

### Assignment 10

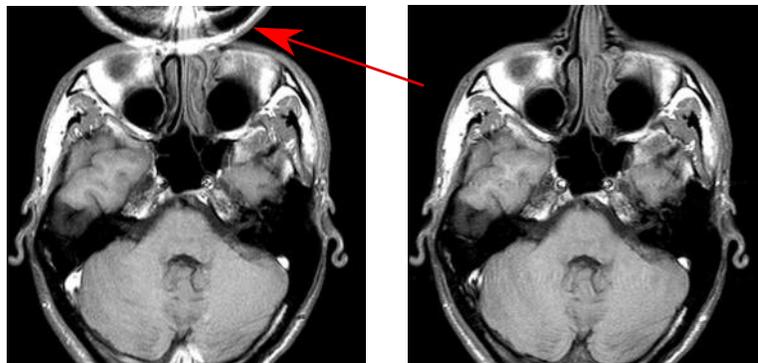
Due April 20th, 2012

This is the last homework in class. Enjoy it.

#### 1) MRI Artifacts and Debugging

In this question you will be given several pairs of images. The images may exhibit artifacts which will be indicated by arrows/circles. Answer the questions as best as you can. First describe the artifact and then provide as much information as possible on the source/s of the artifact and why it appears this way. There might be several possible right answers. A final answer without explanation will not be credited.

- a) Example: What is the source of the artifact in the left image and what is the difference in the acquisition/processing of the two?



*Solution:*

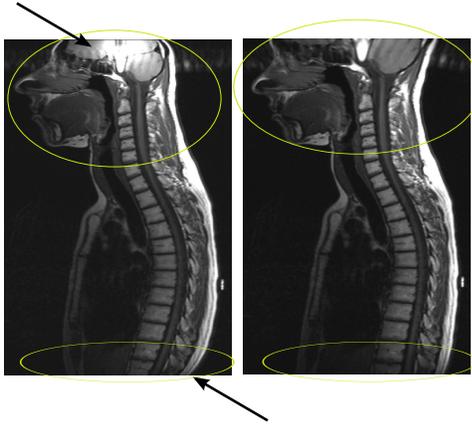
The artifact in the image appears as the back of the brain aliasing over and wrapping to the front. This is typical when the prescribed FOV in the phase encode is smaller than the FOV of the object. In that case the sampling density does not meet the Nyquist rate. The phase encode is obviously Anterior-Posterior.

The image on the right does not exhibit aliasing. One way to avoid aliasing is to increase the number of phase encodes while keeping the resolution the same. The other way is to swap the readout and phase-encode directions. It seems that the latter approach of swapping the readout direction was used since the back of the skull as well as the nose are cropped. Choosing a larger FOV in the phase encode would have shown the entire nose and skull.

Source of Artifact: aliasing in the phase-encode direction (A/P)

Difference: readout and phase encode directions swapped

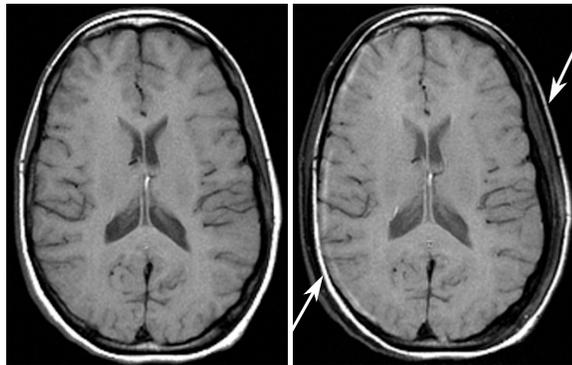
- b) What is the source of the artifact in the left image and what is the difference in the acquisition/processing of the two?



Source of artifact:

Difference:

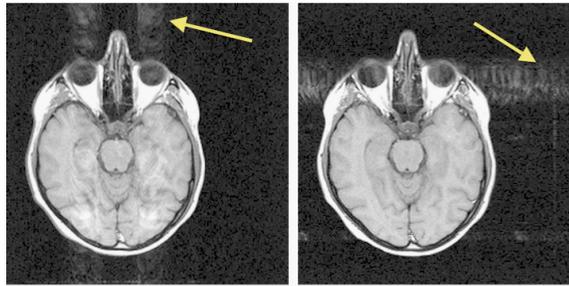
- c) What is the source of the artifact in the right image and what is the difference in the acquisition/processing of the two?



Source of artifact:

Difference:

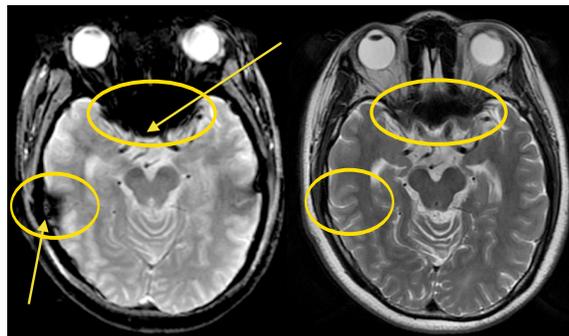
- d) What is the source of the artifact in the two images and what is the difference in the acquisition/processing of the two?



