

HW 2: Medical Imaging Systems

Jake Bergquist, u6010393

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Q1

a)Goal: Derive the following relationship

$$M_0 = \frac{N\gamma^2\hbar^2}{16\pi^2 kT} B_0$$

To begin with I will lay out some of the foundational assumptions and relationships based on class lecture.

$$\hbar = \frac{h}{2\pi} \quad (1)$$

$$N_+ + N_- = N \quad (2)$$

$$N_- \approx \frac{N}{2} \quad (3)$$

$$\mu = \gamma\hbar I \quad (4)$$

$$E = -\mu B_0 = -\gamma\hbar I B_0 \quad (5)$$

$$\frac{N_+}{N_-} = e^{\frac{-\Delta E}{kT}} \quad (6)$$

$$M_0 = \frac{(N_+ - N_-)\gamma\hbar}{2} \quad (7)$$

The first step I will take is to break ΔE into $E_+ - E_-$ by definition. By then substituting the definitions for E from Eq 5 into Eq 6 we get Eq 8 below (after distributing the negative and using $I = 1/2$ for E_+ and $I = -1/2$ for E_-).

$$\frac{N_+}{N_-} = e^{\frac{-(E_+ - E_-)}{kT}} = e^{\frac{\gamma\hbar\frac{1}{2}B_0 + \gamma\hbar\frac{1}{2}B_0}{kT}} \quad (8)$$

The term in the numerator of the exponential combines to give $\gamma\hbar B_0$. When e^x and x is small in magnitude $e^x \approx 1 + x$. Using this approximation we can reduce Eq 8 by converting the exponential as shown here and then multiplying N_- over. This yields Eq 9 below.

$$N_+ = N_- + \frac{N_- \gamma\hbar B_0}{kT} \quad (9)$$

By subtracting N_- over and applying the approximation of Eq 3 we get Eq 10

$$N_+ - N_- = \frac{N\gamma\hbar B_0}{2kT} \quad (10)$$

By substituting this quantity into Eq 7 and substituting out \hbar according to Eq 1 we get our final result.

b) at the given dimensions, each voxel would have dimensions of $24/192\text{cm} \times 24/192\text{cm} \times 1.0\text{mm} = 1.25\text{mm} \times 1.25\text{mm} \times 1.0\text{mm}$ giving us a volume of 1.6mm^3 per voxel. 85% of this volume is occupied by water in our tumor. Water has a density of $1\text{g/cm}^3 = 0.001\text{g/mm}^3$ giving us $1.6\text{mm}^3 * 85\% * 0.001\text{g/mm}^3 = 0.0014\text{g}$ of water. At an atomic mass of 14 g/mole, at a mass of 0.0014 g this gives us $\frac{0.0014\text{g}}{14\text{g/mole}} = 0.0001\text{mole}$ of water molecules. At 2 hydrogen per water molecule this gives us a total of 0.0002mole of hydrogen atoms or $1.2 * 10^{21}$ hydrogen nuclei in water molecules. Assuming a body temperature of 310.15 K (37 Celsius), knowing that $\gamma/2 * \pi = 42.58\text{MHz/T}$ for hydrogen nuclei, with a 3 T magnet:

$$M_0 = \frac{N\gamma^2\hbar^2}{16\pi^2 kT} B_0 = \frac{N\hbar^2}{4kT} \frac{\gamma^2}{4\pi^2} B_0 = \frac{1.2 * 10^{21} * (6.6 * 10^{-34}\text{J/s})^2}{4 * (1.4 * 10^{-23}\text{J/K}) * 310.15\text{K}} (42.58\text{MHz/T})^2 * 3T$$

Q2

a

To begin with, from lecture we know EQ 11

$$M_z(t) = M_{z\text{initial}} e^{-\frac{t}{T_1}} + m_0(1 - e^{-\frac{t}{T_1}}) \quad (11)$$

This can be reformulated in terms of the steady state M_z which will occur at TR time. Once at steady state the $M_{z\text{initial}} = M_z^{ss} \cos(\alpha)$ because at this steady state after each excitation it is the steady state magnitude that will determine what the initial M_z value will be. Thus we get Eq 12. By solving for M_z^{ss} we get Eq 13 which is our desired steady state longitudinal magnetization.

