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Impact of Particle Sizes on MRI Signal Relaxation in Phantoms for

Assessment of Hepatic Steatosis and Iron Overload

by

Sarah Caroline Brasher

A Thesis

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Preface

Chapter III, a comprehensive review on MRI phantom literature, is composed of the following journal article: "Quantitative MRI of Diffuse Liver Diseases: Techniques and Tissue-Mimicking Phantoms", written for submission to *Magnetic Resonance Materials in Physics, Biology and Medicine (MAGMA)*.

Abstract

Magnetic Resonance Imaging (MRI) is emerging as a powerful tool to non-invasively evaluate diffuse liver diseases such as hepatic steatosis and iron overload. To test new MRI techniques, phantom studies are utilized in place of patients but often do not consider the microscopic interactions of particles suspended in the media which may cause a notable difference in the signal. Hence, this study investigates the impact of differing particle sizes on the magnetic resonance signal using phantoms. To accomplish this, steatosis phantoms were created using two different mixing methods to control droplet size and while combination ironfat phantoms featuring iron particles of differing diameters were used to emulate hepatic iron overload. Signal behavior from both sets of phantoms were resolved using linear calibrations to determine values from two known biomarkers, fat fraction and R2*, for steatosis and iron overload, respectively. Overall, evidence showing that particle size impacts the signal to a significant degree remains inconclusive, but fitting model performance in biomarker quantification varied. This study demonstrates different sequencing and post-processing assessments are critical for the analysis of sensitive biomarkers such as R2* and FF.

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Abbreviations

ADC	Apparent Diffusion Constant
ARMA	Autoregressive Moving Average
B-Gel	Bovine Skin Powder
BNF	Bionized NanoFerrite
CAG	Carrageenan-strengthened Agarose/Gadolinium Gel
CPMG	Carr-Purcell-Meiboom-Gill
CRN	Clinical Research Network
СТ	Computed Tomography
DSPIO	Dextran-coated Superparamagnetic Iron Oxide
DWI	Diffusion-Weighted Imaging
ECM	Extracellular Matrix
FDA	Federal Drug Administration
FID	Free Induction Decay
FF	Fat Fraction
FFT	Fast Fourier Transform
FOV	Field of View
FS	Fat Suppression
DWI	Diffusion Weighted Imaging
G	Shear Modulus
G'	Storage Modulus
G _x	Frequency Encoding Gradient
Gy	Phase Encoding Gradient
Gz	Slice Selection Gradient
GRE	Gradient Echo
H&E	Hematoxylin and Eosin
HIC	Hepatic Iron Content
IR-Snapshot	Inversion Recovery Snapshot Fast Low Angle Shot
IR-UTE	Adiabatic Inversion Recovery Prepared Ultrashort Echo Time
ISMRM	International Society for Magnetic Resonance in Medicine
IVIM	Intravoxel Incoherent Motion
mGRE	Multi-Echo Gradient Echo
MOLLI	Modified Look-Locker Inversion Recovery
MR	Magnetic Resonance
MRE	Magnetic Resonance Elastography
MRI	Magnetic Resonance Imaging

MRS	Magnetic Resonance Spectroscopy
NAFLD	Non-Alcoholic Fatty Liver Disease
NASH	Non-Alcoholic Steatohepatitis
NIST	National Institute of Standards and Technology
NLI	Non-Linear Inversion
NLSO	Non-Linear Least Squares
NMR	Nuclear Magnetic Resonance
QIBA	Quantitative Imaging Biomarkers Alliance
QSM	Quantitative Susceptibility Mapping
PAA	Polyacrylamide
PVA	Polyvinyl Alcohol
R2	Rate of Transverse Relaxation
R2*	Rate of Transverse Relaxation with Inhomogeneities
RF	Radio Frequency
ROI	Region of Interest
RSNA	Radiological Society of North America
SASHA	Saturation Recovery Single-Shot Acquisition
S	Stiffness
SE	Spin Echo
SE-EPI	Spin Echo Echo Planar Imaging
SE-IR	Spin Echo Inversion Recovery
SEM	Scanning Electron Microscopy
SIR	Signal Intensity Ratio
SNR	Signal to Noise Ratio
SPIO	Superparamagnetic Iron Oxide
SS-EPI	Single-Shot Echo Planar Imaging
T1	Longitudinal Relaxation
T2	Transverse Relaxation
T2*	Transverse Relaxation with Inhomogeneities
TE	Echo Time
TR	Repetition Time
TSE	Turbo Spin Echo
USPIO	Ultra-small Particles of Iron Oxide
UTE	Ultrashort Echo Time
VFA	Variable Flip Angle

Chapter I

Introduction

Motivation for Study

Iron is an essential mineral in many metabolic processes, particularly for the synthesis of the oxygen transport proteins, hemoglobin and myoglobin [5]. In the case of an imbalance of iron, homeostatic mechanisms can only regulate iron by limiting absorption during digestion as the human body has no natural physiological processes for excreting iron [6]. Thus, excess iron can build up in the liver, the primary organ for iron storage, under conditions of hereditary hemochromatosis or transfusional hemosiderosis [7]. In addition to iron storage, the liver plays a vital role in lipid metabolism [8]. Hepatic steatosis, the abnormal deposition of lipids in the liver, is a hallmark characteristic of many diffuse liver diseases such as viral hepatitis and alcoholic or nonalcoholic fatty liver disease [9,10]. Co-occurrence of hepatic steatosis and iron overload is well documented [11], and the disease progression of either or both diffuse liver diseases can lead to fibrosis, cirrhosis, and ultimately liver failure [10].

Liver biopsy is the current gold standard for evaluating hepatic steatosis and iron overload, but it has a few major limitations. Firstly, steatosis and iron overload can be heterogeneously distributed [12,13], and the small sampling size of the biopsy specimen lacks information on the spatial distribution of the liver, thus causing high sampling variability in quantitative measurements [7,14]. Biopsy is a painful and invasive procedure with a risk of bleeding and infection, leading to suboptimal patient compliance [15,16]. Additionally, frequent assessments of diffuse liver disease are needed to guide treatments, requiring multiple biopsies. Because of these drawbacks, MRI has arisen as a non-invasive alternative to liver biopsy and is

becoming increasingly recognized in the clinical sector to monitor hepatic steatosis and iron overload [17,18].

Quantitative MRI techniques for evaluating hepatic steatosis and iron overload have been developed but have limitations on the range of fat and iron deposition they can measure. To advance these techniques, extensive testing must occur on specified pulse sequences and scanning hardware. MRI examinations routinely take approximately 20-60 minutes per session [19], so acquiring volunteers to test new MRI sequences can be challenging especially across multi-site studies. Hence, researchers utilize imaging phantoms, specialized standards equipped to emulate a certain disease, to develop new MRI techniques. These phantoms contain materials that manipulate the MRI signal to produce characteristic effects that would be seen under certain pathologies, therefore targeting different MRI biomarkers. However, diffuse liver phantoms in the literature display little to no standardization in materials, volume, protocol, and even targeted biomarkers for a particular disease.

For this thesis, a question of particular interest is that concerning the microscopic interactions of iron and fat particles in the phantom media on the MR signal. In vivo, hepatocytes exist in a range of sizes between 20-25 μ m in diameter, while steatotic lipid globules grow no larger than 15 μ m. Additionally, iron is usually stored in ferritin molecules that cluster in sizes of 50-70nm. However, under severe iron overload, hemosiderin is the primary iron storage molecule and clusters to sizes of 0.1-2 μ m. In phantom literature, these physiological sizes are often not considered in the particles used to emulate diffuse liver disease and may show a much larger difference in magnitude than what is seen in vivo. This is a concern as it is theorized that substances such as iron will preferentially affect the signal produced by water over fat due to geographical limitations and solubility. Thus, adjusting the sizes of these particles may allow for

more or less interactions between iron, fat, and water, all of which affect the MR signal significantly.

Hence, there is a gap in our understanding of the impact of particle size on MR signal behavior. The following sections will establish fundamental MR techniques related to diffuse liver disease before culminating into a study investigating the impact of iron and fat particle sizes on the MRI signal. A foundational segment on MRI basics, sampling, and pulse sequences will first be discussed in Chapter II before moving onto a comprehensive literature review on hepatic phantoms in Chapter III. Following this, an analysis of the microscopic interactions in phantom media will be conducted in Chapter IV by varying sizes of fat and iron particles in phantoms emulating hepatic steatosis and iron overload on a scale relative to molecules seen in in vivo liver tissues. These phantoms will use two biomarkers that encompass clinical ranges: fat fraction (FF) for hepatic steatosis and R2* for hepatic iron overload. Additionally, particle size will be validated using light microscopy and scanning electron microscopy (SEM), and various signal fitting models will be tested for accurate and precise quantification of R2* and FF. Using these methods, an investigation on the impact of particle size on MR signal behavior will be performed. Finally, a discussion on potential expansions and adaptations to this study will be elaborated on in Chapter V.

Chapter II

MRI Physics

Basics of MRI Physics

While MRI is favored in most clinical environments due to its ability to provide spatial information, all MRI scanners operate based on the physical phenomenon of nuclear magnetic resonance (NMR). Since its development by Purcell [20] and Bloch [21] in 1945, NMR has been used as an analytical technique for elucidating chemical structures in solids by manipulating specific atomic species using weak oscillating magnetic fields. Only nuclei which have an odd number of neutrons and/or protons possess net spin and are considered NMR active and can display this phenomenon [22]. Many NMR-active species exist, including ¹H, ¹³C, ¹⁵N, and ³¹P, all present in living tissues [23-25]. However, ¹H is utilized in clinical MRI scanners due to its relative biological abundance [26].

The NMR phenomenon begins with the presence of a strong, static magnetic field aligning NMR-active nuclei with or against the field [26]. This will cause the nuclei to precess at a unique Larmor frequency, thereby generating a spinning magnetic moment that can be represented as a net magnetization vector in the direction of the static magnetic field [26,27]. The equation for a given Larmor frequency can be seen below,

$$\omega_0 = \gamma B_0$$

where ω_0 is the Larmor frequency in MHz, γ is the gyromagnetic ratio of the nuclei, and B₀ is the static magnetic field strength in Tesla [27]. A coil in the scanner will then generate a current to produce a radiofrequency (RF) pulse, therefore inducing an oscillating magnetic field. This RF pulse will tip the net magnetization vector of the nuclei from the z-axis to the xy-plane with a flip angle depending on the strength and duration of the RF pulse. A secondary RF receiver coil will

detect the induced current caused by a change in the net magnetic field via electromagnetic induction. Inevitably, the spins of the nuclei will become dephased due to the interference of random thermodynamic interactions with neighboring nuclei, and the signal induced by the magnetic moments of each nucleus will decay as more return to the direction of the static magnetic field [28]. This phenomenon is known as relaxation, where the recovery of the longitudinal magnetization vector is termed T1 relaxation, and the loss of the transverse magnetization vector is termed T2 relaxation which can be shown graphically as a free induction decay (FID) [29]. Additionally, local field inhomogeneities can accelerate the signal decay in an effect known as T2* relaxation [30]. The preceding concepts are illustrated in Figure 1.

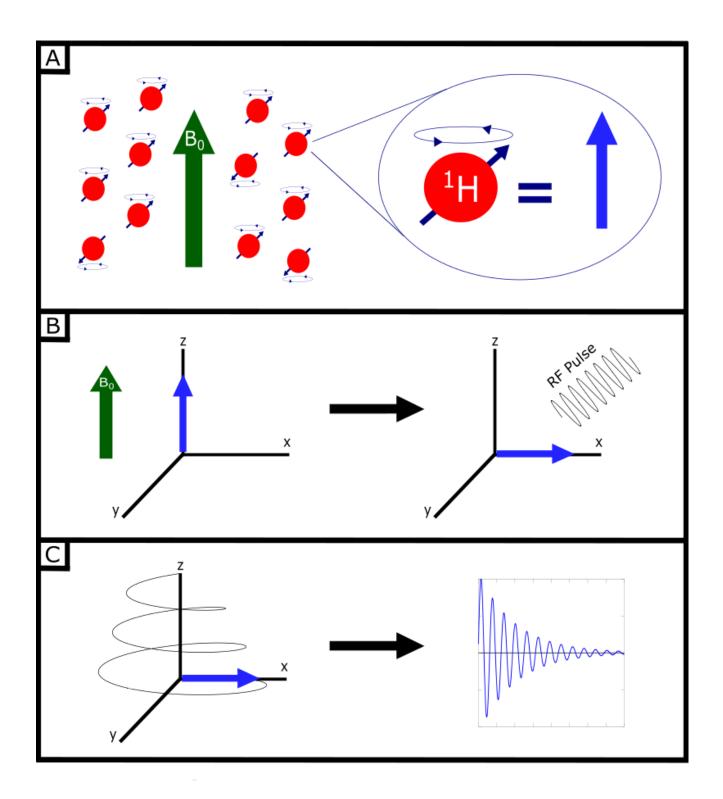


Figure 1. Overview of basic MRI physics. (A) Precession of NMR-active nuclei align toward the static magnetic field, B_0 , and can be represented as a net magnetization vector. (B) The net magnetization vector is tipped from the direction of B_0 on the z-axis to the xy-plane by an RF pulse. (C) Relaxation of the magnetization vector results in an FID signal as the signal is lost in the transverse plane.

Cartesian and Radial Sampling

Using the Fourier transform, any oscillating signal can be broken up into a series of sinusoidal waveforms [31]. Hence, an FID produced via MRI can be translated into the frequency domain using the Fourier transform and represented in two dimensions through k-space [32,33]. In this visual representation of frequencies, the contrast information of the signal primarily resides in the center of k-space while the high-resolution spatial information resides in the periphery [33]. This information is filled into k-space by manipulating MRI gradients that subtly change the field strength [33]. Specifically, phase encoding (G_y) and frequency encoding (G_x) gradients store information on the y and x-axis, respectively. Performing an inverse Fourier transform can therefore recover the spatial MRI image from the frequency domain [34]. However, how one samples the acquisition through k-space plays an important role in MRI image acquisition and can create pronounced impacts on the final MRI image.

The standard method of sampling k-space data in most clinical MRI sequences is through Cartesian (rectilinear) sampling [34]. As seen on Figure 2, this method involves iteratively recording regularly spaced parallel data elements from left to right using independent phase and frequency encoding gradients until the entirety of k-space is filled [34]. This allows for the matrix of data elements to be placed into an array for the efficient processing through fast Fourier transform (FFT) to create MRI images [34]. Alternatively, non-Cartesian methods such as radial sampling have been gaining popularity for their unique benefits. Here, radial sampling involves the generation of center-out vectors in k-space formed by the utilization of simultaneous phase and frequency encoding gradients [1]. However, these vectors contain data elements that do not neatly settle into a rectangular matrix, so they must undergo "re-gridding" into a Cartesian matrix format before undergoing FFT [1,34].

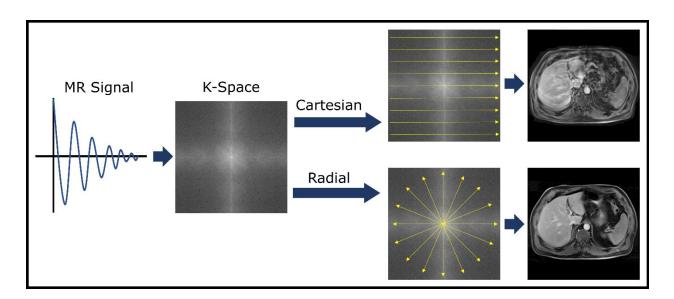


Figure 2. Schematic of Cartesian and radial sampling of k-space. Performing Fourier transform on a spatial 2D signal provides data that can be represented in the frequency domain through K-space. Phase and frequency encoding can be manipulated to perform cartesian (left to right) and radial (center-out) acquisitions followed by inverse Fourier transform to recreate the spatial image. The images provided are of a free-breathing abdominal exam with motion artifacts [1].

In today's clinical environment, Cartesian sampling remains the most dominant modality, but other approaches are becoming more popular in light of recent research developments. The main advantage of Cartesian sampling is that it can be simply transformed into an image with the FFT without additional processing. However, radial sampling is less sensitive to motion artifacts and has better contrast due to the oversampling of the center of k-space [1]. This redundancy allows for the correction of any movement in the signal as more is sampled at the center [1]. Nonetheless, radial sampling is prone to distinctive streaking artifacts that emerge from object motion or bright intensities in the image, but the streaks are relatively easy to identify and usually do not confound a diagnosis if the radiologists/technicians are trained prior [1,35]. Radial sampling also utilizes simultaneous phase and frequency encoding gradients without increasing the scan time, unlike Cartesian methodologies that use the gradients sequentially [1]. However, radial trajectories have less efficient coverage of k-space in the periphery, causing low resolution and image blurring [1].

GRE and UTE Sequences

In MRI, the physical properties of an FID can be manipulated by adjusting a number of parameters in the internal gradients of the scanner including echo time (TE), repetition time (TR), flip angle, echo train length, and field of view (FOV), among others [30,36]. By utilizing presets of these parameters, unique pulse sequences are generated to highlight specific physical phenomena at the discretion of the user. For diffuse liver diseases such as iron overload and steatosis, sequences that utilize the properties of chemical shift, i.e., the difference in resonance frequencies between nuclei, and R2* relaxometry are needed to capture relevant properties of the signal [37].