Source of artifact:

Difference:

e) What is the source of the artifact in the left image and what is the difference in the acquisition/processing of the two? (Ignore the contrast difference between the two)

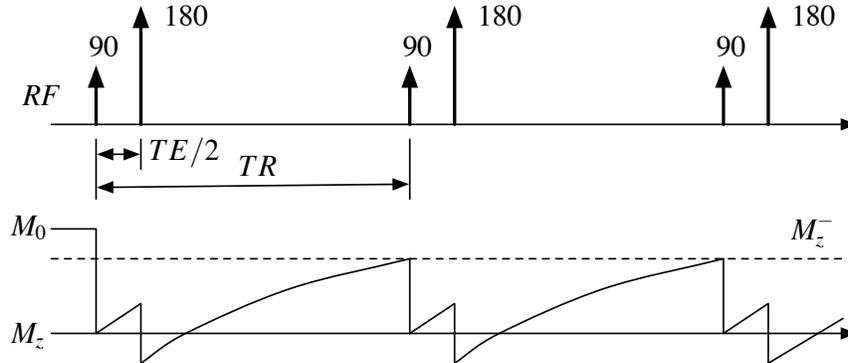


Source of artifact:

Difference:

- 2) In class when we presented the spin-echo saturation recovery pulse sequence, we assumed that the echo time  $TE$  was short relative to the repetition time  $TR$ , so we could ignore it. This is not always a good approximation, particularly if we want to measure  $T_1$ .

In fact, after the 90 there will be some  $M_z$  recovery. The 180 will invert this, and  $M_z$  will continue to recover until the next 90. Let  $M_z^-$  be the magnetization immediately before the 90.



Assume that  $T_E$  is small compared to  $T_1$ , but not negligible, so that we can approximate

$$(1 - e^{-T_E/T_1}) \simeq T_E/T_1.$$

This means that the  $M_z$  recovery curves are approximately linear within a time  $T_E$  after the 90.

- Find an approximate expression for the time when the magnetization cross the  $M_z = 0$  axis.
  - This might be called a  $T_1$  echo. Why?
  - Find an expression for  $M_z^-$  using what you found in (a).
  - Compare  $M_z^-$  calculated using your expression and the expression ignoring  $T_E$  for the case where  $T_R = 200$  ms,  $T_E = 20$  ms, and  $T_1 = 800$  ms. What is the percent error produced by neglecting  $T_E$  in this case?
- 3) Choosing Scan Parameters (We will cover material for this on Tuesday.)

You are designing a pulse sequence to image the upper abdomen. The tissues of interest are

- Liver,  $T_1 = 600$  ms,  $T_2 = 50$  ms.
- Spleen,  $T_1 = 1000$ ms ,  $T_2 = 80$  ms.
- Fat,  $T_1 = 350$  ms,  $T_2 = 60$  ms.
- Gall Bladder,  $T_1 = 2000$  ms,  $T_2 = 300$  ms.

This will be a spin echo acquisition, where you have to choose the repetition time  $T_R$ , the echo time  $T_E$ . In addition, you can also add an inversion recovery pulse *if you need it*, and specify the inversion time  $T_I$ . Otherwise leave the  $T_I$  blank. The minimum  $T_E$  is 15 ms, and the minimum  $T_R$  is 20 ms.

In each case you only need to specify the scan parameters *approximately*. You only need reasonable values, not necessarily the optimum values. However, you need to describe the reasoning behind your choices.

- a) Choose the scan parameters so that you have good contrast between liver and spleen, and low signal from the gall bladder.

$T_R =$	$T_E =$	$T_I =$
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- b) Choose the scan parameters so that you see only the gall bladder, with little signal from the other tissues.

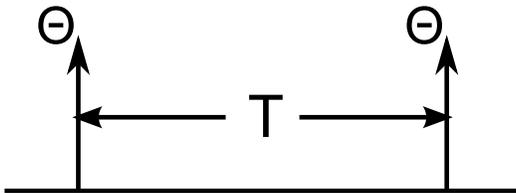
$T_R =$	$T_E =$	$T_I =$
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- c) Choose the scan parameters so that the liver produces no signal at all, but the spleen is bright.

$T_R =$	$T_E =$	$T_I =$
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4) Contrast Preparation:

Consider the following sequence of two  $\theta$  degrees tip-angle RF pulses separated by  $T$  seconds.



- a) Given that the equilibrium magnetization is  $M_0$ , derive an expression for the  $M_z$  component of the magnetization immediately following the second RF as a function of  $T_2$ ,  $T$ , and  $\theta$ . You can neglect  $T_1$  recovery (since  $T_1 \gg T$ ) and off-resonance.

