken for true perfusion maps (i.e. in the brain, CSF is bright, followed by gray matter, then white matter, just as true perfusion would present).

Derivation of semi-log linear regression method

In this derivation, it is assumed that, because the excitation RF flip angle is small, repeated application of these pulses for data acquisition does not disturb $T_1$ recovery of the inverted spins, nor the equilibrium magnetization of the imaged spins if not inverted. Given this assumption, under a spin tagging condition, the steady-state longitudinal magnetization at the end of each $TR$ period just before the next inversion pulse would be

$$M_{10ss} = M_0 (1 - e^{-bTR}) - M_{10ss} e^{-bTR}$$

where, $M_{10ss} = $ the steady-state longitudinal magnetization at the end of each $TR$ period, $M_0 = $ the longitudinal magnetization under fully relax condition, $TR = $ the repetition time, and

$$b = \begin{cases} \frac{1}{T_{1s}} & \text{for the Selective Tagging Condition} \\ \frac{1}{T_{1n}} & \text{for the Non-selective Tagging Condition} \end{cases}$$

Therefore,

$$M_{10ss} = M_0 \frac{1 - e^{-bTR}}{1 + e^{-bTR}} \quad (A.1)$$

The longitudinal magnetization at time $t$ after the spin inversion at steady-state is
\[ M_1(t) = M_0 (1 - e^{-bt}) - M_{0ss} e^{-bt} \]
\[ = M_0 (1 - e^{-bt}) - M_0 \frac{1 - e^{-bTR} e^{-bt}}{1 + e^{-bTR} e^{-bt}} \]
\[ = M_0 (1 - e^{-bt} - \frac{1 - e^{-bTR} e^{-bt}}{1 + e^{-bTR} e^{-bt}}) \]
\[ = M_0 (1 - \frac{2e^{-bt}}{1 + e^{-bTR}}) \]

As a result, the signal measured at echo time \( TE \) is

\[ S_t = M_0 (1 - \frac{2e^{-bTR}}{1 + e^{-bTR}}) \sin (\theta) e^{\frac{TE}{T_2}} \]  \hspace{1cm} (A.3)

where, \( T_I = \) the time of inversion from the inversion pulse to the effective center of k space, \( TE = \) the time of echo at each read-out period, and \( \theta = \) the excitation flip angle.

Under the Regular Tagging Condition, the longitudinal magnetization \( M_{\text{reg}}(t) \) at any time \( t \) stays at equilibrium. Specifically,

\[ M_{\text{reg}}(t) = M_0 \]  \hspace{1cm} (A.4)

The signal measured at echo time \( TE \) becomes

\[ S_{\text{reg}} = M_0 \sin (\theta) e^{\frac{TE}{T_2}} \]  \hspace{1cm} (A.5)

Thus,

\[ \frac{S_t}{S_{\text{reg}}} = 1 - \frac{2e^{-bTR}}{1 + e^{-bTR}} \]  \hspace{1cm} (A.6)

and,

\[ 1 - \frac{S_t}{S_{\text{reg}}} = \frac{2e^{-bTR}}{1 + e^{-bTR}} \]

Thus, considering the \( i \)th inversion time (TI):

\[ \frac{S_{\text{reg}} - S_t(i)}{S_{\text{reg}} - S_t(j)} = \frac{e^{-bTR(i)}}{e^{-bTR(j)}} = e^{-(T_I(i) - T_I(j))} \]  \hspace{1cm} (A.7)

where, \( T_I(i) = \) the time of inversion at the \( i \)th \( T_I \) cycle, \( S_t(i) = \) the signal at a pixel when \( T_I(i) \) is used, \( T_I(j) = \) the time of inversion at the \( j \)th \( T_I \) cycle, \( S_t(j) = \) the signal at a pixel when \( T_I(j) \) is used. Taking the logarithm of both sides yields,
\[
\ln (S_{reg} - S_i(i)) - \ln (S_{reg} - S_i(j)) = -b(TI(i) - TI(j))
\]

and solving for \(b\) gives,

\[
b = -\frac{\ln (S_{reg} - S_i(i)) - \ln (S_{reg} - S_i(j))}{TI(i) - TI(j)}
\]

(A.8)

The above equation shows the semi-log linear regression relationship between \((S_{reg} - S_i)\) and \(TI\). The formula can be applied to both Selective and Non-selective tagging conditions, to derive \(b_s\) and \(b_n\), respectively. The exact values of the effective \(TTs\) are uncertain (due data collection is spread out over 1 second). Nevertheless, it is reasonable to assume that these values each have the same amount of time error relative to their true values. Because \(TI(i)\) and \(TI(j)\) is subtracted in the denominator, the uncertainty of the timing of this “effective” \(TI\) cancels out.

**Elimination of slice profile mismatch effects**

The semi-log relationship eliminates the errors due to mismatch between the slice profiles of the inversion and data acquisition RF pulses. Because the RF pulses used for inversion and data acquisition are different under the Selective Condition, (The Non-selective Condition does not have this problem because the RF inversion pulse inverts spins everywhere.).