One of the most fundamental sequences featured in MRI is the gradient echo (GRE) which involves two rudimentary steps. First, an external dephasing gradient is applied in the frequency encoding step to squelch the FID by slightly altering the resonance frequencies of the sample [38]. After this, a rephasing gradient is implemented with the same magnitude as the before gradient, but in an opposite polarity [38]. This allows the FID to reform as a gradient echo that retains both the T2 and T2* information of the signal, hence GRE sequences can be used to assess pathologies that induce severe field inhomogeneities such as iron overload [30,39]. Additionally, a multi-echo gradient echo (mGRE) sequence shown in Figure 3 will allow for more echoes to be produced after the initial one through repeated applications of the rephasing and dephasing gradients, thus it can be utilized in chemical shift imaging in which multiple TEs are needed to measure "in-phase" and "out-of-phase" fat-water signals [40]. However, GRE sequences do encounter a few problems. Because the T2* information is retained, signal loss is much more rapid with GRE sequences compared to other sequence types such as spin-echo [39]. Also, because GRE sequences utilize Cartesian sampling methods, motion artifacts such as

breathing may become an issue in abdominal and pediatric imaging [41]. Hence, alternative sequences have been explored to better assess diseases such as iron overload and steatosis [35,42].

While GRE sequences are the traditional method of evaluating iron overload, ultrashort echo time (UTE) sequences have shown promising results in quantifying extreme concentrations of iron that GRE sequences inadequately assessed [35,42]. The structure of a UTE sequence optimizes the shortest possible delays between signal excitation and acquisition by utilizing TEs of ≤ 0.1 ms [35]. Compared to the shortest possible TE of GRE, 1.0ms, a tenfold difference in TE allows UTE to sample the signal much earlier and can measure very short-lived signals such as those produced under high iron conditions [35]. This is accomplished by a few factors. First, UTE sequences utilize radial sampling of k-space which allows for simultaneous phase and frequency encoding gradients to reduce scan time while also reducing motion artifacts that might be seen in abdominal imaging [35,43]. As seen on Figure 3, UTE sequences also use "Half RF" pulses, but this requires two acquisitions with inverted slice selection gradients to obtain the entirety of the signal information [35,43]. By utilizing these parameters in tandem, the shortest possible TEs of ~0.1-0.02 ms are obtained in UTE sequences [35,43]. UTE sequences do have a few drawbacks, however. The utilization of half-pulse excitation makes the sequence sensitive to confounding out-of-slice signals due to eddy currents or gradient delays, all which may impede quantification of signal biomarkers [35]. Additionally, UTE sequences have only been validated in magnitude-based data, so quantification methods that utilize complex-based biomarkers require additional verification against biopsy data [42,44].

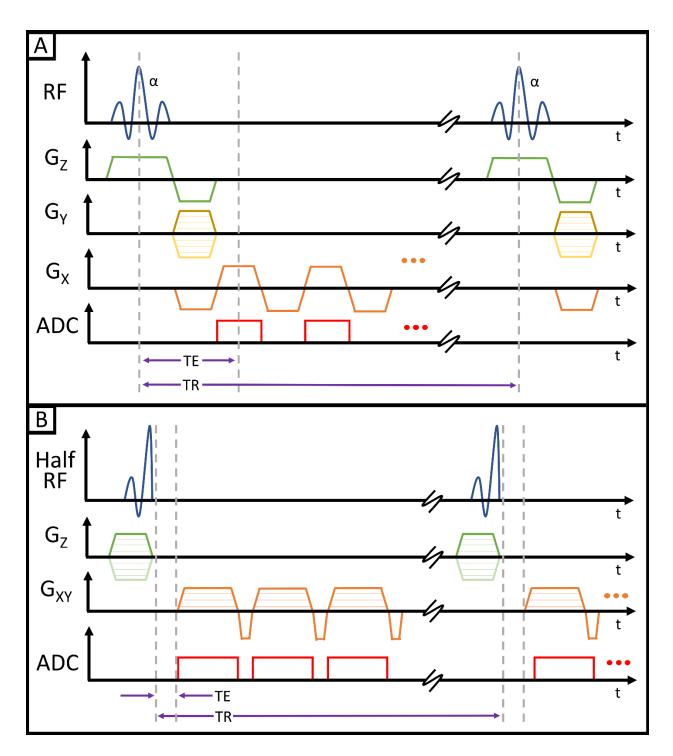


Figure 3. Pulse sequence diagrams of GRE and UTE sequences. (A) A mGRE (multi-echo gradient echo) sequence is shown with separate slice selection (G_z), phase encoding (G_y), and frequency encoding (G_x) gradients, and signal acquisition occurs during the rephasing gradient of the frequency encoding step. (B) A UTE (ultra-short echo time) sequence is shown with a half RF pulse along with an independent slice selection gradient (G_z) and combined phase and frequency encoding gradients (G_{xy}). The acquisition occurs only during the positive G_{xy} to prevent phase encoding errors.

Chapter III

Quantitative MRI of Diffuse Liver Diseases: Techniques and Tissue-Mimicking Phantoms

Introduction

Magnetic resonance imaging (MRI) has evolved as a noninvasive alternative to biopsy for the assessment of diffuse liver diseases, primarily hepatic iron overload, steatosis, and fibrosis [17,18,45]. Hepatic iron overload is a manifestation of hereditary hemochromatosis and transfusional hemosiderosis and is also common in some chronic hepatopathies [46]. Likewise, hepatic steatosis is the abnormal accumulation of fat in the liver and is a hallmark feature of nonalcoholic fatty liver disease (NAFLD), the most common diffuse liver disease that currently affects about 20%-30% of the U.S. population [47]. The co-occurrence of hepatic fat and iron overload, referred to as combined storage disease, is also being increasingly recognized [10,11,48]. Hepatic fibrosis is another common outcome of various liver pathologies, including hepatitis, NAFLD, and iron overload, and has a high prevalence of up to 25% in the general population [2,49,50]. The coexistence of fat, iron overload, and/or fibrosis with any pre-existing liver disease can accelerate disease progression. As early stages of these diseases are reversible conditions, techniques that can provide early diagnosis, allow disease monitoring, and guide therapeutic intervention are crucial for better prognosis.

Over the last two decades, several technical developments in quantitative MRI techniques have emerged that enable more precise and frequent monitoring of these diffuse liver pathologies to guide treatment and prevent disease progression and long-term complications. For testing, validating, and standardizing quantitative MRI methods across sites, vendors, and field strengths, phantoms are used as stand-ins to human tissue to mimic various liver pathologies. Quantitative liver phantoms also play an essential role for quality assurance in multi-center clinical trials and drug discovery studies that rely on quantitative imaging biomarkers as primary endpoints [51-53]. Organizations such as Radiological Society of North America (RSNA) Quantitative Imaging Biomarkers Alliance (QIBA) have laid out guidelines in bringing quantitative liver imaging to the clinical sector by collaborating with researchers and healthcare professionals to perform multi-site and multi-vendor validation studies on some promising MR biomarkers using phantoms [54-57]. However, currently, there is no standardization in the preparation of MRI phantoms for emulating diffuse liver diseases, as different studies utilize various types and sizes of iron and fat particles to mimic iron overload and steatosis respectively, different materials for mimicking fibrosis, and different matrices like water, agarose, or gels [58-64]. Further, with MRI, the co-occurrence of liver iron, fat, and fibrosis can confound or interfere with the quantification of each other. To address these challenges, novel MRI acquisition methods and multi-spectral signal models are being developed for accurate and simultaneous assessment of these diffuse liver diseases [44,65-67]. In addition, recent attempts include creating combination phantoms to mimic coexisting liver pathologies of iron overload, steatosis, and fibrosis observed in vivo for validation and quality assurance of MRI techniques for quantification of diffuse liver diseases [68]. However, simply combining different materials for each pathology to create phantoms mimicking concurrent diffuse liver diseases do not mimic the MRI signal characteristics seen in tissues [68].

The purpose of this study is, therefore, to provide a comprehensive review of the published MRI phantom studies on diffuse liver pathologies of iron overload, steatosis, and fibrosis by first describing the disease morphology and discussing the MRI techniques and biomarkers for each pathology and then examining the phantom materials chosen for mimicking these diffuse liver diseases. In this study, a systematic review is performed using PubMed as the primary database to identify and extract information from publications that constructed MR liver phantoms for mimicking iron overload, steatosis, fibrosis, and the combination of these pathologies. The key words used to search for these publications are phantom, liver, iron overload, steatosis, fibrosis, and MRI.

The following sections describe the desired characteristics and matrix materials of liver phantoms and briefly cover the in vivo morphology and quantitative MRI techniques for each pathology, followed by reviewing the phantom materials used to emulate each disease as well as elucidating their benefits and limitations.

Liver Phantoms

The design requirements of any quantitative tissue-mimicking MRI phantom are typically: (1) characteristic MRI signal behavior and relaxation times that accurately emulate biomarkers of disease while considering field strength dependencies, (2) cover relevant clinical ranges of the disease, (3) multi-site, multi-vendor, and multi-platform reproducibility for standardization, (4) stability over long periods of time, and (5) ease of fabrication and handling [58,69-72]. These tissue-mimicking phantoms can be constructed using a variety of materials to achieve these characteristics. However, phantoms emulating diffuse liver diseases have particular requirements that may limit the materials chosen, and these specifications are described below in the appropriate phantom material sections. Nevertheless, it must be understood that tissue-mimicking phantoms cannot simulate every facet of a tissue, but the phantoms must mimic the MRI signal characteristics observed in vivo of these diffuse liver pathologies.

Matrix Materials

Matrix mediums are a necessary requirement in phantoms for stabilizing the contents used to mimic liver pathologies. The most widely used matrix mediums for making phantoms are water, agar, and agarose. Water is risk-free and easy to handle, but it has some disadvantages as a phantom medium. Water requires a 10-minute settling time to allow suspensions to stabilize and is more susceptible to vibrational effects that may cause motion artifacts [58]. Additionally, water has similar T1 and T2 relaxation times, and therefore, it does not mimic the behavior of human liver tissues in which T1 is much longer than T2 [73-75]. Alternatively, materials such as agarose, carrageenan, polyvinyl alcohol (PVA), polyacrylamide (PAA), and gelatin create a jelly-like medium that will hold the phantom contents in suspension, thereby reducing boundary artifacts and avoiding flow issues [58,76]. Both agarose and agar gel have T2 relaxation times that are similar to human tissue (~30-200 ms) depending on the chosen concentration [58,77]. However, the T1 relaxation times are not comparable to human tissues (~300-1500 ms) but can be adjusted with paramagnetic additives such as CuSO₄, NiCl₂, MnCl₂, or GdCl₃ to reduce T1 [58,77]. An alternative polysaccharide that is utilized in MRI phantoms is carrageenan gel, a more flexible and resistant substance that can be molded into unique shapes, but use of high concentrations can cause the T2 relaxation times to be longer than human tissues [76,78]. Like many polysaccharides, agar and agarose need thermal treatments (80-100°C) during the preparation process, so caution must be taken when using some phantom contents that have restricted temperature ranges [58]. Conversely, PVA-based phantoms benefit from greater longevity and structural stability than most matrix materials, but the need for freezing and thawing cycles to prepare the phantom may pose similar issues [76,78].

While there is no standard guideline to define an acceptable shelf life of a phantom, the phantom stability is greatly affected by the matrix material chosen. Some materials such as agarose and silicone are known to be mechanically stable over long periods of time, assuming no extreme fluctuations in temperature and humidity in storage, while other materials such as PAA are only stable for a few months. [76]. Further, gel phantoms are prone to microbial growth and often use sodium azide or sodium benzoate as preservatives to retard mold formation and increase shelf life [58,61,68,79]. To mimic native tissues, some studies also used homogenized animal liver tissues as a matrix medium instead; however, special handling and storage of these tissue phantoms may be a concern [18,80,81].

Iron Overload

Pathophysiology & Morphology

Hepatic iron overload is a histological hallmark of primary (hereditary) or secondary hemochromatosis [7,82]. Hereditary hemochromatosis is caused by a defect in the genes that control intestinal iron absorption from food, primarily the HFE gene, and affected patients absorb iron at 5–10 times the normal rate (up to 10mg/day) [46,83]. Secondary hemochromatosis is usually the result of another disease or condition that causes iron overload, e.g., severe chronic hemolysis requiring chronic blood transfusions. Because the body has no physiologic mechanism for clearing excess iron, the extra iron accumulates in organs, primarily the liver, leading to hepatic iron overload. As the liver is a central organ in regulating iron homeostasis, iron is also one of the most common parenchymal deposits in other diffuse liver diseases like NAFLD apart from hemochromatosis [84]. The degree of liver dysfunction is directly dependent on the amount of hepatic iron deposition and progressive iron accumulation eventually leads to hepatomegaly, fibrosis, and finally, cirrhosis and liver failure [85-89].

The liver stores excess iron in hepatocytes and reticuloendothelial Kupffer cells, as ferritin and hemosiderin molecules. Ferritin is the principal iron storage molecule found in cells and is readily available to meet physiologic needs. Ferritin molecules in clusters have a size of 50-70 nm and are not visible individually on Perls iron stain, however, they can be seen as a diffuse blue blush in the cytoplasm of hepatocytes on light microscopy [90-92]. If ferritin levels are persistently elevated, hemosiderin deposits can develop as conglomerates of clumped ferritin particles. Hemosiderin appears as granular, golden brown cytoplasmic deposits on hematoxylin and eosin (H&E) stain and as blue deposits on Perls iron stain as shown in Figure 1. During iron overload, aggregates of hemosiderin molecules are deposited in body tissues and range in size from 0.1 to 2 μ m, eventually causing tissue damage and dysregulation of function [93]. Further, with an increase in iron overload severity, iron is deposited as clusters, and the deposition among hepatocytes are not uniform [93].

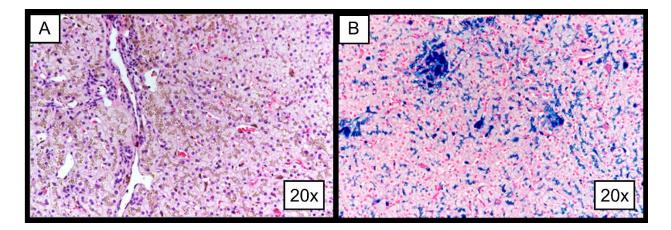


Figure 4. Hemosiderin deposits shown in the liver biopsy samples using hematoxylin & eosin (H&E) stain (A) and Perls Prussian blue iron stain (B) at 20x in a severely iron overloaded patient with biopsy hepatic iron content of 16.5 mg Fe/g. Iron deposits are seen as brown granules in the cytoplasm of the Kupffer cells and hepatocytes on the H&E stain and as blue pigment on Perls iron stain.

The amount of iron in the liver, or hepatic iron content (HIC), is a necessary metric for detecting and staging iron overload as HIC significantly correlates with total body iron stores [94]. The current gold standard to quantify HIC directly is by atomic absorption spectrophotometry on liver biopsy specimens, which measures HIC in milligram of iron per gram of dry liver tissue weight (mg Fe/g dry tissue wt.) [95]. Alternatively, iron deposition in biopsy specimens can also

be evaluated histologically on a semi-quantitative 5-point grading scale based on Prussian Blue staining of iron granules [46].

Quantitative MR Techniques

R2/R2 Relaxometry*

Currently, MRI techniques based on transverse relaxometry (R2/R2*) are considered robust and clinically accepted methods for non-invasively assessing HIC [96]. Iron in the form of ferritin and hemosiderin has paramagnetic properties, causing faster signal decay and increasing the R2 and R2* relaxation rates. R2 can be measured using a series of single spin-echoes, each acquired after a separate excitation [97] or with a train of spin-echoes acquired after a single excitation using a Carr-Purcell-Meiboom-Gill sequence [98-100]. St. Pierre et al. developed a R2-HIC calibration method that is commercially available as FerriScan (Resonance Health, Ltd, Claremont, Australia) for HIC determination [97]. The R2-HIC relationship is however curvilinear which saturates at high HIC, hence making the method less sensitive to severe iron overload.

On the other hand, R2* is measured using a multi-echo gradient-echo (GRE) sequence and fitting a mono-exponential signal model [2,101,102]. The R2*-HIC relationship is linear for the entire pathophysiological range of HIC making R2* measurements desirable for clinical assessments [103,104]. However, for extreme cases of iron overload (HIC > 30 mg Fe/g dry tissue wt. at 1.5T or HIC > 15 mg Fe/g at 3T), the R2* quantification with GRE sequences is limited by the shortest possible echo time (TE) of about 1 ms. To overcome this issue, ultrashort echo-time (UTE) techniques have been developed to improve the dynamic range of R2* relaxometry and have demonstrated high R2* accuracy even for severe iron overload [35,42,105]. One major limitation with R2* techniques is that the presence of fat (steatosis) introduces oscillations in the signal decay and confounds R2* measurements. To overcome this, multi-spectral signal modeling

techniques have been proposed for accounting for both the spectral complexity of fat and R2* decay. These techniques are discussed in more detail in the MR Techniques section under steatosis below.

Quantitative Susceptibility Mapping

Although relaxometry based methods are robust, they are confounded by other pathologies such as fibrosis, thus causing bias in HIC estimations [106,107]. Alternatively, quantitative susceptibility mapping (QSM) techniques can be used to estimate HIC by quantifying magnetic susceptibility, an intrinsic tissue property that is highly sensitive and linearly related to iron content, and is not affected by cellularity changes such as fibrosis [108]. Iron in the form of ferritin and hemosiderin induces a local susceptibility and affects the measured phase on GRE images. The local field map is then extracted from the phase data, and the susceptibility map is generated from the local field map by using a dipole inversion algorithm [109-112]. In contrast to the brain, QSM estimation in the abdomen must account for signals from multiple species (i.e., fat and water) and quantify a relatively wider range of iron contents (5-10 times greater iron deposition in liver iron overload than in brain). Recent studies have incorporated multi-spectral fat-water models to produce a fat-corrected field map estimation and used that in conjunction with QSM processing to generate susceptibility maps and have demonstrated a linear correlation between HIC and susceptibility values [113-117]. However, QSM is still limited to research due to both a lack of standardization of the technique and a not well-established relationship between susceptibility values and HIC [17]. Hence, it is not yet commercially available on MRI scanners for clinical HIC diagnosis [17].