$$M_z^{ss} = M_z(TR) = M_z^{ss} e^{-\frac{TR}{T_1}} + m_0(1 - e^{-\frac{TR}{T_1}}) \quad (12)$$

$$M_z^{ss} = \frac{m_0(1 - e^{-\frac{TR}{T_1}})}{1 - \cos(\alpha)e^{-\frac{TR}{T_1}}} \quad (13)$$

b

The above describes the magnitude of the M_z^{ss} but the observed/measured magnitude must be in the xy plane in order to be measured by the receiving coil. Thus the observed steady state longitudinal magnetization is given by Eq 14.

$$M_{z\text{observed}}^{ss} = M_z^{ss} \sin(\alpha) = \frac{\sin(\alpha)m_0(1 - e^{-\frac{TR}{T_1}})}{1 - \cos(\alpha)e^{-\frac{TR}{T_1}}} \quad (14)$$

In order to find the flip angle (α) that gives the greatest observed signal we must differentiate Eq 14 with respect to α , set this equal to zero and solve for the alpha that maximizes intensity.

We must then check that this α is indeed a maxima. In order to make this process simpler, terms in Eq 14 that are constant and do not change with respect to α will be lumped into constant terms. such that $C_1 = m_0(1 - e^{-\frac{TR}{T_1}})$, and $C_2 = e^{-\frac{TR}{T_1}}$. Additionally the denominator will be brought up as a term with a negative exponent so that the product rule (rather than the quotient rule) can be used (by preference). The resulting simplification is shown in Eq 15.

$$M_{z_{observed}}^{ss} = C_1 \sin(\alpha) (1 - C_2 \cos(\alpha))^{-1} \quad (15)$$

Using the product rule and the chain rule this can be differentiated into the form seen in Eq 16 and simplified to Eq 17 by setting the derivative equal to zero and rearranging.

$$\frac{d}{d\alpha} = (1 - C_2 \cos(\alpha))^{-1} C_1 \cos(\alpha) + C_1 \sin(\alpha) (-(1 - C_2 \cos(\alpha))^{-2}) C_2 \sin(\alpha) \quad (16)$$

$$\frac{C_1 \cos(\alpha)}{1 - C_2 \cos \alpha} = \frac{C_1 C_2 \sin^2(\alpha)}{(1 - C_2 \cos \alpha)^2} \quad (17)$$

Eq 17 can be used to solve for the α that sets the derivative to 0 (either a maxima or minima) by canceling like terms, and applying a trig identity ($\sin^2 = 1 - \cos^2$). These steps are carried out in Eq 18 - 22.

$$\cos(\alpha) = \frac{C_2 \sin^2(\alpha)}{1 - C_2 \cos \alpha} \quad (18)$$

$$\cos(\alpha) - C_2 \cos^2(\alpha) = C_2 (1 - \cos^2(\alpha)) \quad (19)$$

$$\cos(\alpha) - C_2 \cos^2(\alpha) = C_2 - C_2 \cos^2(\alpha) \quad (20)$$

$$\cos(\alpha) = C_2 \quad (21)$$

$$\alpha = \arccos(C_2) = \arccos(e^{-\frac{TR}{T_1}}) \quad (22)$$

In order to confirm that this is the maxima I chose to plot the values of the observed intensities at a range of angles and observe where the maxima occurred as compared to what would be naïve guess of the angle that maximized signals. The code used to do so is shown below and the resulting curve is shown in Figure 1. As can be seen the maxima occurs as calculated at an angle of $\arccos(C_2)$. By comparison I might have expected the value to come at around $\frac{\pi}{2}$ as this would result in the entirety of Mz being excited into the xy plane, this is not the case. This is likely because if 90 degrees was chosen for the flip angle then it steady state longitudinal magnitude would be smaller, leading to a smaller signal when it is flipped into the xy plane.

```
%%
%Q2
C2 = exp(-5/5);
C1 = 5*(1-C2);
alps = [0:.01:3.14];
Mzs = C1*sin(alps)./(1-C2*cos(alps));
figure(1);clf();plot(alps,Mzs,'linewidth',2);hold on;
```

```

scatter(pi/2,C1*sin(pi/2)/(1-C2*cos(pi/2)),'ro');
scatter(acos(C2),C1*sin(acos(C2))/(1-C2*cos(acos(C2))), 'bo');
ylabel('Observed M_z');
xlabel('Alpha Values');
set(gca, 'fontsize',18)