The geometric mismatch between the inversion and read-out profiles would lead to the contamination of signal from non-inverted spins. Thus the apparent signal from the Selective Condition can be modeled as

\[
S_{selapp} = (1 - err)S_{sel} + err S_{reg}
\]

(A.9)

where \(S_{sel}\) = the signal of a pixel if no slice profile mismatch, \(S_{selapp}\) = the signal of a pixel actually measured, and \(err\) = the amount of percent error due to slice profile mismatch. Then,

\[
\frac{S_{selapp}}{S_{reg}} = (1 - err)\frac{S_{sel}}{S_{reg}} + err
\]

and

\[
\frac{S_{sel}}{S_{reg}} = \frac{S_{selapp} - err}{S_{reg} 1 - err}
\]

Thus,
Comparing with Eq. (A.7), we have

\[
\frac{S_{\text{reg}} - S_{\text{sel}}}{S_{\text{reg}}} = \frac{1 - \text{err} - \frac{S_{\text{sel}}}{S_{\text{reg}}}}{1 - \text{err}}
\]

\[
1 - \frac{S_{\text{sel}}}{S_{\text{reg}}} = \frac{1}{1 - \text{err}}\]

Thus,

\[
\frac{S_{\text{reg}} - S_{\text{sel}}}{S_{\text{reg}}} = e^{-b_z (T(i) - T(j))}
\]

where,

\[
b_z = \frac{1}{T_{Is}} = \frac{1}{T_I} + \frac{f}{\lambda} = \text{the } b \text{ value under the Selective Condition}
\]

Thus,

\[
b_z = \frac{\ln (S_{\text{reg}} - S_{\text{sel}} (i)) - \ln (S_{\text{reg}} - S_{\text{sel}} (j))}{T(i) - T(j)}
\]

Eq. (A.11) corresponds exactly as Eq. (A.8). In other words, by using the semi-log linear regression method, the problem due to geometric mismatch between the inversion and read-out slice profiles is eliminated.

**Phantom, normal subject, and patient study notes**

These notes (pages 41-50, presented in two-column landscape mode) were written immediately after each study, and provide brief subject history, and detailed listing of pulse sequences and parameters used in the protocol, and problems noted. Preliminary accounts of studies prior to 11/19/97 have been reported [33,34,35].

**Inventory of Breast Imaging Studies to Date**

<table>
<thead>
<tr>
<th></th>
<th>Test Phantoms</th>
<th>Normal Subject</th>
<th>Patient (no CE-MRI)</th>
<th>Patient (no CE-MRI)</th>
<th>Patient (with CE-MRI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Prior to 10/1/98</td>
<td>9</td>
<td>4</td>
<td>3</td>
<td>3b, 2m</td>
<td>0</td>
</tr>
<tr>
<td>Since 10/1/98</td>
<td>5</td>
<td>1</td>
<td>1</td>
<td>4b</td>
<td>2b</td>
</tr>
</tbody>
</table>

CE-MRI (dynamic first-pass contrast-enhanced MRI). Number preceding b indicates number of subjects in which the biopsy revealed benign disease. Number preceding m indicates number of subjects in which the biopsy revealed malignant disease. Patients without biopsy did not meet clinical criteria and mass was assumed benign.
Subject Study Summary:

Study Descriptions: February 2, 1997 through August 5, 1998

(Procedure Updated on 2/7/97)
Date of study: 2/7/97

Breast Sequence Study
by Michael Buonocore and David Zhu

Patient: Reva (NOMINAL SUBJECT)

Suggest procedures:
1. 2D Spin Echo axial cut (Series 1), TR = 600 ms, 1 acquisition
2. 2D fast spin echo (Series 2)
   TE=128 msec, TR=17 msec, TR=4s.
   5mm with 1 mm interslice, center.
   Increases to 20 slices
3. 3D scan coronal cut (Series 3):
   Location:
4. David's breast sequence coronal cut: fglm_new
   Five TI cycles, br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 150 ms.
   TI decreases from 615 ms to 15 ms.
   Slice thickness: 3 mm
   Frequency direction: R/L for all locations
   Prescan: R1 = 6 , R2 = 15 , TG =169 or 167
   series 4: location: p20
   series 5: location: p15
   series 6: location: a0
   series 7: location: a22
   series 8: location: a30
   series 9: location: a40
   series 10: location: a50
5. Redo 3D scan coronal cut (series II), type: SPGR PROSP

(Procedure Updated on 2/7/97)
Date of study: 3/9/97

Flow Phantom Study
by David Zhu

Coil: breast coil

Suggest procedures:

David's breast sequence axial cut: fglm_new
Five TI cycles, opphases = 55, br_TR = 2.7 s, br_TI = 15 ms, 
ti_inc = 150 ms. TI decreases from 615 ms to 15 ms.
Slice thickness: 3 mm
br_sel_sl = opslthick = 3mm
Frequency direction: R/L
phantom on the left of the breast coil in prone.
Location: A20

Flow rate: Motor mark
series 1: Motor off R1 = 6, R2 =15, TG =107
series 2: 40 ml /91.93 sec 0.5 TO = 112
series 3: 100 ml /79.62 sec 0.8 TO = 114
series 4: 80 ml / 56.47 sec 0.66 TO = 123
series 5: 100 ml / 64.68 sec 0.9 TO = 115
series 6: 100 ml /54.46 sec 1.0 TO = 123

Date of study: 3/9/97

Breast Sequesnt Study
by Michael Buonocore and David Zhu

Subject: Marijana (Fibroadenoma)

Local Coil (lps)
Squire, mamil. fov = 22cm, ciftall, cftall and cftfull are scaled down.