Iron Overload Phantoms

The desired properties of an iron overload phantom are to maintain the characteristic relationship between MRI biomarkers (R2/R2*, susceptibility) and iron concentration, match the clinical ranges of MRI biomarkers in conditions of hepatic iron overload (R2*: ~40-1000 s⁻¹ at 1.5T, susceptibility: ~0.2–6 ppm) [113,115-118] and preserve the size and distribution of iron particles seen in liver tissues throughout the phantom. MRI studies that constructed phantoms to emulate hepatic iron overload using a variety of paramagnetic materials are summarized in Table 1. Of the studies reviewed, the most common materials used for making phantoms mimicking iron overload are iron nanoparticles. Iron oxide nanoparticles exhibit varying levels of paramagnetic and superparamagnetic behavior depending on the oxidation state of iron (ferrous vs. ferric) and particle size, in turn altering the R2* relaxation [119,120]. Ferrous iron, such as that of iron sulfate, can be added to emulate the R2* range [51-434 s⁻¹] of iron content in clinical hepatic iron overload [121]. On the other hand, ferric iron molecules exhibit stronger paramagnetic behavior and have greater susceptibility than ferrous iron [120]. A recent study by Alústiza et al. compared iron oxide nanoparticles with different oxidation states and found that the decrease in MRI signal for ferric iron (FeCl₃) was more linearly correlated with the increase in iron concentration than compared to ferrous iron ((NH₄)₂Fe(SO₄)₂), therefore more closely emulating the linear relationship between R2* and iron content observed in hepatic iron overload [64].

Citation PMID	Phantom Materials	MR Technique (Field Strength)	R2*(s ⁻¹)/QSM (ppm) Range
St. Pierre et al. 2005 15256427 [97]	MnCl ₂ (0.1-3.2 mM) Matrix: Water	Multi-echo SE	10-280 s ⁻¹ (R2)
Szurowska et al. 2010 20105330 [80]	FeCl ₃ Matrix: Pork liver	GRE SIR (1.5T)	n/a
Chandarana et al. 2012 22997371 [122]	Ferumoxytol (Feraheme®, 1/2000-1/500) Matrix: Deionized Water	Multi-echo GRE (1.5T)	25-150 s ⁻¹
Tsai et al. 2014 25120622 [123]	Fe(OH) ₃ (0.1-25 mg/mL) Matrix: Pork liver	Multi-echo GRE (1.5T)	80-400 s ⁻¹
Alústiza et al. 2015 25874207 [64]	FeCl ₃ (0.05-4 mg Fe/mL) Matrix: Water	GRE SIR (1.5T)	n/a
Yokoo et. al 2015 25996989 [124]	MnCl ₂ (0-25 mM) Matrix: Water	3D GRE (1.5T)	~1-1250 s ⁻¹
Krafft et al. 2016 26308155 [59]	 1) BNF (0.4-220 μg/mL), 2) DSPIO (0.2-125 μg/mL) 3) MnCl₂ (0-3.2 mM) Matrix: Agarose 	Multi-echo GRE, FS GRE (1.5T, 3.0T)	1) 20-2700 s ⁻¹ 2) 20-600 s ⁻¹ 3) 3-250 s ⁻¹
Ibrahim et al. 2016 26861202 [121]	FeSO ₄ (0.5-4.5g) Matrix: Agarose	Multi-echo GRE (3.0T)	~51-435 s ⁻¹
Guimaraes et al. 2016 27190762 [125]	MION 48 (1:50 - 1:10000 of 1 mg Fe/mL) Matrix: Saline	FS TSE, FS CPMG (1.5T)	n/a
Brown et al. 2017 27435747 [126]	USPIO (Resovist®, 0–246.48 µg/mL) Matrix: CAG gel	Multi-echo GRE (3.0T)	22–2829 s ⁻¹
Krafft et al. 2017 28090666 [35]	BNF (0.4-220 μg/mL) Matrix - 2% Agar	Multi-echo UTE (1.5T, 3.0T)	20-2640 s ⁻¹
Hong et al. 2017 27495266 [60]	Ferumoxides (Feridex®, 2-45 mM) Matrix: 0.9% Agarose	Multi-echo 2D- UTE, 2D IR- UTE, 3D UTE, 3D IR-UTE (3.0T)	~0.1-12 ms ⁻¹
Lu et al. 2018 29314215 [127]	 1) Gadopentetate Dimeglumine (Magnevist®,1.5-9 mg/mL) 2) Ferumoxides (Feridex®, 2-22mM) Matrix: 0.9% Agarose 	Multi-echo 3D UTE (3.0T)	1) ~0.2-2.8 ppm 2) ~0.4-4.5 ms ⁻¹ ~0-52 ppm

Table 1. MRI iron phantom studies listing phantom materials used, MR techniques, and R2*/QSM ranges.

Tipirneni-Sajja et al. 2019 30358001 [42]	BNF (0.5-220 µg/mL) Matrix - 2% Agar	Multi-echo UTE (1.5T)	~ 20-2500 s ⁻¹
Headley et al. 2020 33207043 [128]	FeCl ₃ (29.4-96.4 mg) Matrix: Water	Multi-echo GRE (1.5T, 3.0T)	103-420 s ⁻¹ (1.5T) 144-616 s ⁻¹ (3.0T)

Abbreviations: BNF, Bionized NanoFerrite; CAG, carrageenan-strengthened agarose/gadolinium gel; DSPIO, dextran-coated superparamagnetic iron oxide nanoparticles; CPMG: Carr-Purcell-Meiboom-Gill; FeCl₃, Iron(III) chloride; Fe(OH)₃, Iron (III)-hydroxide; FeSO₄, Iron sulphate; FS, fat suppression; GRE, gradient recalled echo; MnCl₂, Manganese chloride; MION 48, dextran-coated magnetic iron oxide nanoparticles; SE, spin echo; SIR, signal intensity ratio; TSE: turbo spin echo; USPIO, ultra-small particles of iron oxide; UTE, ultrashort echo time; IR-UTE: adiabatic inversion recovery prepared UTE.

Another component that impacts the efficacy of iron oxide nanoparticles is the organic coating over the iron core, impacting aqueous stability, conjugation potential, and reducing agglomeration and allowing uniform dispersion of iron particles in the medium [129,130]. For example, pure magnetite powder without a coat forms heterogenous clumps in phantoms [79]. Dextran is one of the most popular coatings used for iron nanoparticles as it is water-soluble and biodegradable [130]. BNF (Bionized NanoFerrite) particles are one example of dextran-coated iron nanoparticles and have been reported to produce a wide span of R2* values (20-2700 s⁻¹), thus covering the entire clinical range of R2* at 1.5T and 3T during iron overload [59]. Figure 5 shows representative MRI iron phantoms made using BNF particles in 2% agar, their corresponding R2* maps obtained using mono-exponential model and the excellent linear correlation obtained between R2* values and iron concentrations in these phantoms.

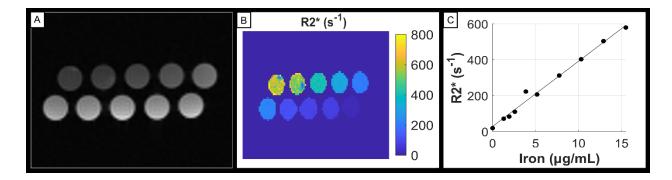


Figure 5. Representative MRI iron phantoms mimicking liver iron overload. (A) MRI magnitude images of iron phantoms consisting of BNF iron nanoparticles of size 80 nm in 2% agar, (B) R2* maps calculated by fitting a mono-exponential model [2], and (C) linear regression plot of R2* (s⁻¹) vs iron concentrations (%) demonstrating an excellent linear correlation with $R^2 = 0.9945$.

Alternatively, SPIO (superparamagnetic iron oxide) nanoparticles, which have emerged as specialized MRI contrast agents, can also be used for mimicking iron overload. The dextrancoated SPIO contrast agent, ferumoxides (Feridex®), is a common choice of iron material used in phantoms, yielding R2* and susceptibility values as high as 100-12000 s⁻¹ and 52 ppm respectively at 3.0T [60,61,127,131]. Some previous studies also used the carboxydextran-coated SPIO contrast agent, ferucarbotran (Resovist®), for mimicking iron overload with R2* values of 22-2829 s⁻¹ at 3.0T [126,132]. Alternatively, ferumoxytol (Feraheme®) can also be used for making iron overload mimicking phantoms by diluting with deionized water, with a reported R2* range of 25 - 150 s⁻¹; however, the R2* range can be increased with smaller dilutions [122].

Certain non-iron contrast agents such as extracellular gadolinium chelates (e.g., gadopentetate dimeglumine (Magnevist®)) and manganese-based contrast agents (e.g., manganese chloride) that are used clinically, also have magnetic properties comparable to those of iron, thereby providing a similar linear relationship between contrast concentration and R2*/susceptibility for mimicking iron overload [127,133,134,124]. Both gadolinium and manganese are water soluble, allowing them to be suspended in solution [133]. However,

compared to manganese-based contrast agents (1-1250 s⁻¹), gadolinium has been reported to provide higher R2* values at 1.5T (up to 3000 s⁻¹) [59,124,135].

One problem with the use of most iron particles is that they are produced on the scale of nanometers. However, in cases of iron overload, accumulated hemosiderin particles are on the scale of micrometers ($0.1 - 2 \mu m$ [93]), a much larger size than iron particles available in commercial products. To mimic in vivo sizes of iron particles, Mobini et al. used insoluble magnetite powder with a mean diameter of $1.17 \mu m$ [$0.5 - 5 \mu m$] in their phantom design to emulate pathological conditions of liver iron overload [79]. However, the inhomogeneous suspension of magnetite resulted in microscopic iron clustering that altered relaxation times, hence leading to overestimated R2* at only mild iron contents [79]. In a recent study, Zhao et al. incorporated soluble iron microspheres with a diameter of $2.9 \pm 0.14 \mu m$, to mimic in vivo hemosiderin particles, and constructed phantoms that produced target R2* values of $100 - 500 \text{ s}^{-1}$ at 1.5T and 3T, which however did not cover the entire clinical spectrum of R2* ranges [68].

MRI phantoms mimicking iron overload are also available ready-made to be purchased from commercial vendors. Calimetrix provides twelve R2* phantoms in the range of 15-1000 s⁻¹ (R2* phantom, Model 300). These phantoms are designed to match clinical hemochromatosis and other iron overload conditions and specifically for testing and quality control of MR-based R2* quantification. CaliberMRI is another commercial vendor that provides a T2 phantom (ISMRM/NIST System Phantom, Model 130) utilizing MnCl₂ with T2 ranges of 5-550 ms (~2-200 s⁻¹ R2 measured at 3T).

Steatosis

Pathophysiology & Morphology

Hepatic steatosis, or fatty liver disease, is the abnormal accumulation of lipids within the cytoplasm of hepatocytes, predominantly in the form of triglycerides. Approximately, 25% of the world population is affected by steatosis [136] and this number is projected to increase further with the increasing epidemics of obesity, diabetes, and metabolic syndrome [137,138]. The liver plays a critical role in lipid metabolism and any disturbances in these metabolic processes can lead to hepatic steatosis, which is a hallmark of many diffuse liver diseases including viral hepatitis and alcoholic and nonalcoholic fatty liver disease [10]. NAFLD can progress to non-alcoholic steatohepatitis (NASH), which is characterized by inflammation, cellular degeneration, and fibrosis and can ultimately lead to cirrhosis [139]. Cirrhosis associated with NASH is estimated to be the leading etiology of liver transplantation in the coming decades [140].

Steatosis can be characterized into two major types: microvesicular and macrovesicular steatosis, depending on the relative size of lipid vacuoles and the displacement of nucleus seen on histology. Macrovesicular steatosis is the predominant form seen in diffuse liver diseases with fat droplet sizes ranging from >1 μ m to the order of the sizes of hepatocytes (20-25 μ m) [141,142]. Steatosis is typically graded semi-quantitatively on histology based on the visually estimated percentage of hepatocytes infiltrated with fat deposits – less than 5% (normal), 5% to 33% (mild), 34% to 66% (moderate) and greater than 66% (severe) as shown in Figure 6 [143]. However, the clinical histological grading estimates only the percentage of affected hepatocytes but does not reflect the absolute fat percentage in the liver.

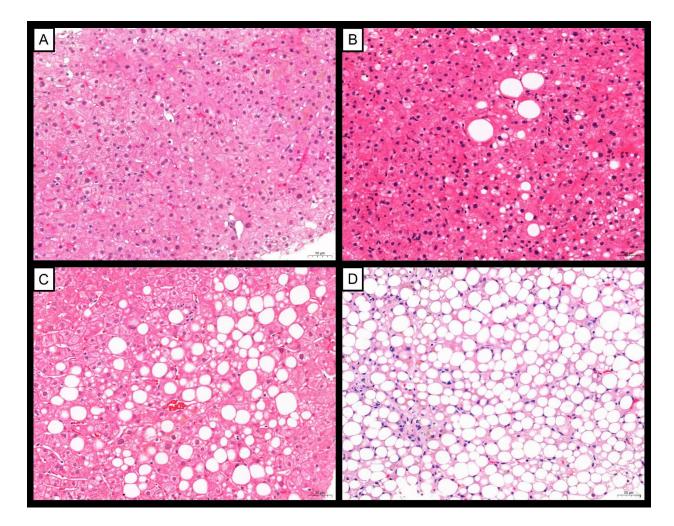


Figure 6. Histology images at 200x of liver biopsy samples stained with hematoxylin and eosin showing the fat droplet morphology and deposition in patients with macrovesicular steatosis. The steatosis grades of these liver biopsy samples based on the NASH CRN scoring system are (A) grade 0 (normal), (B) grade 1 (mild), (C) grade 2 (moderate) and (D) grade 3 (severe).

Quantitative MR Techniques

Magnetic Resonance Spectroscopy

Magnetic resonance spectroscopy (MRS) has been used as a noninvasive alternative to biopsy and is regarded as the most accurate technique to quantify fat [144,145]. MRS is based on the physical principal that MR-active hydrogen protons in specific molecules resonate at a unique Larmor frequency depending on the molecular structure of compounds. For example, fat and water molecules resonate at slightly different frequencies with a frequency difference of 3.4 ppm (224 Hz at 1.5T and 448 Hz at 3T) between the water (-OH) and main lipid peak (methylene, -CH₂) in liver in vivo [146,147]. Based on the priori known chemical shifts, water and fat peaks can be detected using MRS, and their proton densities (proportional to peak amplitudes) can be quantified to estimate fat fraction (FF) [148-150]. However, due to the small sampling volume (~0.2 - 2 cm³ [151]) and the requirement of offline analysis by an experienced user, MRS has limited use for clinical purposes [145]. Alternatively, over the last two decades, chemical shift-based MRI techniques have been developed that provide 2D cross-sectional images or 3D whole liver coverage.

Chemical Shift Based Fat-Water Imaging

Chemical shift-based fat-water imaging is based on the same physical principle as MRS. Due to the difference in the resonance frequencies, fat protons precess at a slower frequency than water protons and develop a phase lag over time compared to that of water protons. For example, the main lipid peak (methylene, -CH₂) and water (hydroxyl, -OH) protons are in-phase at 4.6 ms and out-of-phase at 2.3 ms at 1.5T. The most common chemical shift-based fat-water separation method is 2-point Dixon imaging, which collects images when fat and water signals are in-phase and out-of-phase, and thereby generates fat-only and water-only images from which FF is computed [152,153]. Although 2-point Dixon is a useful technique to assess steatosis [154-157], this method is affected by magnetic field inhomogeneities and T1 bias [158-160], and most importantly, this technique does not consider R2* decay and the spectral complexity of fat which can corrupt FF estimates [158,161]. T1 bias causes overestimation of FF and can be minimized by reducing the flip angle, correcting using variable flip angles, or in post processing [44,162,163]. However, post processing corrections are generally based upon the assumed T1 values of fat and water in the liver, so in vivo pathologies such as iron overload and fibrosis or phantoms that deviate

from these T1 values can produce inaccurate FF results [162,163]. As the signal is collected over multiple echo times during which R2* decay occurs, not accounting for the signal decay can lead to underestimation of FF, especially in tissues with short T2* such as those containing excess iron [152]. Various Dixon techniques have been developed over the years that differ in the number of echoes (or points) that are sampled [164]. While traditional 2-point and 3-point Dixon methods are desirable for a shorter scan time, they typically provide only qualitative information about fat content [164]. On the other hand, using six or more echoes increases the scan time, but they allow for accurate modeling and provide quantitative FF results when fitted with multispectral fat-water R2* models described below. [164-166]. These chemical shift-based MRI techniques are well validated against MRS, showing an excellent agreement between MRS and MRI FF values [167-169]. Further, previous studies demonstrated that clinical MRI FF values can range up to ~50% in patients and showed a good correlation to severity of steatosis graded on histology [170-172].