```

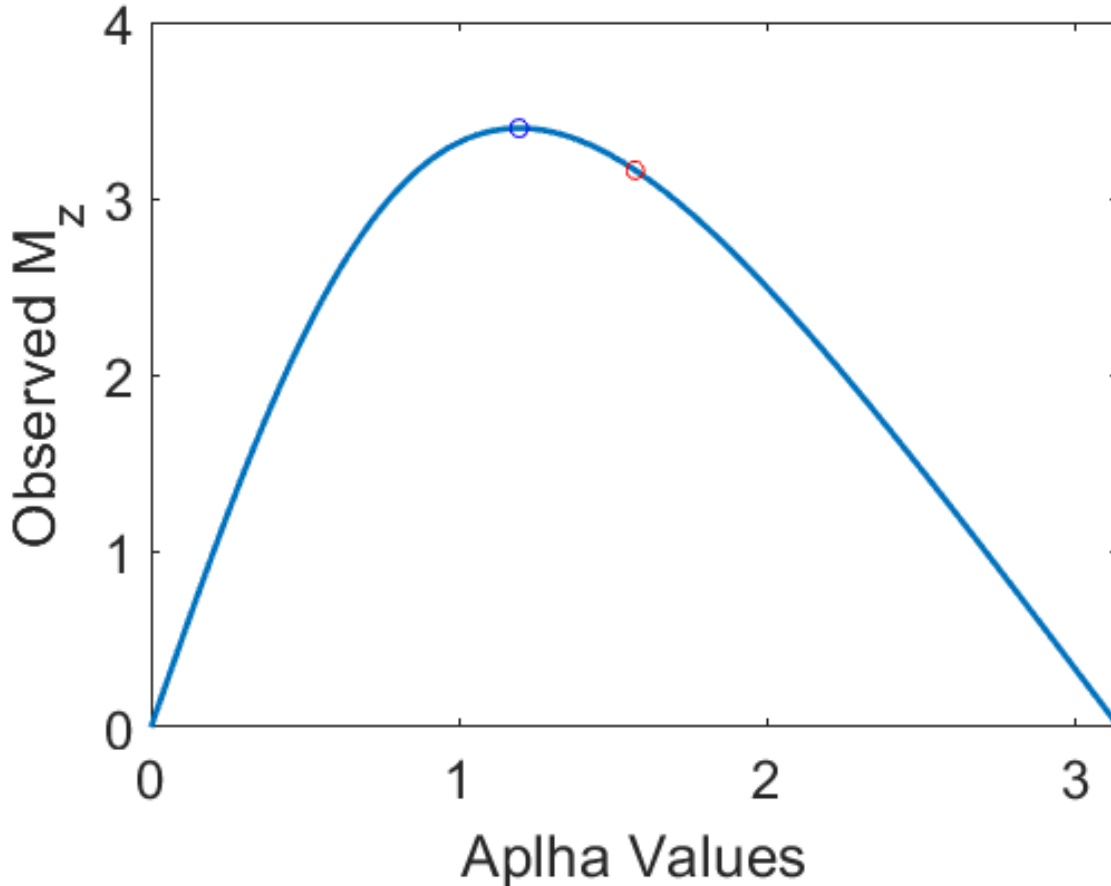


Figure 1: Alpha values vs the observed M_z for each at steady state. The red circle indicates the intensity at $\alpha = \pi/2$ and the blue circle indicates the intensity at $\alpha = C_2 = e^{-\frac{TR}{T_1}}$.