Procedures:
1. 2D Spin Echo coronal cut (Series 1)
2. David's breast sequence coronal cut: fglm_new
   Five TI cycles, br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 150 ms.
   TI decreases from 615 ms to 15 ms.
   Acquisition slice thickness: 3 mm
   Selective slice thickness: 3mm
   Frequency direction: S/I for all locations
   Prescan: R1 = 6 , R2 = 15 , TG = 113
   FOV: 22 cm
   series 2: location: F24
   series 3: location: P15
   series 4: location: A0
   optr = 10444
   opte = 9344

Date of study: 3/9/97

Breast Sequence Study
by Michael Buonocore and David Zhu

Subject: Marijana (Fibroadenoma)

Procedures:
1. 2D Spin Echo axial cut (Series 1)
2. 2D fast spin echo (Series 2)
3. 3D scan coronal cut (Series 3): Patient needed to add
4. David's breast sequence coronal cut: fgtl_new

Five TI cycles, br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 150 ms.
TI decreases from 615 ms to 15 ms.

Acquisition slice thickness: 3 mm
Selective slice thickness: 3 mm
Frequency direction: R/L for all locations
Prescan: R1 = 6, R2 = 15, TG = 177

series 4: location: A10
series 5: location: A13
series 7: Repeat series 5 with the same location, but with a br_TI = 50 ms,
so TI decrease from 650 ms to 50 ms
series 8: location: A16

Date of study: 5/3/97

Breast Sequence Study
by Michael Buonocore and David Zhu

Patient: Nancy Saunders

Procedures:
1. 2D Spin Echo axial cut (Series 1)
2. 2D fast spin echo (Series 2)
3. 3D scan coronal cut (Series 3): Patent is added,
P26.8 to A108.9
Scan thickness: 2.3 mm

4. David's breast sequence coronal cut: fgtl_new

Five TI cycles, br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 150 ms.
TI decreases from 615 ms to 15 ms.

Acquisition slice thickness: 3 mm
Selective slice thickness: 3 mm
Frequency direction: R/L for all locations
Prescan: R1 = 6, R2 = 15, TG = 180 (autoprescan fails with the limit of
TG at 172, manually set 180)

series 4: location: A23.8
series 5: location: A21.5
series 6: location: A05.9
series 7: location: A08.2

comments:
Suspected of fibroadenoma.
Two cysts or tumors at left breast.
Autoprescan fails with the limit of TG at 172, manually set 180.

Question: In inversion set at optimal 180 degree? Now to take care of the
non-optimal inversion pulse?

Date of study: 5/11/97 (May 11, 97)

Phantom study

Date of study: 5/17/97
breast sequence study
7up phantom
A10, R/L freq

40 cm FOV

(1) fgtl_new:
R1 = 6, R2 = 15, TG = 109

(2) fgtl_new
R1 = 6, R2 = 15, TG = 113

Date of study: 5/23/97

Breast Sequence Study
by Michael Buonocore and David Zhu

Patient Name: Jean (66 years old) (TUMOR RIGHT BREAST)

Procedures:
1. 2D Spin Echo axial cut (Series 1)
2. 2D fast spin echo (Series 2)
3. 3D scan coronal cut (Series 3):
P13.3 to A 51.5
Scan thickness: 2.4 mm.
# of scan loc: 28

4. David's breast sequence coronal cut: fgtl_new

Five TI cycles, br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 150 ms.
TI decreases from 615 ms to 15 ms.

Acquisition slice thickness: 3 mm
Selective slice thickness: 3 mm
Frequency direction: R/L for all locations
Prescan: R1 = 6, R2 = 15, TG = 146

series 4: location: A15.5
series 5: location: A20
series 6: location: A26
series 7: location: A32
series 8: location: A38

comments: Tumor locates at right breast

Date of study: 5/24/97

Breast Sequence Study
by Michael Buonocore and David Zhu

Patient Name: Quan (David's mother-in-law) (NORMAL SUBJECT)

age: 55

Procedures:
1. 2D Spin Echo axial cut (Series 1): TE = minimum, TR = 600 ms
2. 2D fast spin echo (Series 2): p60 = a66
3. 3D scan coronal cut (Series 3):
P 22.9 to A 68.9
Scan thickness: 3.4 mm

4. David's breast sequence coronal cut: fgtl_new3

Five TI cycles. br_TR = 2.7 s, br_T1 = 15 ms, ti_inc = 150 ms.
TI decreases from 615 ms to 15 ms.

Acquisition slice thickness: 3 mm
Selective slice thickness: 3mm
Frequency direction: R/L for all locations
Prescan: R1 = 6, R2 = 15, TO = 179

series 4: location: P 2.5
series 5: location: A 17.9
series 6: location: P 5.9
series 7: location: A 28.1
series 8: location: A 48.5
series 9: location: A55.3 (end prescan: R1 = 6, R2 = 15, TO = 171)

comments: Normal subject

To check whether we can run fgr using opal thickness of 1.7 mm.

Result:
(1) opal thickness = 1.7 mm, bw_rf1 = 700 Hz, opflip = 20
I.1 to 1.5
Autoscan successful: R1 = 6, R2 = 15, TO = 68
Scan successful.
(1) opal thickness = 3 mm, bw_rf1 = 1250 Hz, opflip = 20
I.6 to 1.10
Autoscan successful: R1 = 6, R2 = 15, TO = 68
Scan successful. These regular images appear to be darker than those
with opal thickness of 1.7 mm.

Date of study: 7/11/97

Breast: Sequence Study
by Michael Bonacure and David Zhu

Patient Name: Breast biopsy patient (MISSED TUMOR NEXT TO NIPPLE, OTHERWISE NORMAL)

age: about 60

Procedures:
1. 2D Spin Echo axial cut (Series 1): E30 to E30
2. 2D fast spin echo (Series 2)
3. 3D scan coronal cut (Series 3):
   P 112.2 to A 69.8
   Scan thickness: 5 mm

4. David's breast sequence coronal cut: fgtl_new3b

Five TI cycles. br_TR = 2.7 s, br_T1 = 15 ms, ti_inc = 150 ms.
TI decreases from 615 ms to 15 ms.