Multi-spectral Fat-Water Models with R2* Correction

Over the last decade, several multi-spectral signal modeling techniques have been proposed for accounting for the spectral complexity of fat and R2* decay. These techniques use multi-echo GRE acquisitions and perform multi-spectral fitting for simultaneous quantification of R2* and FF and thus, can enable the diagnosis of both hepatic iron overload and steatosis. One popular technique is based on nonlinear least squares (NLSQ) fitting of the fat–water signal model either using magnitude or complex data [67]. These models require priori information about relative frequencies and amplitudes of the multiple lipid peaks. Further, most NLSQ models assume a common R2* rate for fat and water for reducing model complexity [67], known as the single R2* model. However, this assumption will be incorrect if the R2* of fat and water are different, especially in cases of iron overload [173]. A dual R2* model has been developed and was shown to improve the FF quantification accuracy as compared to the single R2* model in phantoms [173]. However, the dual R2* model can be unstable and complex due to the estimation of an additional parameter, and the single R2* model has been described to be more accurate in low SNR conditions for FF estimation [174]. Additionally, recent studies have reported that multipeak fat modeling with complex fitting and single R2* correction model performs better over a range of clinically relevant FF values [174,175].

Another multi-spectral signal modeling technique is based on autoregressive moving averaging (ARMA) that does not use any prior information of fat spectrum and determines FF and separate R2* values for water and lipid species [44,176,177]. A recent study demonstrated that both NLSQ and ARMA models accurately quantified R2* and FF values, but ARMA yielded inaccurate values at both high iron and fat conditions, whereas NLSQ overestimated R2* and FF values at high iron and no fat conditions; both models potentially failing due to rapid T2* decay at high iron overload conditions [44].

Steatosis Phantoms

The design considerations for steatosis phantoms are to emulate the MRI characteristic multi-peak fat spectra seen in adipose tissue, match clinical ranges of hepatic steatosis (~0-50% FF), and maintain the size and distribution of fat deposits seen in liver tissues homogeneously throughout the phantom. Table 2 details the lipid substitutes, FF ranges, and MR techniques used in steatosis phantom studies.

Peanut oil is the most common material used in fat phantom studies due to its availability, ease of preparation, and similar NMR spectrum as triglycerides in adipose tissue [51-53,178-180]. Other types of vegetable oils, such as canola or soybean, have also produced similar FF results to peanut oil despite slight differences in saturated and unsaturated fat content [81,150,181-185]. Animal products such as lard and pork fat are also used as lipid substitutes and are mixed with appropriate proportions of liver tissue to create a homogenate or heated until in a liquid form for making steatosis phantoms [18,186]. As seen on Table 2, mayonnaise is another animal product used in steatosis phantoms and has been shown to display similar MRI signal behavior as other lipid substitutes when used in combination phantoms [117]. Hybrid plant and animal products can also be used to create steatosis phantoms through mixing vegetable oils and liver tissues [81]. Further, clinical fat emulsion supplements such as Intralipid® and MicrolipidTM can also be used for constructing steatosis phantoms [150,187]. Alternatively, commercial steatosis phantoms are available from Calimetrix (Fat Fraction Phantom, Model 300) covering FF ranges from 0-50% and 100% [54].

Citation PMID	Phantom Materials	MR Technique (Field Strength)	FF (%) Range
Chang et al. 2006 16794122 [81]	Canola Oil Matrix: Calf liver	2-echo GRE, MRS (1.5T)	0-80%
Bernard et al. 2008 18064714 [181]	Soya oil Matrix: Carageenan	2-echo Dixon, chemical shift imaging, MRS (3.0T)	0-100%
Hamilton et al. 2011 21834002 [150]	Safflower oil (Microlipid®)	MRS (3.0T)	50%
Peng et al. 2011 21737754 [182]	Vegetable oil Matrix: Water	2-echo Dixon, chemical shift imaging, MRS (7.0T)	0-100%
Mashhood et al. 2013 23172799 [187]	Intralipid® fat emulsion Matrix: Water	2-, 4-, 6-echo Dixon (1.5T, 3.0T)	0-30%
Leporq et al. 2013 23588583 [183]	Olive oil Matrix: Agar	MRS, Multi-echo GRE (1.5T, 3T)	0-50%
Deng et al. 2014 24840769 [18]	Pork visceral fat Matrix: liver tissue	Multi-echo GRE (1.5T)	3.5- 46.6%

Table 2. MRI fat phantom studies listing the phantom materials used, MR techniques, and fat fraction ranges.

Hernando et al. 2017 27080068 [52]	Peanut oil Matrix: 2% Agar	Multi-echo GRE (1.5T, 3.0T)	0-100%
Fukuzawa et al. 2017 28770394 [184] Hayashi et al. 2018 29425901 [185]	Vegetable oil (soybean & rapeseed) Matrix: 1% Agar	MRS, Multi-echo Dixon (3.0T)	0-50%
Pooler et al. 2018 29665193 [178]	Peanut oil Matrix: 2% Agar	MRS, Multi-echo GRE (1.5T)	0-100%
Pickhardt et al. 2018 30016142 [53]	Peanut oil Matrix: 2% Agar	Multi-echo GRE (1.5T, 3.0T)	0-100%
Kim et al. 2019 30430684 [186]	Lard Matrix: Agar	MRS, Multi-echo GRE (1.5T, 3.0T)	0-60%
Jaubert et al. 2020 31736146 [179]	Peanut oil Matrix: 2% Agar	6-echo Dixon (1.5T)	0-100%
Hu et al. 2021 33464181 [54]	Fat Fraction Phantom, Model 300 (Calimetrix)	Multi-echo GRE (1.5T, 3.0T)	0-100%
Schneider et al. 2021 33768291 [180]	Peanut oil Matrix: 2% Agar	Multi-echo GRE (3.0T)	0-100%
Zhao et al. 2021 34105167 [51]	Peanut oil Matrix: 2% Agar	Multi-echo GRE (1.5T, 3.0T)	0-100%

Abbreviations: GRE, gradient recalled echo; MRS, magnetic resonance spectroscopy, SS-EPI, single-shot echo planar imaging.

When emulating steatosis, emulsifying agents are typically needed to disperse oil droplets homogeneously throughout the sample. All emulsifiers have amphiphilic properties that interface between the oil and aqueous phases, but some emulsifiers are more soluble in oil than water and vice versa. In the case of phantoms, the most commonly used emulsifier is sodium dodecyl sulfate (SDS), an anionic surfactant [52,61,79,131]. SDS is an extremely hydrophilic emulsifier with a much higher affinity for aqueous media compared to oil-based media [188]. However, some researchers have opted for using oil-soluble emulsifiers such as Span 80 or lecithin [184,189-191]. This allows a lipophilic emulsifier to be added directly to oil during the phantom preparation to

stabilize water-in-oil emulsions rather than dissolving a hydrophilic emulsifying agent in water to stabilize oil-in-water emulsions [190]. However, caution must be taken when using emulsifiers in phantom studies as some emulsifying agents may alter the T1 and T2 relaxation times of the phantom [192].

During the production of steatosis phantoms, it is important to note that the mixing temperature of fat and matrix emulsions and other environmental conditions can affect the quality of the final phantom [189]. Notably, caution must be taken to mix oil and polysaccharide emulsions such as agar/agarose in a limited temperature range of $\sim 40-50^{\circ}$ C as mixing them at higher temperatures can cause creaming (i.e., separation of oil and aqueous media) while lower temperatures can solidify polysaccharide matrix solutions, both cases producing inhomogeneous suspensions [189]. When scanning, changes in room temperature will greatly change the resonance frequency of water, but the fat resonance frequency will remain relatively unaffected [146]. In fact, the frequency difference between main fat peak and water at room temperature is 3.5 ppm whereas it is 3.4 ppm in in vivo liver imaging [146,147]. However, changing the temperature of the scanning room to mimic body temperature is not practical, so creation of steatosis phantoms that retain the frequency difference of fat and water as in vivo liver is therefore an unmet need. Nevertheless, maintaining the temperature of the phantom throughout the duration of the scan and adjusting the chemical shift accordingly can produce higher reproducibility and accuracy in FF quantification [146].

Another issue in constructing fat phantoms emulating hepatic steatosis is the discrepancies in sizes between fat globules in in vivo hepatocytes and steatosis phantoms. Hines et al. acquired photomicrographs of a fat-water-SPIO phantom at fat fractions of 10% and 50%, and found that average diameters of the fat droplets were $47 \pm 19 \,\mu\text{m}$ and $234 \pm 66 \,\mu\text{m}$, respectively, which were much larger than the average diameter of the hepatocytes, $20-25 \ \mu m$ [61,193]. This dissimilarity is not only different from histologic conditions, but it may also contribute to faster R2* decay than in vivo samples [61,79]. Some recent studies have incorporated alternative mixing methods such as hand-held or high-pressure homogenization to reduce the sizes of fat droplets [68,191]. However, microscopy validation has not been performed in these studies to ensure that the diameter of the fat droplets match those of in vivo hepatocytes [68,191].

Fibrosis

Pathophysiology & Morphology

In cases of persistent liver injury such as in alcoholic liver disease, viral hepatitis, iron overload, and steatosis, functional liver tissue is progressively replaced primarily by extracellular matrix (ECM) components in a process called fibrosis [194]. In fibrosis, abnormal amounts of scar tissue replace functional hepatic cells and disrupt internal organ structure that over time limits the blood flow as well as the synthetic capabilities of the liver. Collagen is the most abundant ECM protein, and as fibrosis progresses, the type and composition of collagen changes dramatically with loss and over expression [195,196]. Predominantly, nanofibers of type I collagen with diameters of 300–500 nm replace functional hepatic tissues in fibrosis [197,198]. Initially, fibrosis does not cause noticeable symptoms, but as it worsens, it can lead to portal hypertension and cirrhosis [199]. Cirrhosis is the most advanced stage of liver fibrosis and is evidenced by fibrotic bands, parenchymal nodules, and vascular distortion – all of which derail hepatic function leading to liver failure [200]. The severity of fibrosis is graded semi-quantitatively on histology using the NASH CRN scoring system as follows: 0 (no fibrosis), 1A (mild zone 3 perisinusoidal fibrosis), 1B (moderate zone 3 perisinusoidal fibrosis), 1C (portal/periportal fibrosis only), 2 (perisinusoidal and portal/periportal fibrosis), 3 (bridging fibrosis), and 4 (cirrhosis) [201]. Accordingly, Figure 7

shows representative histology images and fibrosis morphology in patients with different fibrosis grades.

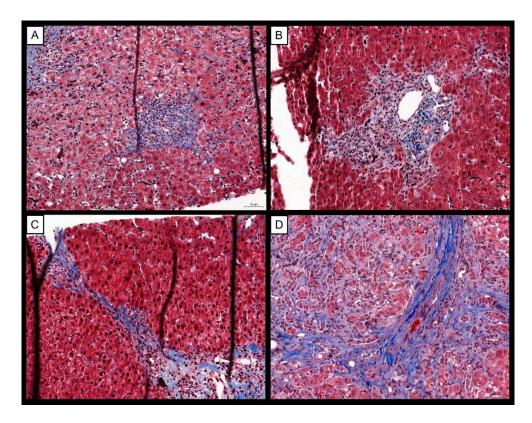


Figure 7. Histology images at 20x of liver biopsy samples showing the deposition of extracellular matrix (primarily collagen) with Mason's trichrome stain in patients with different fibrosis grades. The fibrosis grades of these histology samples based on the NASH CRN scoring system are stage 1, perisinusoidal (A), stage 2, perisinusoidal and periportal (B), stage 3, bridging fibrosis (C), and stage 4, cirrhosis (D).

Quantitative MR Techniques

Elastography

One of the most widely validated MRI techniques used in staging liver fibrosis is magnetic resonance elastography (MRE) [202]. Elastography takes advantage of the relationship between tissue stiffness and the speed of mechanical wave propagation, with waves propagating faster in harder tissues compared to soft tissues, to evaluate collagenous stiffening due to hepatic fibrosis [203]. Specifically, MRE methods use a low frequency vibration source (40-80 Hz) to generate mechanical waves, followed by the utilization of phase contrast techniques with motion encoding

gradients to capture microscopic displacements caused by wave propagation [204,205]. The resulting wave information is then processed to produce tissue stiffness maps using an inversion algorithm to recover mechanical parameters from the displacement data. [204,205]. MRE has been shown to have a high sensitivity and specificity for distinguishing among NAFLD, NASH, fibrosis, and cirrhosis [202,206] and can image longitudinal changes in stiffness in patients over time [207]. Previous studies demonstrated good correlation between MRE stiffness values with both ultrasound elastography (Fibroscan) [62,208,209] and biopsy fibrosis grades [206,210,211], which are often used for clinical diagnosis, thus, making MRE a promising non-invasive alternative for assessing hepatic fibrosis.

Although hepatic MRE is accurate, it can fail in some pathologies such as iron overload in which significant signal decay occurs when using gradient echo based MRE sequences [204,212]. The spatial resolution of MRE can be improved by increasing the shear wave frequency; however, high-frequency waves attenuate quickly, so there is a tradeoff between spatial resolution and propagation distance, thus restricting the depth that can be imaged [213]. Due to this, MRE techniques are limited in use in obese subjects and patients with ascites [214].

T1 Imaging

T1 imaging is arising as an alternative in staging fibrosis to overcome the limitations with MRE. In fibrotic conditions, widespread hepatocyte injury occurs, consequently inducing the recruitment of inflammatory cells. The progression of fibrosis thus leads to chronic inflammation and increases extracellular volume throughout the liver [215]. Because T1 relaxation is altered by the molecular environment of water molecules in tissue, the increased extracellular volume in fibrosis and cirrhosis also prolongs liver T1 values [216]. Therefore, T1 imaging could be used as a noninvasive MRI biomarker for evaluating fibrosis. In the literature, T1 mapping has been used

in staging myocardial fibrosis [217] as well as in many independent studies for assessing hepatic fibrosis [216,218-221]. While T1 has had high diagnostic accuracy, there are some significant drawbacks to this approach. Notably, T1 values can be confounded by other liver pathologies such as inflammation, steatosis, iron overload and portal hypertension [222-225].

Diffusion

Diffusion-weighted imaging (DWI) is one of the first MR techniques used for assessing liver fibrosis. DWI assesses the degree of Brownian motion of water protons by using motion sensitizing gradients and produces image contrast based on the fact that freely moving protons lose signal faster than restricted molecules [226]. Because ECM remodeling leads to a much higher expression of collagen, the flow of water is restricted in fibrotic tissues thus reducing the apparent diffusion coefficient (ADC) values [227,228]. The normalized ADC values for healthy livers are typically ~1.50-1.85 (x10⁻³ mm²/s) while for fibrotic livers are ~1.30-1.45 (x10⁻³ mm²/s) [227,229-232]. However, these ADC measurements can vary between vendors and scanners as they are sensitive to b-values and image noise, thus the cutoff values for staging liver fibrosis are difficult to establish [233]. More importantly, ADC values are also affected by iron overload and steatosis, hence limiting their usage for assessing co-occurring diffuse liver diseases [122,234].

Fibrosis Phantoms

The design requirements of fibrosis phantoms are to maintain characteristic relationship between MRI biomarkers (stiffness, T1, and ADC) and fibrosis biopsy grades and cover a wide range of clinical values of MRI biomarkers seen in conditions of liver fibrosis, stiffness: ~2.5-8.0 kPa [206,210,211,235], T1: ~500-1000 ms [45,74], ADC: $1.3 - 1.85 (x10^{-3} mm^2/s)$ [227,229-232]. A variety of phantom materials are used to emulate the correlation with the three MRI biomarkers used for the evaluation of liver fibrosis as detailed in Table 3. Most frequently, fibrosis phantoms are created with varying stiffnesses by changing the concentration of the medium. The differing Young's moduli between each sample allows MRE techniques to quantify the degree of stiffness in the phantom, thus allowing the emulation of different fibrosis grades [62]. In vivo, collagen is the primary component that makes up fibrosis pathologies, so some phantom studies have incorporated differing types and concentrations of collagen into their design [63,117]. Apart from collagen, gelatins, hydrogels, and polymers such as B-gel (bovine gelatin), agar, tofu, grenetine gel, Phytagel, Zerdine, silicone, polyvinyl alcohol, and polyacrylamide can also be used in MRE studies for emulating fibrosis [62,65,236-243]. Elastography phantoms are also available from commercial vendors such as Computerized Imaging Reference Systems, Inc. (Zerdine shear wave liver fibrosis phantom), which are designed specifically for ultrasound elastography but have been used in liver MRE studies [239]. Alternatively, commercial MRE phantoms with stiffness values between 2-4 kPa are also available from Resoundant, an MRE manufacturer, for verification and validation of the MRE system. Figure 8 shows a representative set of MRE phantoms with increasing stiffness values made using different concentrations of Phytagel covering the range of stiffness values similar to those seen in liver fibrosis.