Q3

A simple method can be used to calculate the T_1 and T_2 using Eq 23. I_0 seems like a troubling quantity but it can be eliminated by taking multiple measurements and using their ratio to cancel out common terms. By making three measurements both T values can be measured. The first measurement uses essentially a random TE and TR. These should be selected to be within normal ranges, and their values should be reasonable but it is not important what specific values are chosen from a purely theoretical point of view. Technical limitations and practicalities of measurement will likely inform their selection. What is important is what is done next. These first values are called TE_1 and TR_1 respectively. A second set of parameters must be selected that are different

from TE_1 and TR_1 which we call TE_2 and TR_2 respectively. Again their selection will be based on practical concerns about the measurement system, the only important thing is that they be different from TE_1 and TR_1 . Three measurements of intensity will be taken using these values. One measurement using TE_1 and TR_1 , one using TE_1 and TR_2 , and a final one using TE_2 and TR_1 . Notice how in the second two measurements either TE or TR was changed from the first measurement but not both. These measurements produce intensity values of I_1 , I_2 , and I_3 respectively. By forming Ratios of either $\frac{I_1}{I_2}$ or $\frac{I_1}{I_3}$ we create scenarios where only TR or TE vary respectively. In the case of ratio 1 ($\frac{I_1}{I_2}$), all terms cancel out from Eq 23 except for those that involve TR. Thus the ratio 1 is dictated by T_1 and thus the ratio equation can be rearranged to solve for T_1 . The same principle is applied for Ratio 2 ($\frac{I_1}{I_3}$) to get a scenario that depends only on T_2 . In this way three measurements can be used to calculate both recovery times. This scenario of course assumes minimal/no noise. Real noise would necessitate multiple recordings in order to average the noise and minimize its impact on the relaxation time calculations.

$$I = I_0(1 - e^{-\frac{TR}{T_1}})e^{-\frac{TE}{T_2}} \quad (23)$$

Q4

a:

Eq 23 can be used to calculate the "T2-weighted" spin echo intensity. By fixing TR the second term (involving T1) becomes constant and can be combined with I_0 . This term I call $I^* = I_0(1 - e^{-\frac{TR}{T_1}})$. This gives us Eq 24.

$$I = I^*e^{-\frac{TE}{T_2}} \quad (24)$$

Taking the natural logarithm of both sides of this function and separating out I^* and T2 dependent parts using one of the logarithm rules we get Eq 25 which simplifies to Eq 26

$$\ln(I) = \ln(I^*) + \ln(e^{-\frac{TE}{T_2}}) \quad (25)$$

$$\ln(I) = -\frac{1}{T_2}TE + \ln(I^*) \quad (26)$$

Eq 26 shows a familiar $y = mx + b$ linear form where the slope is given by $-\frac{1}{T_2}$ and the y intercept is given by $\ln(I^*)$. The $\ln(I)$ term is the dependent variable while the TE is the independent variable. The $-\frac{1}{T_2}$ and I^* terms are unknown that could be solved for by fitting measured data to this equation form.

b

The code below was used to calculate the T2 and I^* for both the background and the tumor data. For the tumor T2 was found to be 84.1 ms and I^* was 1.4×10^3 . For the background/surrounding tissue the T2 was 79.0 and the I^* was 187.9. We can evaluate the fits by looking at the measured and fit curves in both the linearized and non-linear views. For the tumor data, Figure 2 shows the fit and measured curves. As can be seen the linearized fit agrees well with the data. Roughly 50% of the data appears to be above and 50% below the fit line and the distance between fit and measured is very small. When observing the non-linear view (Figure 2a) we see again that the

fit seems to match well, almost appearing as a spline interpolation between the measured points as opposed to the red linear interpolation. The fit results for the background/surrounding tissue are worse. Figure 3 shows these fitting results. In the linearized view, while still roughly 50% of the measured points lie on either side of the fit line the distance to the fit line is much higher. The linearized data itself is very spread out, implying a higher amount of noise in this area. This could be due to the lower amplitude of the signal and therefor lower SNR. The resulting fit on the non-linear view is very poor, and it seems that fitting this data without linearizing may have actually resulted in a better fit.