Acquisition slice thickness: 3 mm
Selective slice thickness: 3mm
Frequency direction: R/L for all locations
Prescan: R1 = , R2 = , TO =

series 4: location: A27.8, R1 = 4, R2 = 14, TO = 170 (might not be good)
series 5: location: A2, R1 = 6, R2 = 15, TO 179
series 6: location: A21
series 7: location: A30

comments: Tumor locates at right breast, about 3 o'clock near nipple, not biopsy yet.
We might have missed it. (Correct, did not scan at correct location)

Date of study: 7/26/97
7-up phantom testing

001: fgtl_new3b, default setting
   autoscan: R1 = 6, R2 = 15, TO = 109

001: fgtl_new3c, default setting
   autoscan: R1 = 6, R2 = 15, TO = 107
Date of study: 7/26/97
Breast Sequence Study
by David Zhu

Patient Name: Grandma (NORMAL)
age: 81

Procedures:
1. 2D Spin Echo axial cut (Series 1)
2. 3D scan coronal cut (Series 3):
   P 21.4 to A 59.6
   Scan thickness: 3 mm
4. David's breast sequence coronal cut: fgtl_new3c
   Seven TI cycles, br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 100 ms.
   TI decreases from 615 ms to 15 ms.
   Acquisition slice thickness: 3 mm
   Selective slice thickness: 3 mm
   Frequency direction: R/L for all locations
   Prescan: R1 = 6 , R2 = 15 , TG = 178
series 3: location: A6.6
series 4: location: A23.6
series 5: location: A35.6
series 6: location: A44.6

comments: Subject did not complain of any abnormality.

Date of study: 8/1/97:
Flow phantom study using fgtl_new3c:
Series Number Flow Mark
001 0
002 0.5
003 0.8
004 1.0

Date of study: 8/6/97
Breast Sequence Study
by Michael Buonocore and David Zhu

Patient Name: Josie (SIGNET CELL CARCINOMA)
age: 44

Procedures:
1. 2D Spin Echo axial cut (Series 1)
2. 2D fast spin echo (Series 2):
   P 2.9 to A 64.6
   Scan thickness: 2.5 mm
3. 3D scan coronal cut (Series 3):
   P 30 - A60
4. 3D scan coronal cut (Series 4):
   P 2.9 to A 64.6
   Scan thickness: 2.5 mm

whole breast
Scan thickness: 3.7 mm
5. Redo No. 2 due to patient movement. (Series 5)
6. David's breast sequence coronal cut: fgtl_new3c
   Seven TI cycles, br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 100 ms.
   TI decreases from 615 ms to 15 ms.
   Acquisition slice thickness: 3 mm
   Selective slice thickness: 3 mm
   Frequency direction: R/L for all locations
   Prescan: R1 = 6 , R2 = 15 , TG = 153
series 6: location: A14
series 7: location: A24
series 8: location: A30

comments:
Tumors are at right breast from various locations.
One big biopsy site at inner bottom of the right breast.
Probably cancer patient.

Aug 9, 1997: Heat phantom
Date of study: 8/9/97
Breast Sequence Study
by David Zhu

Phantom study (heat phantom always on left, prone position)

Procedures:
Place only the Heat phantom.
1. 2D Spin Echo axial cut (Series 1) : L10 to a30, 11 locs, thick: 5, inter: 1
   R1 = 6 , R2 = 15 , TG = 75
2. 2D fast spin echo (Series 2) : thick:5, inter:1, a30-a60, 8 locs.
   R1 = 6 , R2 = 15 , TG = 75
3. 3D scan coronal cut (Series 3):
   P 19.2 to A 59.7
   Scan thickness: 1.5 mm
   No of loc: 28
   R1 = 6 , R2 = 15 , TG = 76

4. David's breast sequence coronal cut: fgtl_new3c
   Seven TI cycles, br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 100 ms.
   TI decreases from 615 ms to 15 ms.
   Acquisition slice thickness: 3 mm
   Selective slice thickness: 3 mm
   Frequency direction: R/L for all locations
   series 4: location: a30 , R1 = 6 , R2 = 15 , TG = 116
series 5: location: a38 , R1 = 6 , R2 = 15 , TG = 115
series 6: location: a50, R1 = 6 , R2 = 15 , TG = 116

Place the Heat phantom with the get phantom
series 7: location: a38, R1 = 6 , R2 = 15 , TG = 106
Place the 7-up phantom with the 7-up phantom

series 8: location: A8  R1 = 6 , R2 = 15 , TG = 111

Purpose:
(1) Check perfusion with meat with no blood flowing.

(2) Observe how the prescan result changes with the placement of different tissue type.

Aug 10, 1997
Date of study: 8/10/97
Breast Sequence Study
David Zhu

Subject Name: Tracy
age: 19

Procedures:
1. 2D Spin Echo axial cut (Series 1)
   T1 84 - #44, loc: 29. Scan thick = 5 mm, Inten = 1 mm.