Citation PMID	Phantom Materials	MR Technique (Field Strength)	Stiffness(kPa)/T1(ms)/ ADC(mm ² /s) Range
Girometti et al. 2007 17440695 [244]	1) Water 2) Acetone	SE-EPI DWI (1.5T)	1) $2.45 \pm 0.05 (x10^{-3} mm^2/s)$, 2) $4.21 \pm 0.05 (x10^{-3} mm^2/s)$ (ADC)
Yin et al. 2007 17654577 [236]	15% B-gel	GRE MRE (1.5T)	11.2 kPa (G)
Oudry et al. 2009 19856447 [62]	B-gel (5-17.5%) Copolymer-in-oil (3-8.5%)	GRE MRE (1.5T)	~1.0 – 8.0 kPa (G)
Cui et al. 2015 26393236 [245]	Ice water	IVIM DWI (1.5T, 3.0T)	$\begin{array}{l} 1.13 \pm 0.03 \ (x10^{-3} \\ mm^2/s), \\ 1.21 \pm 0.02 \ (x10^{-3} \\ mm^2/s) \end{array}$
Salameh et al. 2016 26915977 [240]	7% Polyvinyl alcohol gel	Ultra-low field MRE (6.5 mT)	2.3 ± 1.0 kPa (G')
Morisaka et al. 2016 27662640 [65]	Polyacrylamide gel (6.2-7.7%)	SE-EPI, GRE MRE (3.0T)	2.4 - 4.2 kPa (S)
Kishimoto et al. 2017 28057657 [241]	Acrylamide gel	SE-EPI MRE (3.0T)	2.1 - 25 kPa (G')
Solamen et al. 2018 29877194 [237]	 1) 55-70% A341 Silicone soft gel 2) 70%,100% A341 Silicone soft gel 3) Soft Tofu and 0.75% Agar 4) Soft Tofu and 0.65% Agar 	NLI-MRE (3.0T)	1) 2.5 - 4.6 kPa (G') 2) 4 -15 kPa (G') 3) ~4 kPa (G') 4) 2.7 - 4.2 kPa (G')
Tirkes et al. 2019 31165353 [246]	System Phantom Model 130 (High Precision Devices, Inc.)	VFA, MOLLI, IR-SNAPSHOT, SASHA (1.5T)	20 - 1500 ms (T1)
Cho et al. 2020 33110134 [45]	Gadoterate meglumine (0.1-3 mmol/L) Matrix: Water	SE-IR, VFA, MOLLI (1.5T)	83 - 1310 ms (T1)
Usumura et. al 2021 34019551 [242]	Polyacrylamide gel (12 wt % Acrylamide, 45% Glycerin)	SE-EPI MRE (3.0T)	4.8 - 5.01 kPa (G')
Andoh et al. 2021 34037285 [239]	Zerdine® solid hydrogel (Computerized Imaging Reference Systems, Inc.)	SE MRE (1.5T, 3.0T)	1.7-14.9 kPa (G)

Table 3. MRI fibrosis phantom studies listing the phantom materials used, MR techniques, and stiffness/T1/ADC ranges.

Meneses et al. 2021 34209547 [247]	Human serum albumin (0-20%)	T1 mapping (3.0T)	1800-4000 ms (T1)
Kishimoto et al. 2022 35061118 [243]	Polyacrylamide gel (13%)	SE-EPI MRE (0.3T)	3.1 kPa (G')

Abbreviations: B-gel, bovine skin powder; DWI, diffusion-weighted imaging; GRE, gradient recalled echo; IR-SNAPSHOT, inversion recovery snapshot FLASH; IVIM, intravoxel incoherent motion; MRE, magnetic resonance elastography; MOLLI, modified look-locker inversion recovery; NLI, non-linear inversion; SASHA, saturation recovery single-shot acquisition; SE-EPI, spin-echo echo planar imaging; SE-IR, spin echo inversion recovery; VFA, variable flip angle. **Note:** Elastography results are published using different variables: S, stiffness; G, shear modulus; G', storage modulus

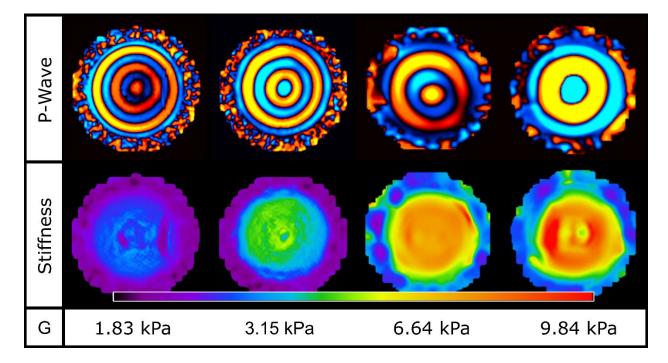


Figure 8: P-wave, stiffness maps, and shear moduli (G) obtained from four magnetic resonance elastography (MRE) phantoms made of Phytagel (1.25 - 1.875%) for emulating liver fibrosis [3]. The stiffness values obtained with MRE ranged from 1.83 - 9.84 kPa.

MRI fibrosis phantoms can also be created using T1 modulators to manipulate T1 times and emulate fibrosis conditions at different grades. Gadoterate meglumine (Dotarem®) is an FDA approved paramagnetic contrast agent capable of shortening T1 relaxation times, hence making it suitable for constructing fibrosis phantoms that are assessed using T1 mapping [45,248]. In contrast, a recent study incorporated human serum albumin as a biomarker for fibrosis as it has similar relaxation times to human tissues and used it to modulate T1 times by varying its concentration for emulating fibrosis [247]. T1 phantoms are also commercially available from CaliberMRI (ISMRM/NIST System Phantom, Model 130) and Calimetrix (T1 Phantom, Model 300) [246,249]. Both phantoms use NiCl₂ as a T1 modulator, but the CaliberMRI System Phantom provides a wider coverage of T1 values of 20-1900 ms, whereas the Calimetrix T1 phantom provides a more limited range of 100-1200 ms; both phantoms' T1 values reported are at 3.0 T.

DWI phantoms are produced from a variety of materials including aqueous solutions (polyvinylpyrrolidone, sucrose, agarose, etc.), polymers (liquid paraffin and alkanes), and pure water. DWI phantoms are also available from commercial vendors such as CaliberMRI (Diffusion Phantom Model 128) and by organizations such as the National Institute of Standards and Technology (NIST) [56,250]. For DWI phantoms mimicking hepatic fibrosis, aqueous solutions of water, acetone or ice water are the most common materials used [228,244,245]. However, phantoms using water and acetone produced higher ADC values than observed in healthy or fibrotic livers but in contrast, using ice water produced ADC values closer to liver tissue [228,244]. Diffusion coefficients have a dependency on field strength and temperature, hence the phantom temperature should be maintained for consistent results [56]. Consequently, DWI phantoms that encompass clinical ranges of fibrosis are still an unmet need in the literature.

Combination Phantoms

Because of the common co-occurrence of iron overload, steatosis, and fibrosis, there have been efforts to combine these pathologies into a standardized set of phantoms, as documented in Table 4. Combination phantoms incorporate the same materials as a single pathology phantom. However, due to the confounding effects between iron, fat, and fibrosis on MRI signals, there have been attempts to find unique biomarkers for each pathology, namely R2* or susceptibility for iron, FF for fat, and stiffness or T1 for fibrosis. Despite this effort, there are still challenges concerning these confounding effects that must be overcome to create an ideal set of combination liver phantoms. **Table 4:** MRI iron-fat and iron-fat-fibrosis phantom studies listing the phantom materials, MR techniques, R2*, fat fraction and T1 ranges.

Condition Mimicked	Citation PMID	Phantom Materials	MR Technique (Field Strength)	R2*(s ⁻¹)/QSM (ppm) Range	FF (%) Range	T1 (ms) Range
Iron & Fat	Sharma et al. 2009 19243059 [251]	Iron: Ferumoxides (Feridex®, 0- 0.5 mM) Fat: Vegetable oil Matrix: 2% Agar	MRS (1.5T)	11-133 s ⁻¹ (R2)	10, 30%	n/a
Iron & Fat	Hines et al. 2009, 19856457 [61]	Iron: Ferumoxides (Feridex®, 0- 50 µg/mL) Fat: Peanut oil Matrix: 2% Agar	2-echo Dixon, Multi-echo GRE, MRS (1.5T)	~25-375 s ⁻¹	0-100%	n/a
Iron & Fat	Boll et al. 2010 20308498 [252]	Iron: Ferumoxides (Feridex®, 0.002-0.027 mL) Fat: Caprylic/capric acid (3-12 mL) Matrix: Polyacrylamide	2-echo Dixon (3.0T)	n/a	10-40%	n/a
Iron & Fat	Lee et al. 2011 21591008 [131]	 Iron: Ferumoxides (Feridex®, 0, 11.2 μg/mL) Fat: Canola oil (0-50%) Matrix: 2% Agar Iron: Feriumoxides (Feridex®, 0-14 μg/mL) Fat: Canola oil (10%) Matrix - 2% Agar 	Multi-echo GRE, MRS (1.5T)	1) n/a 2) ~45–100 s ⁻¹	0–50%	n/a
Iron & Fat	Fukuzawa et al. 2017 28770394 [184]	Iron: SPIO (0-0.4 mM) Fat: Vegetable oil Matrix: 1% Agar	MRS, Multi- echo Dixon (3.0T)	33–217 s ⁻¹	0, 20%	n/a

Iron & Fat	Tipirneni-Sajja et al. 2019 30761652 [44]	Iron: BNF (0-60 µg/mL) Fat: Peanut oil Matrix: 2% Agar	Multi-echo GRE (1.5T, 3.0T)	0-800 s ⁻¹	0–40%	n/a
Iron & Fat	Wang et al. 2019 30859893 [191]	Iron: Iron dextran (0-30 µg/mL) Fat: Vegetable oil Matrix: Water	2-echo GRE (3.0T)	~0-900 s ⁻¹	0–80%	n/a
Iron & Fat	Mobini et al. 2019 31732894 [79]	Iron: Magnetite (0-723.6 µg Fe/mL) Fat: Peanut oil Matrix: 0.9% Agar	Multi-echo GRE (3.0T)	~45-1000 s ⁻¹	0–100%	n/a
Iron & Fat, Iron & Fibrosis	Li et al 2018 29566449 [117]	 Iron: Gadolinium (0-10.0 mM) Fibrosis - Collagen (0-30%) Matrix – Water Iron: Gadolinium (0-10.0 mM) Fat: Mayonnaise Matrix: Water 	Multi-echo GRE (1.5T, 3.0T)	1) 0–150 s ⁻¹ 0–3 ppm 2) 0–350 s ⁻¹ 0–3 ppm	0– 43.7%	n/a
Iron, Fat & Fibrosis	Zhao et al. 2021 32783200 [68]	 Iron: MnCl₂ (0.17- 1.17 mM) & Magnetite (1.03-34.53 μg/mL) Fat - Peanut oil Matrix - 2% Agar Iron: Magnetite (1.03-53.65 μg/mL) Fat - Peanut oil Fibrosis: NiCl₂ (0.28-2.10 mM) Matrix: 2% Agar 	Multi-echo GRE, MRS (1.5T, 3.0T)	1) 100–500 s ⁻¹ 2) 50–600 s ⁻¹	1) 5– 30% 2) 0– 40%	1) n/a 2) 500–1500 ms
Iron & Fat	Colgan et al. (2021) 33783066 [253]	Iron: Magnetite Fat: Peanut oil Matrix: 2% Agar	Multi-echo GRE (1.5T, 3.0T)	200-1000 s ⁻¹	0-30%	n/a

Abbreviations: BNF, Bionized NanoFerrite; GRE, gradient recalled echo; MnCl₂, manganese chloride; NiCl₂, nickel chloride; SPIO, superparamagnetic iron oxide nanoparticles.

Iron-Fat Phantoms

Most literature on combination phantoms features two coexisting pathologies: iron overload and steatosis. For iron overload, iron nanoparticles are most frequently used to emulate the pathology [44,184,191], but contrast agents such as ferumoxides (Feridex®) [61,131,251] and magnetite microspheres [79,253] have been reported as well. Likewise, the most popular substitution for steatosis is peanut oil [44,61,68,79,253], but other vegetable oils such as canola and soybean oil [131,184,191,251] have also been used in iron-fat phantoms. Iron overload and steatosis combination phantoms, however, present additional complications compared to singlepathology phantoms. When placed in an aqueous environment, lipids will form large globules with signal-generating hydrophobic tails in the interior of the bilayer [61]. Conversely, some iron particles are only water-soluble and remain in aqueous solution [61]. Because of this compartmentalization, iron particles, especially of nanometer sizes, may preferentially affect the R2* of water molecules but not that of the R2* of large fat droplets, thereby causing inaccurate R2* and FF measurements, especially at high iron concentrations when a single R2* fitting model is used [61,173]. However, some recent studies used iron microparticles to produce single-R2* behavior in the presence of fat and water to emulate the in vivo conditions and yield accurate R2* and FF measurements [68,79]. Figure 9 shows a representative set of MRI iron-fat phantoms and their R2* and FF maps obtained using the NLSQ fat-water model with single R2* correction. As shown, both measured R2* and FF values showed an excellent linear correlation with iron concentrations and true FFs respectively, except for the highest iron concentration.

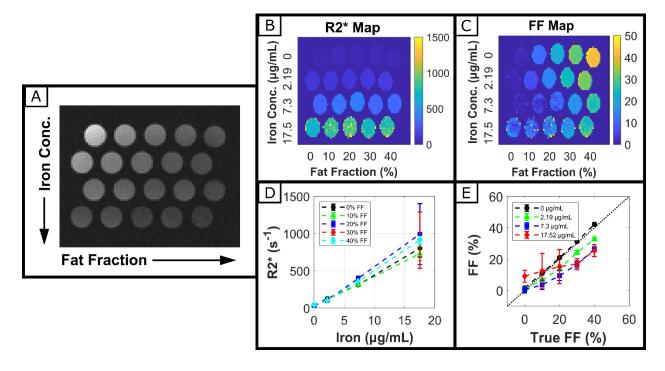


Figure 9. Representative MRI iron-fat phantoms mimicking both liver iron overload and steatosis. BNF iron particles (size: 80 nm) and peanut oil are used to emulate iron overload and steatosis, respectively. MRI magnitude images (A), R2* and fat fraction (FF) maps calculated by fitting a non-linear least squares (NLSQ) multi-spectral fat-water model with R2* correction (B, C) [4], and scatter plots (with error bars) of R2* (s⁻¹) vs iron concentrations (D) and measured fat fraction (FF) vs true FF values (E) are shown for these iron-fat phantoms. Both measured R2* and FF values showed an excellent linear correlation with iron concentrations and true FFs respectively, except for the highest iron concentration.

Iron-Fat-Fibrosis Phantoms

To date, very few studies have integrated fibrosis into iron-fat combination phantom designs to emulate all three diffuse liver pathologies. Zhao et al. incorporated NiCl₂ as a T1 modulator for emulating fibrosis and used peanut oil and iron microspheres for mimicking steatosis and iron overload, respectively [68]. NiCl₂ is suitable for combination phantoms as it is less sensitive to causing R2/R2* changes in combination iron phantoms [254]. However, iron shortens T1 times, so iron-fibrosis combination phantoms using iron particles in tandem with NiCl₂ should correct or account for the decrease in T1 values due to iron, which in fact mimics the shortening of liver T1 in the presence of iron overload [68,255-257]. Further, in this study, the R2* and T1

values in phantoms did not show the similar field strength dependency of these parameters between 1.5T and 3T that is observed in liver in vivo [68]. To investigate the effectiveness of QSM compared to R2* relaxometry for quantifying iron in coexisting pathologies, Li et al. made a set of iron-fibrosis phantoms using gadolinium and collagen, and iron-fat phantoms using gadolinium and mayonnaise [117]. Although it is well documented that iron and fat both influence R2* decay, this study reported that collagen used in fibrosis phantoms also has a positive association with R2* decay [117] and thus, can confound R2*-based HIC estimations. In contrast, susceptibility measurements were not confounded by the presence of fat or fibrosis in these combination phantoms, demonstrating that QSM is a better biomarker for iron estimation in coexisting pathologies [117].

Outlook

The morphology of in vivo tissues is often overlooked when creating liver phantoms emulating iron overload, steatosis, and fibrosis pathologies. Commercial iron particles are only available on the nano scale, but micron-sized particles are needed to mimic the sizes of hemosiderin deposits. The sizes of fat droplets are currently unvalidated in phantoms studies and may be much larger than the size of fat globules in hepatocytes. Therefore, homogenization of fat droplets is needed to reduce their size and mimic in vivo hepatocyte fat storage. The inclusion of materials that simulate fibrosis is important due to its frequent incidence in iron overload and/or steatosis pathologies. Currently, most phantom studies only focus on mimicking iron overload, steatosis or both. However, there is little research on combining these pathologies with the presence of fibrosis. Likewise, most fibrosis phantoms concentrate primarily on stiffness without considering the coexisting pathologies such as iron overload or steatosis. Hence, there is a gap in our understanding of imaging the complex interplay between these diffuse liver diseases in a standardized phantom model using MRI. Further studies are needed in combining all three pathologies to produce a more precise set of phantoms that correspond to realistic disease progression in in vivo conditions and to investigate and validate MRI techniques for clinical use. Lastly, this study only focused on MRI phantoms mimicking diffuse liver pathologies of iron overload, steatosis, fibrosis and their combination, however there are other features of diffuse liver disease such as inflammation and ballooning as well as other hepatic pathologies (portal hypertension, focal lesions, etc.) and MR techniques such as dynamic contrast enhanced methods that are beyond the scope of this work.

Conclusion

Iron overload, steatosis, and combined overload are common manifestations of diffuse liver disease that may lead to further injury in the form of fibrosis. MRI has opened new avenues for noninvasive quantification of iron overload, steatosis and fibrosis as an alternative to liver biopsy. To standardize MRI techniques, liver phantoms that mimic these pathologies are developed using matrix materials such as water and agar. To simulate iron overload, iron particles are used as R2* modulators or as sources of magnetic susceptibility. Likewise, fatty oils or animal tissues are used to create steatosis phantoms using emulsifiers to ensure homogeneous dispersion. For emulating fibrosis, phantom materials that change stiffness values based on concentration or T1 modulators are used. However, the current phantom studies suffer from morphological discrepancies in phantom design when compared to in vivo sizes of these particles and also from confounding effects on MRI biomarkers in cases of co-occurring iron, fat, and fibrosis pathologies. Future studies should be aimed at selecting phantom materials that emulate the size and distribution of these diffuse liver pathologies in vivo and also consider the complex microenvironment of these materials and how they impact the MRI signal for creating realistic morphological quantitative liver MRI phantoms.