%Q4

```
TE = [10 20 30 50 100 250];
tumor = [1213.4 1099.3 994.7 783.2 447.6 70.9];
background = [448.1 306.7 239.1 119.0 1.1 30.4];

%Fit the tumor unknowns
coeffs_tumor = polyfit(TE,log(tumor),1);
T2_tumor = -1/coeffs_tumor(1);
IndependedntIntensity_Tumor = exp(coeffs_tumor(2));
%visualize both linearize dand original
figure(1);clf();hold on;plot(TE,tumor,'r','linewidth',2);
plot([10:1:250],IndependedntIntensity_Tumor*exp(-[10:1:250]/
    T2_tumor),'b','linewidth',2)
xlabel('TE (msec)');ylabel('intensity');set(gca,'fontsize',18);
figure(2);clf();hold on;plot(TE,log(tumor),'r','linewidth',2);
plot([10:1:250],coeffs_tumor(1).*[10:1:250]+coeffs_tumor(2),'b'
    ,'linewidth',2);
xlabel('TE (msec)');ylabel('intensity');set(gca,'fontsize',18);

%Fit for background
coeffs_background = polyfit(TE,log(background),1);

T2_background = -1/coeffs_background(1);
IndependedntIntensity_background = exp(coeffs_background(2));
%visualize both linearize dand original
figure(3);clf();hold on;plot(TE,background,'r','linewidth',2);
plot([10:1:250],IndependedntIntensity_background*exp
    (-[10:1:250]/T2_background),'b','linewidth',2)
xlabel('TE (msec)');ylabel('intensity');set(gca,'fontsize',18);
figure(4);clf();hold on;plot(TE,log(background),'r','linewidth'
    ,2);
plot([10:1:250],coeffs_background(1).*[10:1:250]+
    coeffs_background(2),'b','linewidth',2);
xlabel('TE (msec)');ylabel('intensity');set(gca,'fontsize',18);
```

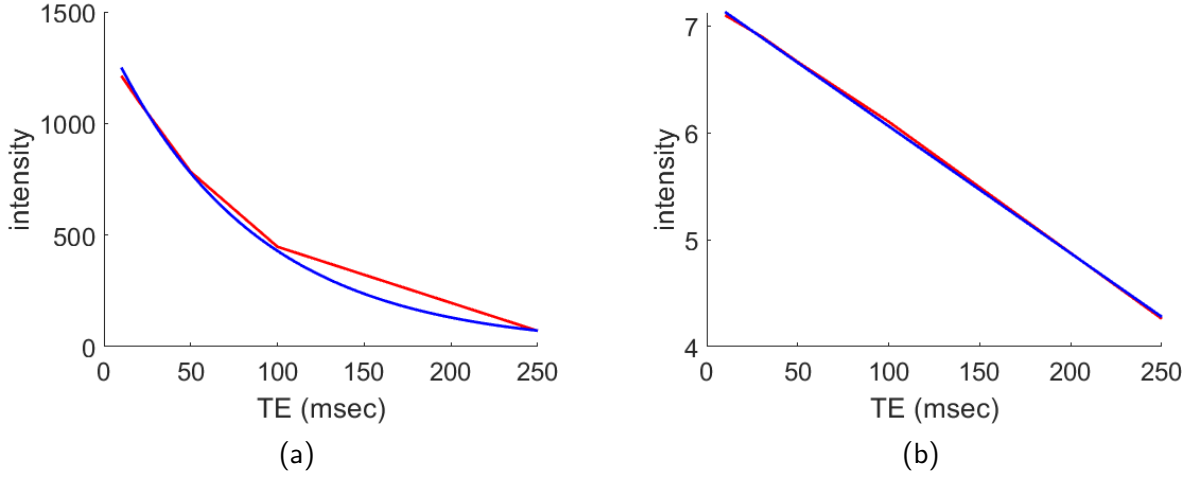


Figure 2: Results of fitting the tumor data (red) and plotting using the fit parameters (blue) and Eq 24 (a) or 26 (b). a shows the non-linearized measured data and fitline, while b shows the linearized data and fitline.