2. 2D fast spin echo (Series 2)
   P10 -> A56. Scan thick = 5 mm, inter = 1 mm

3. 3D scan coronal cut (Series 3):
   P 2.3 to A 49.2
   Scan thickness: 2 mm

4. David's breast sequence coronal cut: fg1-05c
   Seven TI cycles, br_T1 = 2.7 s, br_T2 = 15 ms, ti_inc = 100 ms.
   TI decreases from 515 ms to 15 ms.
   Acquisition slice thickness: 3 mm
   Selective slice thickness: 3 mm
   Frequency direction: R/L for all locations
   Prescan: R1 =  , R2 =  , TG =  

series 4: location: A2  R1 = 6 , R2 = 15 , TG = 140
series 5: location: A7  R1 = 6 , R2 = 15 , TG = 139
series 6: location: A10  R1 = 6 , R2 = 15 , TG = 140
series 7: location: A14  R1 = 6 , R2 = 15 , TG = 145
series 8: location: A20  R1 = 6 , R2 = 15 , TG = 149
series 9: location: A32  R1 = 6 , R2 = 15 , TG = 141
Series 10: repeat No. 2 (2d fat).
   P15 - A21, scan thick = 3 mm, inter = 1mm.

comments:
   Normal healthy subject. 10 months after giving birth.
   Subject complains that she has a mass on both breasts above nipple; the masses are harder than other tissues; they are about 2 cm in diameter; they have been there for 4 to 5 years; the size did not change.

Sep 13a, 97
Date of study: 9/13/97
Breast Sequence Study
David Zhu

Subject Name: Tracy
age: 19

Redo Study of 8/30/97
Ask subject to have a better fit of the foam cast, tighter strap, and not to move, and to breath smoothly with stomach.

Procedures:
1. 2D Spin Echo axial cut (Series 1)
   T1 60 - #44, loc: 21. Scan thick = 5 mm, Inter = 1 mm.

2. 2D fast spin echo (Series 2)
   R2 = A56, loc: 16, scan thick = 3 mm, inter = 1mm

4. David's breast sequence coronal cut: fg1-05c
   Seven TI cycles, br_T1 = 2.7 s, br_T2 = 15 ms, ti_inc = 100 ms.
   TI decreases from 515 ms to 15 ms.
   Acquisition slice thickness: 3 mm
   Selective slice thickness: 3 mm
   Frequency direction: R/L for all locations
   Prescan: R1 =  , R2 =  , TG =  

series 3: location: A3  R1 = 6 , R2 = 15 , TG = 138
series 4: location: A10  R1 = 6 , R2 = 15 , TG = 139
series 5: location: A14  R1 = 6 , R2 = 15 , TG = 140
series 6: location: A18  R1 = 6 , R2 = 15 , TG = 137
series 7: location: A22  R1 = 6 , R2 = 15 , TG = 138
series 8: location: A26  R1 = 6 , R2 = 15 , TG = 139
series 9: location: A30  R1 = 6 , R2 = 15 , TG = 137

comments:
   Normal healthy subject. 10 months after giving birth.
   Subject complains that she has a mass on both breasts above nipple; the masses are harder than other tissues; they are about 2 cm in diameter; they have been there for 4 to 5 years; the size did not change. Sometimes there is slight pulling kind of pain.

The motion-resist procedure shows to be effective based on the data acquired.

Sep 13b, 97
9/13/97
Study with 7-up bottles at location A = 0. See if the calculated Tin and f/lambda
different at lower locations.
Sequence: fg1-05c

9/20/97, (Sep 20, 97)
Heat phantom experiment using fg1-05c to check the image quality close to chest wall.
Sequence: fg1-05c

Sep 27, 97
9/27/97
Eat
7-up phantom study
by David Zhu
series 001:
Number of Images = 10, br_norm = 2, br_tino = 2, br_nons = 2, br_T1 = 100 ms, ti_inc = 200 ms, br_TR = 2.7 s
FOV = 34 cm

(1) I.001 = I.010; fgt1_new0d, set 256 x 128, others default
(2) I.011 = I.020; Addition to (1), set rhyracy = 128.
(3) I.021 = I.030; Addition to (2), set rhyracy = 0.5.
(4) I.031 = I.040; Addition to (3), set phasefov = 128.
(5) I.041 = I.050; fgt1_new0d, 256 x 256

series 002:
fgt1_new0d, default to 61 images, and for 34 cm

series 003:
fgt1_new0d, default to 61 images as in fgt1_new0c, but set 256 x 128, and
set rhyracy = 128, rhyracy = 0.5, phasefov = 128.

Purpose of study: See if fgt1_new0d better. It offers better resolution.

Date of study: 10/4/97

Different phantom study by David Zhu

(1) comparing fgt1_new0c with fgt1_new0d
(2) Crude flow phantom study.
(3) Orientation of the object and image study.

Date of Study: Oct 11, 97

10/11/97
flow phantom study
by David Zhu

Test sequence fgt1_new0d
FOV = 34

series 001: axial cut using spin echo

The following has FOV = 34 and A24

fgt1_new0d, default setting

series 002: pump mark flow measurement
002 0 0
003 0.5 40 ml/128
004 0.66 100 ml/128
005 0.8 100 ml/128
006 0.9 100 ml/128
007 1.0 120 ml/128
008 1.1 140 ml/128

Date of study: 10/12/97
Breast Sequence Study
Michael Buonocore and David Zhu

Subject Name: Joy Watts
Age: 72

Ask subject to have a better fit of the foam cast, tighter strap, and not to move, and to breath smoothly with stomach.

Procedures:
1. 2D Spin Echo axial cut (Series 1) FOV = 40

2. Fast Spin Echo (Series 2) FOV = 34

3. David's breast sequence coronal cut: fgt1_new0d, fov = 34

Seven TI cycles: br_TR = 2.7 s, br_T1 = 15 ms, ti_inc = 100 ms.
TI decreases from 615 to 15 ms.