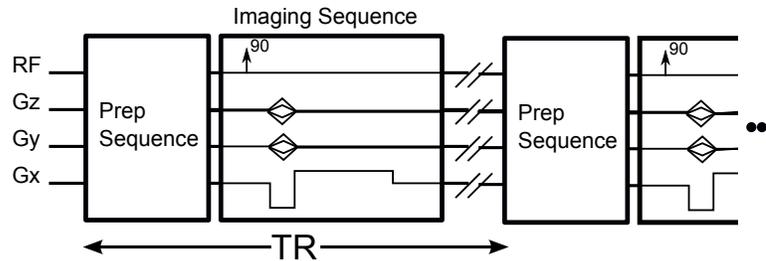
$M_z =$
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- b) Given  $T = T_0$ , find the flip angle  $\theta$  for which the  $M_z$  component is zero for spins with a desired  $T_2$  transverse relaxation value.

$\theta =$
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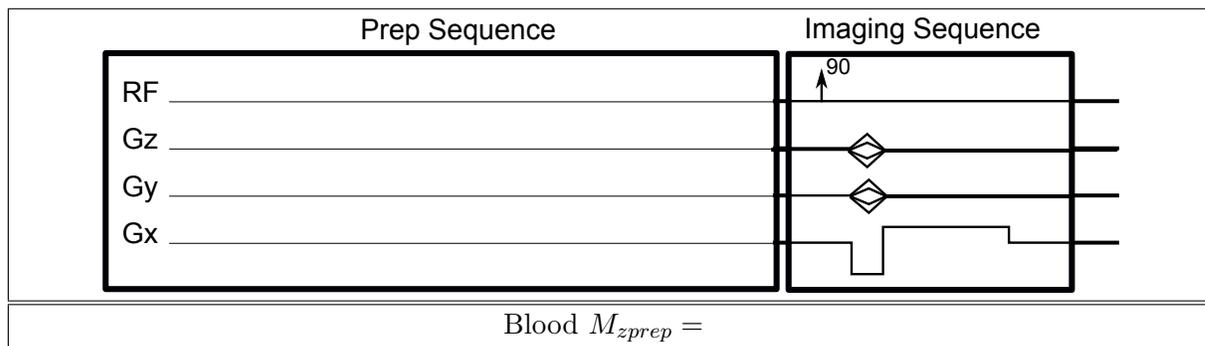
MR Angiography is an important tool in assessing vascular diseases in patients. Often,  $T_1$  shortening Gadolinium contrast agents are used in combination with short TR sequences to increase the blood-muscle contrast. However, using Gadolinium based contrast agents can result in a life threatening syndrome, called NSF, in patients that have renal disease.

Consider the contrast preparation imaging paradigm below. For each preparation sequence a single phase encode is collected. Also, assume  $TR \gg T_1$ .



The  $T_1/T_2$  of blood are 1000/220ms and the  $T_1/T_2$  of muscle are 870/50ms. We would like to design a non-contrast enhanced preparation pulse that will ideally have good blood signal and no muscle signal at all. In addition, we would like the preparation pulse to not be much longer than 50ms.

- c) Based on your previous derivations, design a preparation sequence that nulls the muscle signal while producing signal from blood. Draw the sequence pointing out the relevant parameters. (Extra points will be given for those coming with solutions that are insensitive to off-resonance). what is the blood  $M_z$  magnetization after the prep-pulse?



- 5) Nishimura 7.3
- 6) In order to produce a very high resolution image of the brain, you specify a scan that provides 0.25 mm resolution over a 25.6 cm FOV. Unfortunately, the SNR is very low.
- You lowpass filter the image to increase the SNR, using an ideal lowpass filter in both  $x$  and  $y$ . If the resulting resolution is 0.5 mm in both  $x$  and  $y$ , what is the final SNR compared to the original image?
  - How does this SNR compare to an image with 0.5 mm resolution, and the same total imaging time as the 0.25 mm resolution image? Assume the same A/D duration for each readout interval, and that the 0.5 mm scan averages to provide the same number of readouts.
- 7) You have a working 2DFT gradient-recalled echo pulse sequence that produces an image with an  $SNR_1$  of 100. What is the new  $SNR_2$  after you make *one* of the following changes. Add a brief explanation as well.

- Double the TBW of the slice selective RF pulse, while the RF pulse duration and the slice select gradient remain the same.

$SNR_2 =$
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- Double the readout gradient strength, while keeping the A/D duration and sampling rate the same.

$SNR_2 =$
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- Double both the readout gradient strength and the sampling rate, while halving the duration of the A/D window. Assume the anti-aliasing filter bandwidth matches the sampling rate.

$SNR_2 =$
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- Double the number of phase encodes, while keeping the maximum phase-encode gradient amplitude the same.

$SNR_2 =$
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- 8) Nishimura 7.10