Chapter IV

Impact of Particle Sizes on MRI Signal Relaxation in Phantoms for Assessment of Hepatic Steatosis and Iron Overload

Introduction

Current hepatic MRI phantoms have little to no standardization and do not consider the morphological sizes of iron particles and fat deposits seen in human tissues which may affect the MRI signal behavior differently [61,79]. Hepatic iron overload is characterized by differing ratios of ferritin (50-70nm) [91] and hemosiderin $(0.1-2\mu m)$ [93], yet phantom studies typically employ iron oxide particles far larger or smaller than these iron storage proteins [68,122]. Notably, hemosiderin is the primary iron storage protein under conditions of severe iron overload and tends to aggregate in sizes $\geq 1 \mu m$ [91], but most commercial iron particles are produced on the nano- scale rather than micro-. Further, the size of fat droplets in hepatic steatosis phantoms are often not controlled and are larger than the average diameter of hepatocytes (20-25 μ m) [193]. Iron is speculated to preferentially effect water molecules over lipid molecules due to geographic limitations as the signal-generating lipid tails are located on the interior of globule bilayers and are less accessible to iron particles [61]. As seen on Figure 10 below, increasing the size of lipid globules will thereby allow less interactions between fat and iron as iron is soluble in water, but insoluble in fat [258]. Therefore, iron will be more isolated from lipid molecules than water, and this distinction becomes more apparent with a lesser emulsification. Likewise, increasing the size of iron particles causes a greater proportion of the particle volume to be consisted of the magnetic core rather than the magnetically dead polymer coating, thereby causing greater magnetic field inhomogeneities to be imposed in larger particles [259]. Therefore, the R2* decay rates of fat and water caused by the superparamagnetic effects of iron

particles on surrounding lipid and water molecules respectively in phantoms may be different than what is seen in vivo.

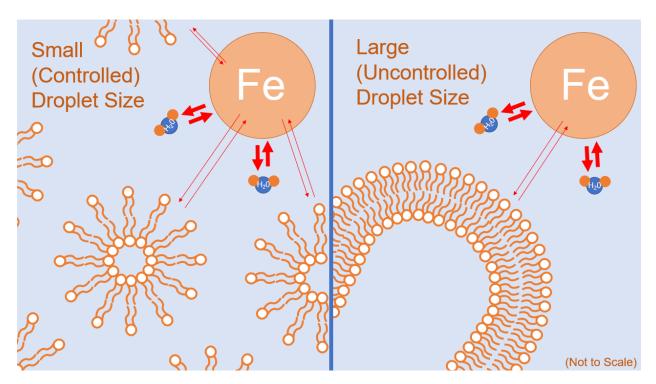


Figure 10. Overview of molecular interactions between iron, water, and lipids. (Left) lipid globules are controlled to be a smaller size via emulsification and/or homogenization, therefore allowing more interactions between iron particles. (Right) uncontrolled lipid globules form via self-assembly and are more geographically isolated from iron due to the large globule size.

Hence, the purpose of this study is to develop and evaluate quantitative iron and fat phantoms with the following goals: i) emulate morphological accuracy of particle size with respect to in vivo hepatic tissues; ii) assess the efficacy of mono-exponential, single-R2*, and dual-R2* fitting; iii) combine materials representing fat and iron to emulate co-occurring hepatic steatosis and iron overload.

Methods

Study Design

Fat Phantom

Fat-water emulsion phantoms were produced at fat fractions of 0%, 10%, 20%, 30%, and 40%. Previous studies claim that traditional methods produce fat droplets that are much greater than in vivo sizes [61]. To control the size of fat droplets, one set of phantoms was created using a stir bar for mixing while another set was mixed more vigorously with a handheld homogenizer to potentially reduce the size of fat droplets. A total of 10 phantoms were created (5 fat fractions x 2 sets).

Iron-Fat Phantom

Homogeneous fat-water-iron emulsion phantoms were produced at fat fractions of 0%, 10%, 20%, 30%, and 40% at increasing iron concentrations. Three different iron-oxide particles were utilized to emulate iron overload over a wide range of particle sizes: BNF (Bionized Nanoferrite, 0.08 μ m), MagnefyTM (0.85 μ m), and COMPELTM (3.3 μ m). A total of 40 phantoms were created (5 fat fractions x 5 iron concentrations x 3 sets).

Materials

- *Fat Source:* Peanut Oil (LouAna, Brea, CA, USA) was chosen to mimic hepatic fat for the Iron-Fat phantom study because it has a similar NMR proton spectrum as triglycerides in adipose tissue.
- Iron Nanoparticles: Three different iron nanoparticles were used to emulate iron storage molecules in this study. Bionized Nanoferrite-Starch (Micromod, Rostock, Germany) with a diameter of 80 nm were used to emulate the lower bounds of hemosiderin (0.1-3 µm [93,260]). Hemosiderin is typically observed to be closer to the micro-scale, so

Magnefy[™] (Bangs Laboratories, Fishers, Indiana) particles of 0.85µm and COMPEL[™] (Bangs Laboratories, Fishers, Indiana) particles of 3.3 µm were chosen to encompass larger scale hemosiderin proteins.

- *T1 Modulator:* The spin-lattice relaxation rate T1 was modulated by adding 0.5 mM anhydrous CuSO₄ powder (Alfa Aesar, Tewksbury, MA, USA) to decrease the T1 times of the phantom to mimic T1 times seen in human tissues.
- Base Components: Each phantom was suspended in a matrix composed of agar powder (Fisher Scientific, Pittsburgh, PA, USA) to create a 2% w/v agar gel. Within this suspension included 43 mM sodium dodecyl sulfate (SDS) (Fisher Bioreagents, Pittsburgh, PA, USA) as a surfactant and 43 mM sodium chloride (NaCl) (Fisher Chemical, Pittsburgh, PA, USA) as a preservative.

Construction

For both studies, a 2% water-agar solution (including SDS, NaCl, and CuSO₄) was heated to boil at 90°C and later allow to settle as a gel upon cooling to room temperature after the mixture was completed.

Fat Phantom

Appropriate volumes of peanut oil were allotted into separate beakers from the agar solution. When the temperature of the agar solution fell to an appropriate range (45-55°C) agar was dispensed into the allotted beaker to fill a 50 mL volume. Exceeding this temperature range runs the risk of causing precipitation of the oil (creaming). One set of phantoms was homogenized using a hand-held homogenizer (D1000, Benchmark Scientific, Sayreville, NJ, USA) with a 10mm x 115mm saw tooth generator probe at approximately 15000 rpm, while the second set was mixed with a stir bar and magnetic hot plate at 1400 rpm. After approximately 60

seconds of blending, the mixture of each beaker was poured into 50 mL vials and allowed to settle to a solid state.

Iron-Fat Phantom

Iron-fat-water emulsion phantoms were produced at fat fractions of 0%, 10%, 20%, 30%, and 40% and with varied concentrations of iron nanoparticles. Again, volumes of peanut oil were allocated to separate beakers complementary to the fat fraction needed. When the temperature of the agar solution fell to 45-55°C, agar was dispensed into the allotted beakers to fill a 50 mL volume. Additionally, iron nanoparticles were added with concentrations that match clinical R2* ranges (~0-1000s⁻¹) using a micro-pipettor. A hand-held homogenizer was used to control the size of the fat droplets and distribute the iron nanoparticles homogeneously in solution. After approximately 60 seconds of blending, the mixture of each beaker was poured into 50 mL vials and allowed to settle at room temperature.

Data Acquisition

Light Microscopy

During phantom creation, a 100 μ L sample of each phantom was extracted and placed on a glass microscopy slide. The fat droplet morphology of each phantom was then characterized at 200x using bright-field light microscopy with an Olympus BX63 microscope (Olympus, Waltham, MA, USA).

Scanning Electron Microscopy

Scanning electron microscopy (SEM) was used to quantify the size and distribution of the different iron nanoparticles used in this study. Each sample was sputter coated with a gold/palladium (80:20) coat using an EMS550x Sputter Coater (Quorum, Laughton, East Sussex, England) and mounted on 25 mm diameter aluminum specimen stubs. SEM analysis was

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performed with a Nova NanoSEM 650 (FEI Company, Hillsboro, Oregon, USA) at a voltage of 20.00 kV, spot size of 3.0, working distance of 5.5-5.2 mm, and a field-free lens mode. *MRI Analysis*

Magnetic resonance imaging was performed at 1.5T (MAGNETOM Avantofit, Siemens) using 2D GRE sequences with a monopolar gradient and 2D UTE sequences for iron and fat quantification. Single R2* correction was performed with NLSQ (Non-linear Least Squares) while the dual R2* model was implemented with ARMA (Autoregressive Moving Average) [4,44]. Imaging parameters used for the GRE sequence included: TE1 = 1.2 ms, echo spacing = 1.44 ms, TR = 200 ms, echo train length = 20, flip angle = 35° , 104×128 matrix, and 5 mm slices. Likewise, the imaging parameters of the UTE sequence are as follows: TE1 = 0.1 ms, echo spacing = 0.5 ms, TR = 172.2 ms, echo train length = 12, flip angle = 20° , 192×192 matrix, and 10 mm slices.

Data Processing and Statistical Analysis

Using MATLAB (2020b), segmentation algorithms based upon binary thresholding were incorporated to determine the radius and area of all fat droplets on light microscopy images.

MATLAB was also used to perform phantom image analysis by performing ROI selection, map generation, and R2* & fat fraction quantification. The code used was an adapted form of the Fat-Water toolbox provided by Hu et al. at ISMRM 2012 [261], Krafft et al. at ISMRM 2013 [177], and Tipineni-Sajja et al [44].

Results

Fat Phantom

The microscopy samples extracted from each steatosis phantom illustrated a considerable discrepancy in size and variability between the two mixing methods. Figure 11 displays a sample from a 40% FF stir bar and homogenized phantom paired against a liver biopsy sample of a patient with moderate steatosis. Visually, it appeared that the size and distribution of the fat globules in the stir bar phantom were more similar to the liver biopsy sample. However, statistical analysis was performed to confirm this assumption as shown on Table 12. Here, the droplet radii in the stir bar data more closely matched the biopsy sample than the homogenized data for the mean, median, and standard deviation. A one-way ANOVA and Tukey's HSD were performed and showed that the homogenized and biopsy data were significantly different with a p-value of 0.01989. Hence, if considering only particle size, the stir bar method would be most appropriate in matching in vivo size and distribution of fat globules.

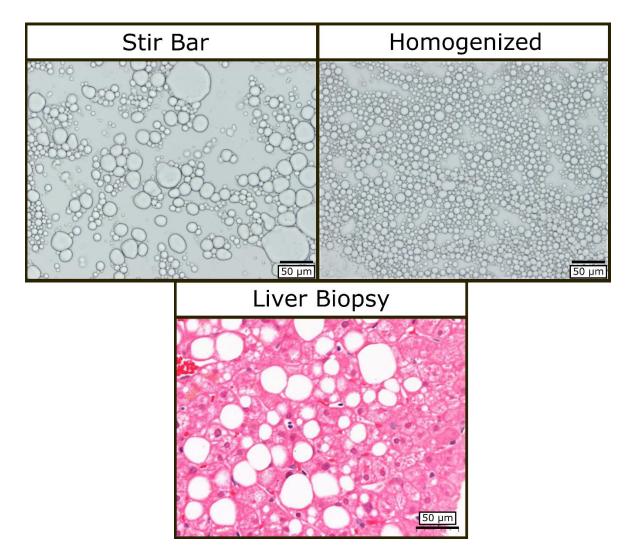


Figure 11: Light microscopy samples of a (top left) stir bar phantom, (top right) homogenized phantom, and (bottom) liver biopsy sample are shown. The phantom samples contain 40% FF while the biopsy sample is graded to be that of a moderately steatotic liver. The stir bar phantom is visually more similar to the biopsy sample that the homogenized phantom.

Table 12: Descriptive statistics results from lipid globules of steatosis phantoms mixed with a stir bar and homogenizer and liver biopsy samples from patients with mild to severe steatosis. The stir bar data more closely aligned with the mean, median, and standard deviations of the biopsy sample than the homogenized data.

Radius (µm)	Stir Bar	Homogenized	Biopsy
Mean	5.563	3.445	5.581
Median	5.629	3.405	4.957
St. Dev.	1.162	0.236	1.519

After evaluating the phantoms via microscopy, MRI analysis was performed. Figure 12 displays FF plots and maps of the steatosis phantoms mixed with a stir bar and homogenizer. Here, it can be seen that the theoretical and experimental FF values of both the stir bar and homogenized phantoms matched closely. However, the NLSQ model slightly underestimated the homogenized phantom FF while ARMA slightly overestimated the stir bar phantom FF. Accordingly, the FF plots reveal that ARMA FF maps were more inhomogeneous than NLSQ and therefore created more standard deviation of FF values in the plots. Overall, the difference in FF signal behavior between both mixing methods is negligible regardless of fitting method.

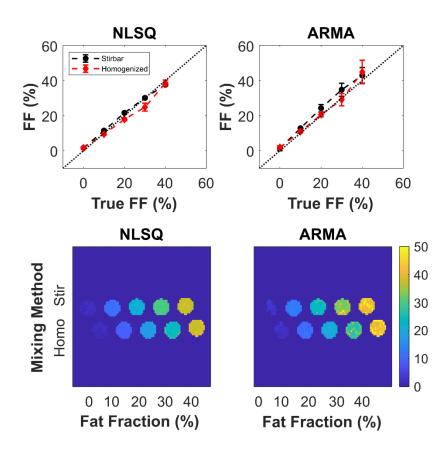


Figure 12: FF plots (top) and maps (bottom) of steatosis phantoms mixed with a stir bar and homogenizer with increasing FF values (0%, 10%, 20%, 30%, and 40%) using NLSQ and ARMA fitting models. The theoretical and experimental FF values of the phantoms matched well.

R2* was also calculated for these phantoms as illustrated in the plots and maps in Figure 13. Here, it can be seen that increasing the FF generally increases R2*. Additionally, the stir bar method produces R2* values slightly higher than the homogenized method, though this is not always the case. All three fitting models largely agree on the R2* values, unlike the FF maps previously.

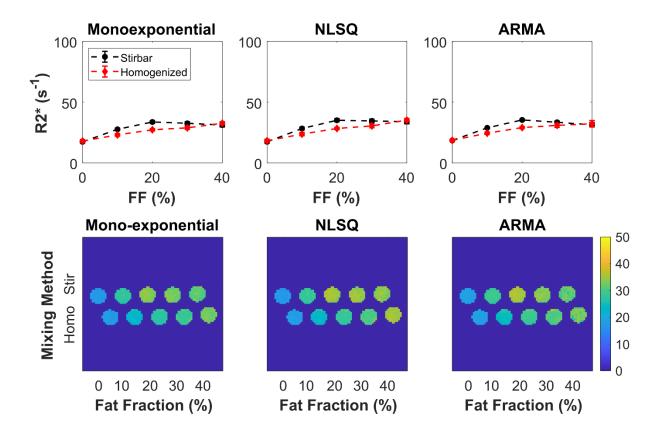


Figure 13: R2* plots (top) and maps (bottom) of steatosis phantoms mixed with a stir bar and homogenizer with increasing FF values (0%, 10%, 20%, 30%, and 40%). All three fitting models (monoexponential, NLSQ, and ARMA) largely agreed on the R2* values.

Iron-Fat Phantom

Scanning electron microscopy confirmed the sizes of the Compel, Magnefy, and BNF particles as per manufacturer specifications. Figure 14 displays each of these three particles at 25000x to further emphasize the difference in size. The Compel and BNF particles tended to distribute homogenously over the slide mount, but the Magnefy particles were more spaced out

and had smaller fragments surrounding them. Element analysis confirmed the presence of iron in each of these particles, including the fragmented Magnefy particles.

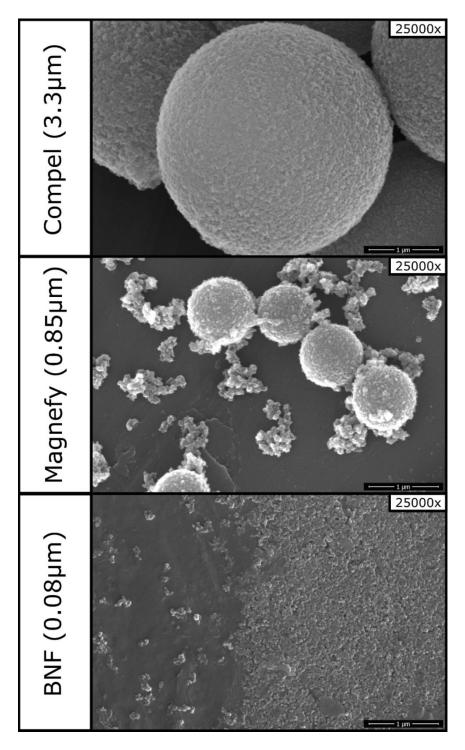


Figure 14: Scanning electron microscopy images of Compel, Magnefy, and BNF nanoparticles are shown at 25000x. Each particle type is noticeably different in size and matches manufacturer specifications.

After verifying the size of each iron particle, all three iron phantom sets were scanned using a GRE sequence and subsequently analyzed using ROI analysis in MATLAB. Figure 15 displays R2* plots for all three particle types using monoexponential, NLSQ, and ARMA models with a GRE sequence. Here, the Compel (3.3μ m) particle delivered the most consistent results with higher FF's producing greater R2* values as expected. However, Magnefy (0.85μ m) and to a lesser degree, BNF (0.08μ m), yielded R2* values that increased with iron concentration, but increasing FF did not consistently increase R2* in a linear manner. For the models, ARMA produced the most linear results with the least number of inconsistencies. Mono-exponential evaluated the Compel phantoms well but underperformed on the BNF and Magnefy phantoms. NLSQ completely fails to fit values above $500s^{-1}$ and overestimates the R2* values with large standard deviations.