Acquisition slice thickness: 3 mm
Selective slice thickness: 3 mm
Frequency direction: R/L for all locations

series 3: location: A18 R1 = 5, R2 = 14, TO = 185
series 4: location: A17 R1 = 5, R2 = 14, TO = 185
series 5: location: A13 R1 = 6, R2 = 15, TO = 185
series 6: location: A8 R1 = 6, R2 = 15, TO = 185
series 7: location: A28 R1 = 6, R2 = 15, TO = 185
series 8: location: A3 R1 = 6, R2 = 15, TO = 185
series 9: location: A23 R1 = 6, R2 = 15, TO = 185
series 10: location: P2 R1 = 6, R2 = 15, TO = 185
series 11: location: A38 R1 = 6, R2 = 15, TO = 185

Comments: The tumor is a breast above the nipple.
series: 12:  
location: P7  
R1 = 6 , R2 = 15 , TG = 185  

Comments:  
Tumor locates at left breast, 6 o'clock, about 1/3 in from the nipple.  

Prescan cannot set TG, manually set TG to 185 for all.  
R1 = 5 and R2 = 14 seem to be too low. Manually set most prescan to  
R1 = 6 and R2 = 15. Hope R1 = 5 and R2 = 14 do not have problem.  

(Procedure Updated on 11/15/97)  
Date of study: 11/16/97  
Breast Sequence Study  
by Michael Buonocore and David Zhu  

Type of study: phantom( ), subject ( x ) , Patient ( )  
Name: Kim Buonocore  
Age: 38  

Preparation:  
Ask subject to have a better fit of the foam cast, tighter strap,  
and not to move, and to breath smoothly with stomach.  

Procedures:  
1. 2D Spin Echo axial cut (Series 1)  
POV = 40  
I 84 to s 84 , loc: 29  . Scan thick = 5 mm, Inter = 1 mm.  
2. 2D Fast Spin Echo (Series 2)  
POV = 34  
P 7 to A 55 , loc: 12  , scan thick = 5 mm, inter = 1 mm.  
3. David's breast sequence coronal cut: fgt1_new12, fov = 34  

Seven TI cycles: br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 100 ms.  
TI decreases from 615 ms to 15 ms.  
Acquisition slice thickness: 3 mm, Selective slice thickness: 3mm  
Frequency direction: R/L for all locations  

series 3:  
location: P7  
R1 = 6 , R2 = 15 , TG = 153  
series 4:  
location: P1  
R1 = 6 , R2 = 15 , TG = 154  
series 5:  
location: A5  
R1 = 6 , R2 = 15 , TG = 153  
series 6:  
location: A11  
R1 = 6 , R2 = 15 , TG = 154  
series 7:  
location: A17  
R1 = 6 , R2 = 15 , TG = 160  
series 8:  
location: A23  
R1 = 6 , R2 = 15 , TG = 160  
series 9:  
location: A29  
R1 = 6 , R2 = 15 , TG = 160  
series 10:  
location: A35  
R1 = 6 , R2 = 15 , TG = 167  

Comments:  
Fibroadenoma is at left breast, about 1/3 in from nipple, about 3 o'clock  
position.  

(Procedure Updated on 11/19/97)  
Date of study: 11/19/97  
Breast Sequence Study  
by Michael Buonocore and David Zhu  

Type of study: phantom( ), normal subject ( ), subject w/ suspicious lesion (x)  
Name: Leslie Deard  
Age: 55  

Preparation:  
Ask subject to have a better fit of the foam cast, tighter strap,  
and not to move, and to breath smoothly with stomach.  

Procedures:  
1. 2D Spin Echo axial cut (Series 1)  
POV = 40  
I 84 to s 84 , loc: 29  . Scan thick = 5 mm, Inter = 1 mm.  
2. 2D Fast Spin Echo (Series 2)  
POV = 34  
P 7 to A 55 , loc: 12  , scan thick = 5 mm, inter = 1 mm.  
3. David's breast sequence coronal cut: fgt1_new12, fov = 34  

Seven TI cycles: br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 100 ms.  
TI decreases from 615 ms to 15 ms.  
Acquisition slice thickness: 3 mm, Selective slice thickness: 3mm  
Frequency direction: R/L for all locations  

series 3:  
location: A0  
R1 = 6 , R2 = 15 , TG = 159  
series 4:  
location: A4  
R1 = 6 , R2 = 15 , TG = 159  
series 5:  
location: A10  
R1 = 6 , R2 = 15 , TG = 160  
series 6:  
location: A13  
R1 = 6 , R2 = 15 , TG = 159  
series 7:  
location: A16  
R1 = 6 , R2 = 15 , TG = 160  
series 8:  
location: A19  
R1 = 6 , R2 = 15 , TG = 160  
series 9:  
location: A22  
R1 = 6 , R2 = 15 , TG = 160  
series 10:  
location: A25  
R1 = 6 , R2 = 15 , TG = 160  
series 11:  
location: A34  
R1 = 6 , R2 = 15 , TG = 160  
series 12:  
location: A40  
R1 = 6 , R2 = 15 , TG = 160  

Comments:  
Tumor at left breast, close to chest wall, lateral.
Procedure Updated on 11/19/97  
Date of study: 5/16b/98  
Breadth Sequence Study  
by David Zhu  
Type of study: potato phantom (x)  
Name: potato phantom  
Age: about 40  

Date of study: 5/16b/98  
Breadth Sequence Study  
by David Zhu  
Heat phantom  
Type of study: Heat phantom  

Procedures:  
1. 2D Spin Echo axial cut (Series 1)  
    FOV = 40  
    Seven TI cycles: br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 100 ms.  
    Acquisitoin slice thickness: 3 mm, Selective slice thickness: 3mm  
    Frequency direction: R/L for all locations  

2. David's breast sequence coronal cut: fgsl_new3d, fov = 34  
    Seven TI cycles: br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 100 ms.  
    Acquisitoin slice thickness: 3 mm, Selective slice thickness: 3mm  
    Frequency direction: R/L for all locations  

3. David's breast sequence coronal cut: fgsl_new3d, fov = 34  
    Modified the following parameters to have: ti_inc = 10ms, br_TI = 550 ms  
    Purpose: To test the validity of motion correction algorithm when the  
    the change of signal intensity is small.

Date of study: 5/16b/98  
Breadth Sequence Study  
by David Zhu  
Type of study: potato phantom (x)

Procedures:  
1. 2D Spin Echo axial cut (Series 1)  
    FOV = 40  
    Seven TI cycles: br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 100 ms.  
    Acquisitoin slice thickness: 3 mm, Selective slice thickness: 3mm  
    Frequency direction: R/L for all locations  

2. David's breast sequence coronal cut: fgsl_new3d, fov = 34  
    Seven TI cycles: br_TR = 2.7 s, br_TI = 15 ms, ti_inc = 100 ms.  
    Acquisitoin slice thickness: 3 mm, Selective slice thickness: 3mm  
    Frequency direction: R/L for all locations  

3. David's breast sequence coronal cut: fgsl_new3d, fov = 34  
    Modified the following parameters to have: ti_inc = 10ms, br_TI = 550 ms  
    Purpose: To test the validity of motion correction algorithm when the  
    the change of signal intensity is small.

(Date Procedure Updated on 11/19/97)  
Date of study: 4/13/98  
Breadth Sequence Study  
by Michael Buonocore and David Zhu  
Type of study: phantom (x), normal subject (x), subject w/ suspicious lesion (x)  
Name: Debra  
Age: about 40  

Preparation:
Ask subject to have a better fit of the foam cast, tighter strap, and not to move, and to breathe smoothly with stomach.

Procedures:
1. 2D Spin Echo axial cut (Series 1) FOV = 40 (named axial)
   I 84 to 6 84 , loc: 29 . Scan thick = 5 mm, Inter = 1 mm.
2. 2D Fast Spin Echo (Series 2) FOV = 34 (named higher)
   P 10 to A 74 , loc: 15 . Scan thick = 5 mm, Inter = 1 mm.
3. SPGR dynamic contrast study (scan 81 in protocol): (series 3 -> dynamic)
   Each scan sequence takes total 30 sec with scan time 25 sec, and press time 5 sec.
   Note: The contrast study was not that successfully since the injection was not given successfully in terms of time.
   Total of 13 scan sequences with a total images of 169.
4. 3D scan (scan 92 in protocol, 2'37") with flip angle 20 degree. (series 4)
   28 slices with the center at A33.6.
5. David's breast sequence coronal cut: fgtl_new3d, fov = 34
   Seven TI cycles: br_TR = 2.7 s, br_TI = 15 ms, br_inc = 100 ms.
   TI decreases from 615 ms to 15 ms.
   Acquisition slice thickness: 3 mm, Selective slice thickness: 3mm
   Frequency direction: R/L for all locations
   series 5: location: A25 R1 = 1 , R2 = 15 , TO = 183
   series 6: location:A14 R1 = , R2 = , TO =
   series 7: location:A26 R1 = , R2 = , TO =
   series 8: location:A26 R1 = , R2 = , TO =
   series 9: location:A26 R1 = 6 , R2 = 15 , TO =
   series 10: location:A26 R1 = 6 , R2 = 15 , TO =
   series 11: location:A26 R1 = 6 , R2 = 15 , TO =
6. 3D scan (scan 92 in protocol, 2'37") with flip angle 65 degree. (series 12 -> post_3d)
   28 slices with the center at A33.6.
7. A phantom study to check the coil (series 13)

Comments:
   The suspicious mass is located at the right breast about two o'clock.
   This subject later had a NEGATIVE biopsy result.

(Procedure Updated on 07/25/98)
Date of study: 7/27/98
Breast Sequence Study
by Michael Buonocore and David Zhu
Type of study: phantom(), normal subject (), subject w/ suspicious lesion (x)
Name: Eileen D. Morrison Age: 64
Preparation:
Ask subject to have a better fit of the foam cast, tighter strap, and not to move, and to breathe smoothly with stomach.

Procedures:
1. 2D Spin Echo axial cut (Series 1 or axial) FOV = 40
   I 84 to 6 84 , loc: 29 . Scan thick = 5 mm, Inter = 1 mm.
2. 2D Fast Spin Echo (Series 2 or higher) FOV = 34
   P 10 to A 74 , loc: 15 . Scan thick = 5 mm, Inter = 1 mm.
3. SPGR dynamic contrast study (scan 81 in protocol): (series 3 -> dynamic)
   Each scan sequence takes total 30 sec with scan time 25 sec, and press time 5 sec.
   Note: The contrast study was not that successfully since the injection was not given successfully in terms of time.
   Total of 13 scan sequences with a total images of 169.
4. 3D scan (scan 92 in protocol, 2'37") with flip angle 20 degree. (series 4)
   28 slices with the center at A33.6.
5. David's breast sequence coronal cut: fgtl_new3d, fov = 34
   Seven TI cycles: br_TR = 2.7 s, br_TI = 15 ms, br_inc = 100 ms.
   TI decreases from 615 ms to 15 ms.
   Acquisition slice thickness: 3 mm, Selective slice thickness: 3mm
   Frequency direction: R/L for all locations
   series 5: location: A25 R1 = 1 , R2 = 15 , TO = 183
   series 6: location:A14 R1 = , R2 = , TO =
   series 7: location:A26 R1 = , R2 = , TO =
   series 8: location:A26 R1 = , R2 = , TO =
   series 9: location:A26 R1 = 6 , R2 = 15 , TO =
   series 10: location:A26 R1 = 6 , R2 = 15 , TO =
   series 11: location:A26 R1 = 6 , R2 = 15 , TO =
6. 3D scan (scan 92 in protocol, 2'37") with flip angle 65 degree. (series 12 -> post_3d)
   28 slices with the center at A33.6.
7. A phantom study to check the coil (series 13)