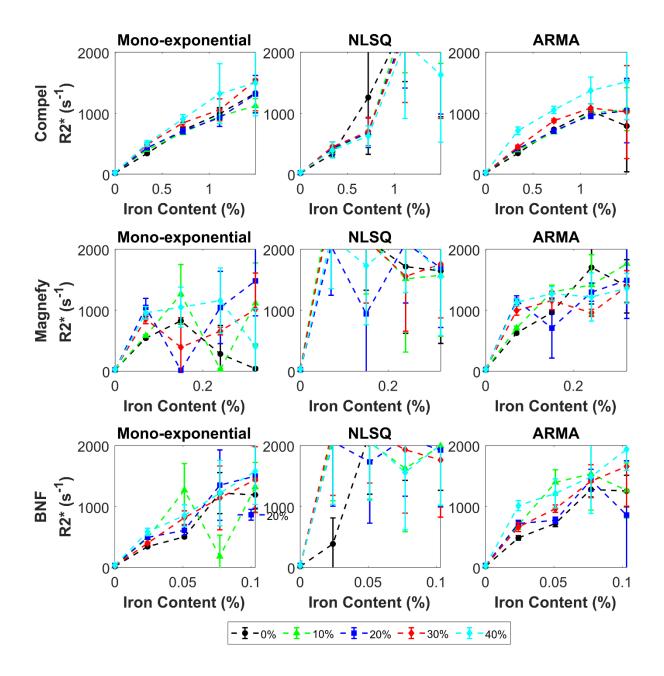


Figure 15: R2* plots of iron-fat phantoms with increasing iron concentrations and increasing FF values (0%, 10%, 20%, 30%, and 40%) are shown. These phantoms were analyzed with a GRE sequence. The three different iron particles used are as follows: Compel (top), Magnefy (middle), and BNF (bottom).

Iron-only phantoms (0% FF) were also of interest to investigate iron-water molecule interactions, so phantoms of each iron type with no oil added were analyzed separately as seen on the plots of Figure 16. Here, the Compel and BNF phantoms generated R2* values that were relatively linear. However, mono-exponential performed better for the Compel phantoms, while ARMA was better for BNF. Conversely, the Magnefy phantoms did not show a linear trend for R2* and increasing iron in all three models as each failed to quantify the phantom with the highest iron. Notably, NLSQ failed to accurately calculate the R2* in all three phantom sets despite the absence of fat.

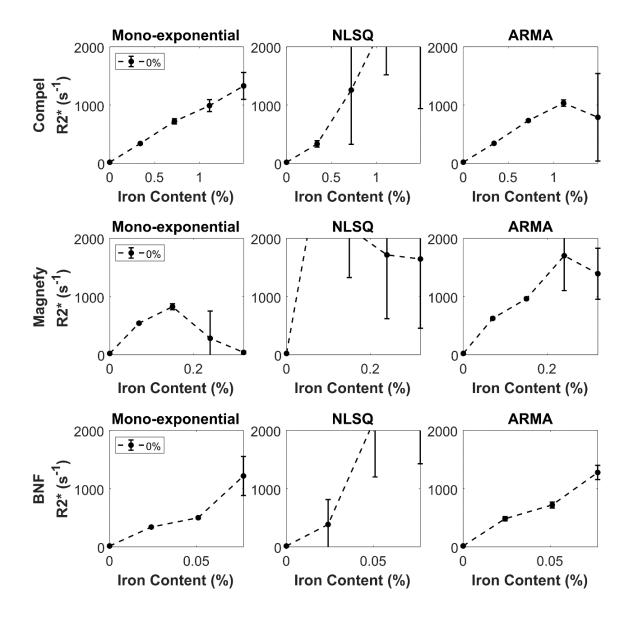


Figure 16: R2* plots of iron-only phantoms with increasing iron concentrations are shown. A GRE sequence was used. The three different iron particles used are as follows: Compel (top), Magnefy (middle), and BNF (bottom). Due to being an outlier, the BNF phantom with the highest concentration was excluded.

To get a top-down view, visual R2* maps were generated from each of these iron-fat phantoms and are displayed on Figure 17. Here, the inhomogeneities in each phantom set are more apparent, but the data still show the trends seen from Figure 15. Mono-exponential performed relatively well, especially for the Compel phantoms, but tended to outright fail to quantify R2* in some BNF and Magnefy phantoms. NLSQ successfully quantified the Compel phantoms, but completely overestimated R2* values in BNF and Magnefy phantoms that had iron in them. Finally, ARMA produced the most homogeneous results of all three models but still has inaccuracies as the iron concentration goes higher.

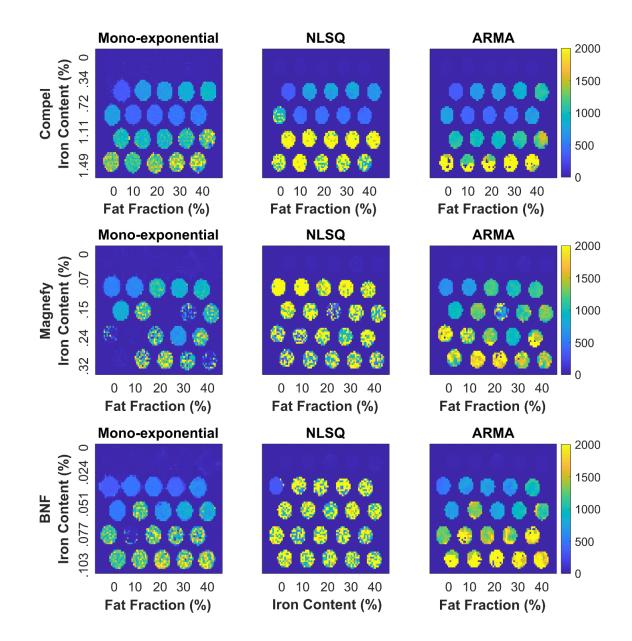


Figure 17: R2* maps of iron-fat phantoms with increasing iron concentrations and increasing FF values (0%, 10%, 20%, 30%, and 40%) are shown. These phantoms were analyzed with a GRE sequence. The three different iron particles used are as follows: Compel (top), Magnefy (middle), and BNF (bottom).

While GRE sequences are traditionally used for quantifying iron and fat, recent literature has investigated the potential for using UTE sequences to accurately quantify extremely high iron contents. Because the highest iron concentrations produced poor results with the GRE sequence, a UTE sequence was also utilized on the iron-fat phantoms. Figure 18 displays R2*

plots for all three particle types using mono-exponential, NLSQ, and ARMA models with a UTE sequence. All three iron-fat phantoms and fitting models performed much better for UTE acquisition, especially NLSQ model. The Compel phantoms still displayed the most linear and consistent results, and this phantom was the only one that correctly illustrated the known phenomenon of increasing FF also increasing the R2* value. UTE acquisition demonstrated substantial improvements for BNF phantoms in showing R2* linearity for the mono-exponential and NLSQ models, but there were few outliers for the 10% FF phantoms. The UTE sequence also revealed that the Magnefy phantoms reached a higher R2* value (~2500s⁻¹) than the Compel and BNF phantoms (1500s⁻¹), which may play a part in the larger standard deviations seen in this phantom. For the models, the mono-exponential and ARMA models agreed closely for most phantoms, but the NSLQ model produced more inconsistencies, albeit with similar trends.

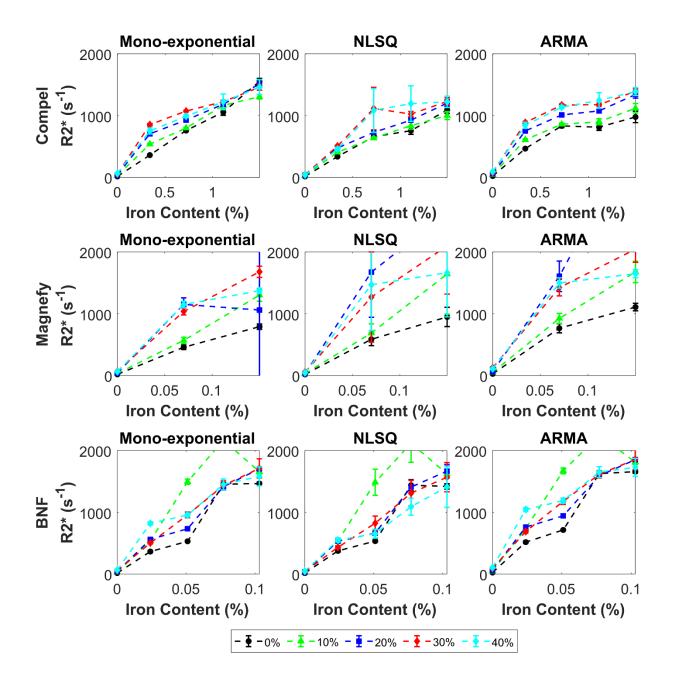


Figure 18: R2* plots of iron-fat phantoms with increasing iron concentrations and increasing FF values (0%, 10%, 20%, 30%, and 40%) are shown. These phantoms were analyzed with a UTE sequence to accurately quantify phantoms with high iron contents. The three different iron particles used are as follows: Compel (top), Magnefy (middle), and BNF (bottom).

Again, iron-only phantoms were independently investigated to analyze iron-water molecule interactions as seen on the plots of Figure 19, with a UTE sequence. The most notable

difference is the drastically improved R2* linearity for the Magnefy phantoms, though all three phantom sets improved. NLSQ and ARMA calculated that the two phantoms of the highest Compel concentration were lower than predicted by the mono-exponential model but also with more standard deviation. However, each model largely agreed on the values for the BNF and Magnefy phantoms which were not as linear, but still showed increasing R2* with increasing iron concentrations. Overall, NLSQ displayed the most uncertainty.

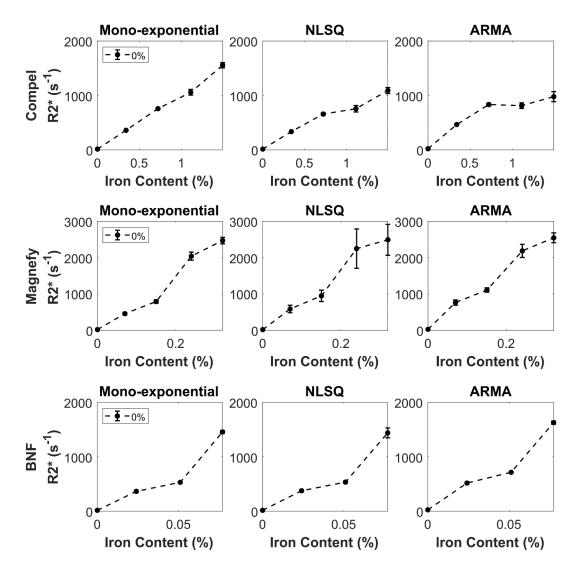


Figure 19: R2* plots of iron-only phantoms with increasing iron concentrations are shown. A UTE sequence was used to accurately quantify phantoms with high iron contents. The three different iron particles used are as follows: Compel (top), Magnefy (middle), and BNF (bottom). Due to being an outlier, the BNF phantom with the highest concentration was excluded.

In addition to the plots, UTE R2* maps were generated to improve visibility of the phantoms and are displayed on Figure 20. Here, some outliers are more apparent such as BNF (0.077%, 10% FF) and Magnefy (0.15%, 20% FF). With the UTE sequence, the trend of R2* increasing with increasing iron is more evident, and the increase in R2* with increasing FF is negligible in comparison. Overall, these maps produce ROI's that are much more homogenous compared to GRE counterparts.

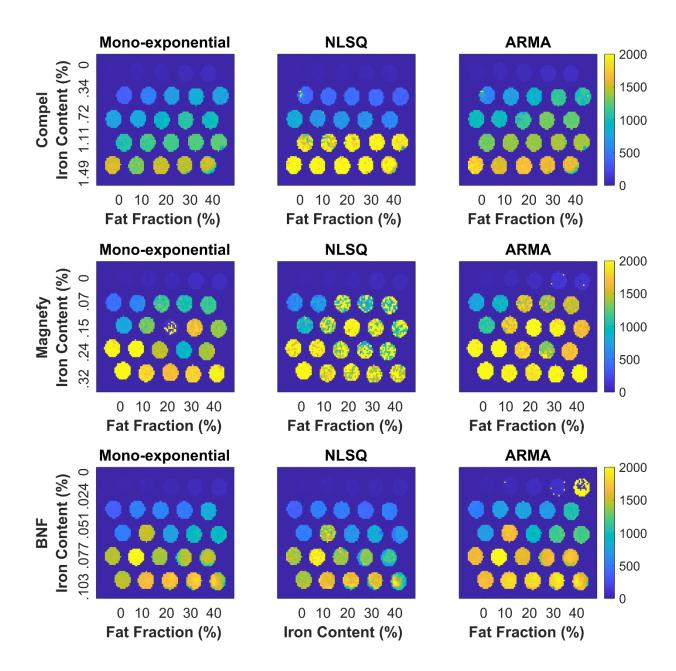


Figure 20: R2* maps of iron-fat phantoms with increasing iron concentrations and increasing FF values (0%, 10%, 20%, 30%, and 40%) are shown. These phantoms were analyzed with a UTE sequence. The three different iron particles used are as follows: Compel (top), Magnefy (middle), and BNF (bottom).

UTE sequences have only been validated in R2* fitting models based on magnitude data [42]. Therefore, complex-based multispectral fat models like ARMA may be unsuitable for mapping FF data. Because of this, GRE sequences were used for the plots seen in Figure 21. In

the Compel phantoms, the quantification of FF becomes increasingly worse as iron increases. At high iron concentrations, NLSQ overestimates FF while ARMA underestimates FF, both of which phenomena have been observed in the literature before [44]. These effects are more extreme in the BNF and Magnefy phantoms in which both models fail to quantify FF in phantoms with any iron present.

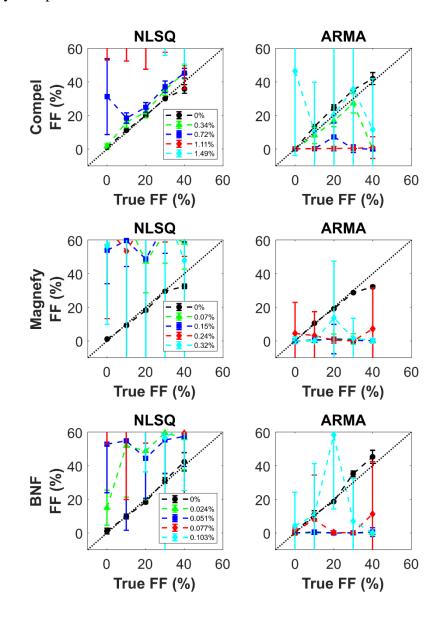


Figure 21: FF plots of iron-fat phantoms with increasing iron concentrations and increasing FF values (0%, 10%, 20%, 30%, and 40%) are shown. These phantoms were scanned with a GRE sequence and analyzed with NLSQ and ARMA fitting models. The three different iron particles used are as follows: Compel (top), Magnefy (middle), and BNF (bottom).

In addition to the FF plots, FF maps were generated to highlight areas of inhomogeneities and estimation failures in Figure 22. Here, it can be seen that ARMA either completely fails to quantify most phantoms or severely overestimates (by > 50% FF) regions of high iron phantoms. The exception to this was the row of 0.34% iron Compel phantoms in which most had an accurate quantification of FF. NLSQ had moderate success in quantifying the Compel phantoms up to 0.72% iron but failed to properly read iron phantoms from the BNF and Magnefy batches. However, both models accurately quantified phantoms without iron, but NLSQ performed better than ARMA on all other phantoms.

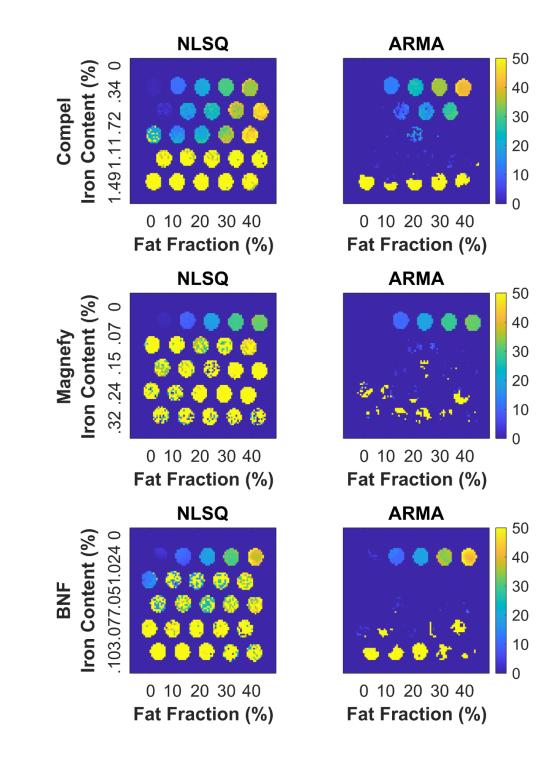


Figure 22: FF maps of iron-fat phantoms with increasing iron concentrations and increasing FF values (0%, 10%, 20%, 30%, and 40%) are shown. These phantoms were scanned with a GRE sequence and analyzed with NLSQ and ARMA fitting models. The three different iron particles used are as follows: Compel (top), Magnefy (middle), and BNF (bottom).