Comments:
   The contrast agent can last for hours.
   The lesion is at the right side of the breast at about 9 o'clock.
   Problems: (1) One side of the cable was broken and led to that the one side did not receive the signal.
   Replacing the cable with a feed-first cable would be ok. The coil is fine.
   (2) Contrast agent has not been performed correctly.

(Procedure Updated on 07/25/98)
Date of study: 8/7/98
Breast Study with Contrast
by Michael Buonocore and David Zhu
Type of study: phantom(), normal subject (), subject w/ suspicious lesion (x)
Name: Marcia Drake Age: 37 Weight: 135
Preparation:
Ask subject to have a better fit of the foam cast, tighter strap, and not to move, and to breathe smoothly with stomach.

Procedures: (Protocol: # 64) (Study # 04817)
1. 2D Spin Echo axial cut (Series 1 -> axial) FOV = 40
   I 84 to 6 84 , loc: 29 . Scan thick = 5 mm, Inter = 1 mm.
2. 2D Fast Spin Echo (Series 2 -> higher) FOV = 34
   P 10 to A 74 , loc: 15 . Scan thick = 5 mm, Inter = 1 mm.
3. David's breast sequence coronal cut: fgtl_new3d, fov = 34
   Seven TI cycles: br_TR = 2.7 s, br_TI = 15 ms, br_inc = 100 ms.
   TI decreases from 615 ms to 15 ms.
   Acquisition slice thickness: 3 mm, Selective slice thickness: 3mm
   Frequency direction: R/L for all locations
   series 3: location: P10 R1 = 1 , R2 = 15 , TO = 146
   series 4: location: P4 R1 = , R2 = , TO =
   series 5: location: A1 R1 = 6 , R2 = 15 , TO = 150
<table>
<thead>
<tr>
<th>series</th>
<th>location</th>
<th>R1</th>
<th>R2</th>
<th>TG</th>
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<tbody>
<tr>
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<td>A14</td>
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<tr>
<td>8</td>
<td>A20</td>
<td>6</td>
<td>15</td>
<td>146</td>
</tr>
<tr>
<td>9</td>
<td>A25</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>10</td>
<td>A32</td>
<td>6</td>
<td>15</td>
<td>149</td>
</tr>
<tr>
<td>11</td>
<td>A38</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>12</td>
<td>A43</td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>13</td>
<td>A50</td>
<td>6</td>
<td>15</td>
<td>151</td>
</tr>
</tbody>
</table>

4. Pre-contrast 3D scan (scan #2 in protocol, 2.37°) (flip angle = 65) (series 14 -> pre_3d.a) (Center at A22.7 with 28 locs, 3 mm slice thickness)

5. More David's breast sequence coronal cut: fgtl_new3d, fov = 34

<table>
<thead>
<tr>
<th>series</th>
<th>location</th>
<th>R1</th>
<th>R2</th>
<th>TG</th>
</tr>
</thead>
<tbody>
<tr>
<td>15</td>
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<td>6</td>
<td>15</td>
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<td>16</td>
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<tr>
<td>17</td>
<td>A29</td>
<td>6</td>
<td>15</td>
<td>148</td>
</tr>
</tbody>
</table>

The patient is pulled out to set up contrast injection in the following scans:

6. Pre-contrast 3D scan (scan #2 in protocol, 2.37°) (flip angle = 65) (repeat the pre_contrast) (Center at A22.7 with 28 locs, 3 mm slice thickness)

7. SPGR dynamic contrast study (total 7 minutes) (scan #1 in protocol):
   (Series 19 -> dynamic)
   Each scan sequence takes total 10 sec with scan time 25 sec, with approximate interscan press time of 5 sec.
   (1) 3 scan sequences before injection
   (2) Begin injection and press SCAN to begin the 4th scan
   (3) Continue on the same fashion to finish the 14th scan.

8. Post-contrast 3D scan (scan #2 in protocol, 2.37°), same as #6 above.
   (Series 20 -> post_3d) (Center at A22.7 with 28 locs, 3 mm slice thickness)

Comments:
Patient claims that the tumor locates at 10 o'clock in the right breast.
The patient has small breasts.
Foam packs have been adjusted to avoid motion artifacts. Image displacement correction might need to consider.
Based on the analysis of the perfusion study, this patient appears to be malignant in the right breast.
Biopsy indicated that the lesion(s) was benign.
MEMORANDUM FOR Administrator, Defense Technical Information Center (DTIC-OCA), 8725 John J. Kingman Road, Fort Belvoir, VA 22060-6218

SUBJECT: Request Change in Distribution Statement

1. The U.S. Army Medical Research and Materiel Command has reexamined the need for the limitation assigned to technical reports written for this Command. Request the limited distribution statement for the enclosed accession numbers be changed to "Approved for public release; distribution unlimited." These reports should be released to the National Technical Information Service.

2. Point of contact for this request is Ms. Kristin Morrow at DSN 343-7327 or by e-mail at Kristin.Morrow@det.amedd.army.mil.

FOR THE COMMANDER:

[Signature]

Encl