Discussion

In this study, MRI analysis of steatosis and iron overload phantoms was performed using GRE and UTE sequences to investigate the effects of particle size on the MR signal as well as the efficacy of MR fitting models. Evidence of particle size significantly impacting the signal came back inconclusive. Mixing via stir bar and homogenization revealed that stir bar lipid globules matched biopsy data in size and distribution more closely than homogenization, yet the difference in MR signal between both methods was negligible. For iron, SEM analysis validated the sizes listed by the manufacturer for each iron particle type. However, there was no apparent trend of particle size impacting MRI performance. Out of the fitting models, ARMA performed the most optimally for R2* quantification while using a UTE sequence. Conversely, running the NLSQ model with a GRE sequence was preferred for FF quantification.

In the fat phantom, it was found that the stir bar mixing method produced droplets that matched liver biopsy lipid globules more closely in size and distribution than homogenization. However, MRI analysis showed that the difference in fat-water signal behavior between the stir bar and homogenized phantoms was negligible. It is not known if the size and distribution of differing fat droplet sizes would affect the signal behavior of iron on fat differently. The iron-fat phantoms produced for this study were created exclusively with the homogenization method, which was shown to generate fat droplets that were of a more consistent size. Hence, it is possible that wider variations in fat droplets could in turn affect iron-fat interactions which were only partially investigated in this study. Ideally, another batch of phantoms could be created with the stir bar method to be used in comparison against homogenization. However, this is logistically impossible as iron will stick to the magnetic stir bar instead of remaining in solution. Therefore, alternate mixing methods would have to be considered in future work.

From the iron-fat phantoms, there is no clear indication that size of iron particles will noticeably affect the MR signal. Compel (3.3μ m) was the largest particle used and produced the most homogeneous values for both R2* and FF out of all the iron particle types. However, BNF (0.08μ m) outperformed the larger Magnefy (0.85μ m) particles despite a tenfold difference in size. Nonetheless, there is a case to be made about the distribution of the particles affecting the signal as Compel and BNF were consistently seen to be of the same shape and size via SEM, and they dispersed homogeneously over the slide. On the other hand, the Magnefy particles had smaller fragments located alongside particles of the proper size that also contributed to the signal produced by iron. Consequently, it is possible that the inhomogeneities seen in the Magnefy phantom are partially caused by inconsistencies in particle size.

Traditionally, clinical R2* values reach up to ~1000 s⁻¹ but can go as high as 2000s⁻¹ [35,44,262]. The Compel and BNF phantoms sat adequately in this range by reaching ~1500 s⁻¹. However, the Magnefy phantoms overshot this with a maximum R2* value of 2492 s⁻¹. While theoretically possible in patients, an R2* value this high would be quite rare. Hence, the concentration of Magnefy particles would be cut down in future work to both satisfy an appropriate clinical range and allow for a better comparison to the Compel and BNF particles in phantoms.

Traditionally, T2*-weighted GRE sequences have been employed to image iron via MRI. Because GRE sequences require a highly uniform magnetic field (B₀), any local field inhomogeneities, such as that produced by superparamagnetic iron, will greatly affect the signal output [263]. In short, GRE is very sensitive to iron. However, in cases of severe iron overload, the precision of GRE-based estimations of iron may fail as the signal decays too quickly to be reliably measured [42]. One reason for this failure is the inability of conventional GRE to image

echo times (TE) shorter than 1.0 ms, in which a significant degree of the MR signal may already be lost [42]. Alternatively, UTE sequences can reach TE values as low as 0.1-0.19 ms, thus can measure extremely short-lived signals [42].

As stated previously, UTE sequences have only been validated in R2* fitting models like NLSQ which are based on magnitude data, so models that use complex data such as ARMA may be unsuitable for fitting FF data. Traditionally, UTE sequences utilize fat suppression pulses to reduce streaking artifacts that make arise from subcutaneous fat [42]. However, the UTE sequence used in this study did not have fat suppression built into the sequence, and therefore could theoretically be used for imaging FF maps. Indeed, NSLQ could quantify FF when tested with the UTE sequence; however, ARMA completely failed to quantify FF from all phantoms that contained iron, likely due to low SNR.

From the results of the FF quantification with the GRE sequence, it was seen that NLSQ performed better than ARMA, but it tended to overestimate FF at high iron concentrations. NLSQ has multiple assumptions regarding the fat peaks built into the model as priori information. These assumptions will persist even when no fat is present or under rapid signal decay induced by iron ($R2^* > 500 \text{ s}^{-1}$) which may interfere with the quantification of parameters that produce the smallest least square error, therefore leading to false FF values [44]. While NLSQ did successfully quantify the FF of the Compel phantoms of up to ~800 s⁻¹, significant error was displayed for all other phantoms. ARMA behaved similarly as the phantoms without iron or low iron were accurately quantified, but all other phantoms failed. This is likely because ARMA has no assumptions about the fat peaks built into the model, thus it fails in the presence of a low signal. Hence, NSLQ performed slightly better than ARMA.

R2* quantification revealed that the Compel particles followed the most linear distribution out of the three iron particle types, and this linearity is phenomenon that is also seen in the human liver [262]. Out of the fitting models, mono-exponential and ARMA fit the iron-fat phantoms more linearly than NLSQ, with mono-exponential model performing slightly better than ARMA. However, the mono-exponential model does not consider the presence of fat [2], so ARMA may be constructing the signal behavior more accurately despite the difference in linearity. NLSQ suffers from the same issues as before, as increasingly high iron in tandem with high FF causes more inhomogeneities in R2* quantification. Because of this, some investigators have tried using the NLSQ model to measure R2* without incorporating priori information on the fat peaks [264]. Because our phantoms feature fat, this was not a feasible alternative.

Conclusion

Despite controlling the size of fat and iron particles in steatosis and iron overload MRI phantoms, evidence showing that particle size impacts the signal to a significant degree remains inconclusive. Reproducing these results with different mixing methods and utilizing particles with homogeneous distributions may be necessary to give a definitive conclusion. UTE sequences improved the estimation of R2* in the presence of fat, but GRE sequences are still necessary for accurate FF quantification in the presence of iron. All three fitting models had their merits with R2* quantification most accurately being portrayed under the mono-exponential model under conditions of no fat and ARMA under conditions of iron and fat. Additionally, NLSQ handled FF quantification more consistently than ARMA. This study demonstrates different sequencing and post-processing assessments are critical for the analysis of sensitive biomarkers such as R2* and FF.

Chapter V

Conclusion and Future Work

Clinical quantitative MRI techniques are at their infancy, yet they provide non-invasive and sensible data that rival established methodologies such as biopsy. This work imitates realistic MRI signal behavior and clinical ranges of biomarkers in imaging phantoms while simultaneously addressing discrepancies between phantom media particle size and molecular interactions inside in vivo tissues. Ultimately, definitive conclusions regarding the impact of particle size on the MR signal cannot be drawn from this study, but several improvements in the design may reveal more solutions.

While the preceding analysis was theoretically sound, there are a few key changes that could be made to the experimental setup to improve upon the validity of this work. This study was fairly limited in scope to the available mixing methods and iron particles available on the market. To control the size of fat droplets, a magnetic stir bar and hand-held homogenizer were used as they are the most prevalent instruments used in hepatic phantom literature. However, there have been reports of other methods being used to develop phantoms [61,265]. In future studies, several more mixing instruments and methodologies could be used to compare against liver biopsy samples, including gentle inversion, overhead stirring, and ultrasonication. Also, while no apparent trend could be observed on the MR signal due to particle size, the polymer coating of each type of iron particle contained differing compositions and thus could have impacted internuclear interactions. Ideally, the same coating should be used for each size as the varying polymers may have differing solubilities, crosslinking agents, and charges, among other variables. Unfortunately, commercial particles that both covered the clinical ranges of hemosiderin and were coated with the same polymer could not be located for this study. Future

studies could either synthesize particles with a wide range of sizes and a consistent polymer composition or utilize commercial particles that meet these conditions.

In this study, there are several more pathologically relevant features that could be incorporated into the phantom design. As stated previously, iron overload is a heterogeneous disease with a mottled distribution across the liver. The severity of hepatic iron overload also impacts the amount of ferritin converted to hemosiderin with more hemosiderin forming in severe iron overload. Hence, the particle size of iron overload phantoms should theoretically reflect the ratios of ferritin to hemosiderin if preserving the integrity of tissue emulation. The effect this may have on the MRI signal may be minuscule, but could be investigated, nonetheless. Additionally, this work was primarily aimed at studying hepatic iron overload and steatosis as they often co-occur, and the resulting MR signal from both pathologies are known to confound quantification of each other. However, fibrosis often results from advanced cases of these diffuse liver diseases and can be traced with other MR biomarkers such as elastography or T1w imaging, so incorporating this pathology into combination iron overload and steatosis phantoms may be worthwhile.

In summary, more mixing methods along with a more consistent polymer coating could reveal additional insights on the impact of particle size in steatosis and iron overload phantoms, all of which could be expanded on to further mimic other hepatic tissues and severities.

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Appendices

Table 6. One-way ANOVA results from lipid globules of steatosis phantoms mixed with a stir bar and homogenizer and liver biopsy samples from patients with mild to severe steatosis. A significant difference between groups was found at a p-value of 0.02587.

/	l			
	Stir Bar	Homogenized	Bio	psy
Radius (µm):	6.859	3.204	4.062966401	4.780511095
	6.028	3.362	4.702170354	4.355492429
	5.23	3.447	7.268451196	6.665140275
	4.133	3.765	4.197894114	6.56897126
			4.441882233	4.675188192
			4.82505242	4.72726259
			9.224104445	3.846604582
			5.674506909	8.78644816
			4.372134251	4.026578345
			5.596148197	4.567648516
			7.599731753	5.707555069
			4.525249029	6.350870351
			5.186230513	5.947062098
			5.783878949	6.360781628
			4.234846866	4.442131402
			5.090082197	4.811299563
			9.785633706	7.657573946
			5.522910222	4.562867232
Skewness:	-0.287874	0.933369	1.29	2792
Excess	-0.596959	1.4319	1.07	0118
Kurtosis:	-0.590959	1.4519	1.07	0118
Normality:	1	0.9533	0.000)2731
Outliers:			9.224104445	9.785633706
Mean:	5.5625	3.4445	5.5	815
S:	1.16213	0.2362	1.51	1874

Source	DF	Sum of Square	Mean Square	F Statistic	P-value
Groups (between groups)	2	16.5782	8.2891	4.0007	0.02587
Error (within groups)	41	84.9488	2.0719		
Total	43	101.5269	2.3611		

ANOVA

Table 7. Tukey's HSD results from lipid globules of steatosis phantoms mixed with a stir bar and homogenizer and liver biopsy samples from patients with mild to severe steatosis. A significant difference was found between the biopsy and homogenized groups at a p-value of 0.01989.

Pair	Difference	SE	Q	Lower Cl	Upper Cl	Critical Mean	p-value
Stir Bar- Homogenized	2.118	0.7197	2.9429	-0.357	4.593	2.475	0.1064
Stir Bar - Biopsy	0.019	0.5364	0.03541	-1.8258	1.8637	1.8447	0.9997
Biopsy - Homogenized	2.137	0.5364	3.9837	0.2922	3.9817	1.8447	0.01989

Tukey's HSD

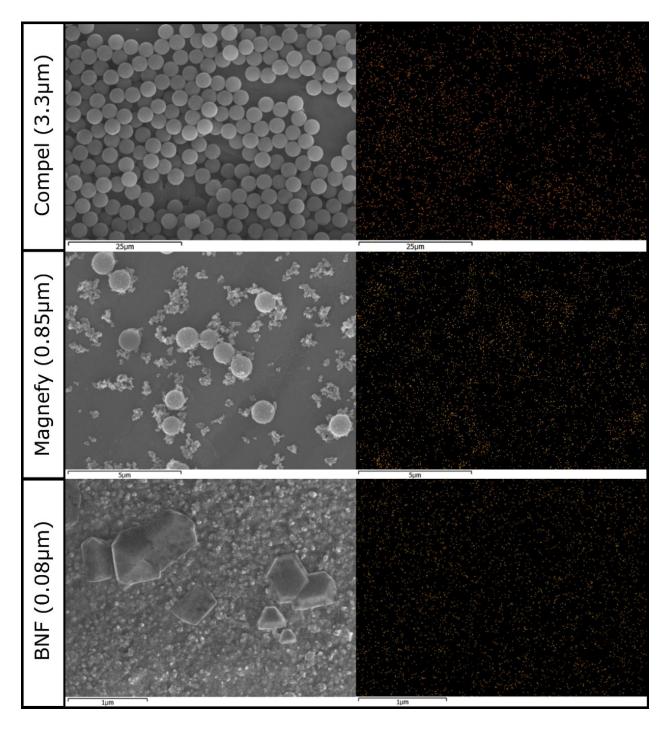


Figure 23. Scanning electron microscopy images of (top left) Compel, (middle left) Magnefy, and (bottom left) BNF iron nanoparticles are shown. Element analysis (right) was performed on each particle to confirm the presence of iron in each sample. The distribution of iron in the element analysis matches the shape of the nanoparticles, thus demonstrating the presence of iron. Notably, the large crystals shown the BNF image are composed of NaCl.

Table 8. Concentrations of oil and iron particles needed to make iron-fat phantoms of BNF concentrations 0%, 0.024%, 0.051%, 0.077%, & 0.103% and fat fractions of 0%, 10%, 20%, 30%, & 40%.

BI	NF	Oil Needed					
		0% FF	10% FF	20% FF	30% FF	40% FF	
Iron Needed	0%	<mark>0 mL</mark> 0 μL	4.6 mL 0 μL	9.2 mL 0 μL	13.8 mL 0 μL	18.4 mL 0 μL	
	0.024%	<mark>0 mL</mark> 70.3 μL	4.6 mL 70.3 μL	9.2 mL 70.3 μL	13.8 mL 70.3 μL	18.4 mL 70.3 μL	
	0.051%	<mark>0 mL</mark> 146.8 μL	<mark>4.6 mL</mark> 146.8 μL	<mark>9.2 mL</mark> 146.8 μL	13.8 mL 146.8 μL	<mark>18.4 mL</mark> 146.8 μL	
	0.077%	<mark>0 mL</mark> 223.3 μL	4.6 mL 223.3 μL	<mark>9.2 mL</mark> 223.3 μL	13.8 mL 223.3 μL	18.4 mL 223.3 μL	
	0.103%	<mark>0 mL</mark> 299.8 μL	<mark>4.6 mL</mark> 299.8 μL	<mark>9.2 mL</mark> 299.8 μL	<mark>13.8 mL</mark> 299.8 μL	<mark>18.4 mL</mark> 299.8 μL	

Table 9. Concentrations of oil and iron particles needed to make iron-fat phantoms of Magnefy concentrations 0%, 0.07%, 0.15%, 0.24%, & 0.32% and fat fractions of 0%, 10%, 20%, 30%, & 40%.

Magnefy		Oil Needed					
		0% FF	10% FF	20% FF	30% FF	40% FF	
	0%	<mark>0 mL</mark> 0 μL	4.6 mL 0 μL	9.2 mL 0 μL	13.8 mL 0 μL	18.4 mL 0 μL	
Iron Needed	0.07%	<mark>0 mL</mark> 70 μL	4.6 mL 70 μL	9.2 mL 70 μL	13.8 mL 70 μL	18.4 mL 70 μL	
	0.15%	<mark>0 mL</mark> 153.3 μL	<mark>4.6 mL</mark> 153.3 μL	<mark>9.2 mL</mark> 153.3 μL	13.8 mL 153.3 μL	18.4 mL 153.3 μL	
	0.24%	<mark>0 mL</mark> 236.6 μL	<mark>4.6 mL</mark> 236.6 μL	<mark>9.2 mL</mark> 236.6 μL	13.8 mL 236.6 μL	18.4 mL 236.6 μL	
	0.32%	<mark>0 mL</mark> 320 μL	<mark>4.6 mL</mark> 320 μL	<mark>9.2 mL</mark> 320 μL	13.8 mL 320 μL	18.4 mL 320 μL	

Table 10. Concentrations of oil and iron particles needed to make iron-fat phantoms of Compel concentrations 0%, 0.34%, 0.72%, 1.11%, & 1.49% and fat fractions of 0%, 10%, 20%, 30%, & 40%.

Compel		Oil Needed					
		0% FF	10% FF	20% FF	30% FF	40% FF	
	0%	<mark>0 mL</mark> 0 μL	4.6 mL 0 μL	9.2 mL 0 μL	13.8 mL 0 μL	18.4 mL 0 μL	
Iron Needed	0.34%	<mark>0 mL</mark> 338.4 μL	4.6 mL 338.4 μL	<mark>9.2 mL</mark> 338.4 μL	13.8 mL 338.4 μL	18.4 mL 338.4 μL	
	0.72%	<mark>0 mL</mark> 723 μL	4.6 mL 723 μL	<mark>9.2 mL</mark> 723 μL	13.8 mL 723 μL	18.4 mL 723 μL	
	1.11%	<mark>0 mL</mark> 1107.6 μL	<mark>4.6 mL</mark> 1107.6 μL	<mark>9.2 mL</mark> 1107.6 μL	13.8 mL 1107.6 μL	<mark>18.4 mL</mark> 1107.6 μL	
	1.49%	<mark>0 mL</mark> 1492.3 μL	<mark>4.6 mL</mark> 1492.3 μL	<mark>9.2 mL</mark> 1492.3 μL	<mark>13.8 mL</mark> 1492.3 μL	<mark>18.4 mL</mark> 1492.3 μL	