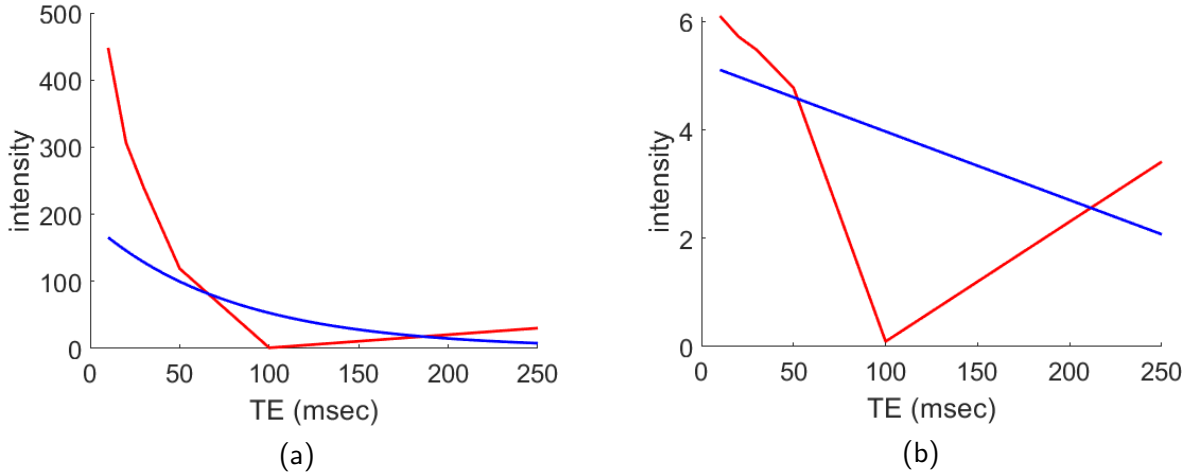


Figure 3: Results of fitting the background data (red) and plotting using the fit parameters (blue) and Eq 24 (a) or 26 (b). a shows the non-linearized measured data and fitline, while b shows the linearized data and fitline.

c

To maximize contrast we must maximize Eq 27 with respect to TE.

$$\frac{I_{tumor}}{I_{background}} = \frac{e^{-\frac{TE}{T_{2tumor}}}}{e^{-\frac{TE}{T_{2background}}}} = e^{-\frac{TE}{T_{2tumor}} + \frac{TE}{T_{2background}}} = e^{TE(-\frac{1}{T_{2tumor}} + \frac{1}{T_{2background}})} \quad (27)$$

By taking the derivative and setting it equal to zero we can get the maximizing value of TE by solving for TE. This is given by Eq 28

$$\frac{I_{tumor}}{I_{background}} = \frac{e^{-\frac{TE}{T_{2tumor}}}}{e^{-\frac{TE}{T_{2background}}}} = e^{-\frac{TE}{T_{2tumor}} + \frac{TE}{T_{2background}}} = e^{TE(-\frac{1}{T_{2tumor}} + \frac{1}{T_{2background}})} \quad (28)$$

Q5

a

Generically we have Eq 11. In our case of a 180 degree pulse $M_{initial} = m_0 \cos(\alpha) = -M_0$. Substituting this into Eq 11 and distributing M_0 in the second term gives the relationship and simplification shown in Eq 29 which is the longitudinal magnetization as a function of T_1 , TI , and M_0 where $t = TI$.

$$M_z(t) = -M_0 e^{-\frac{t}{T_1}} + m_0 - M_0 e^{-\frac{t}{T_1}} = M_0(1 - 2e^{-\frac{t}{T_1}}) \quad (29)$$

b

Before we can get to curve fitting we must first address a problem. Because the intensities measured are absolute value, and we have done a 180 degree pulse there is an artifact in the signal due to this absolute value measurement. This makes the curves look more like check marks. This can be seen in the table as the values first are decreasing then increasing as TI increases. To fix this the first three values in both rows of the table must be negated. Otherwise fitting will fail. The code below was used for this fitting using `fminsearch` and an objective function that produced the squared difference between the measurements and Eq 29, optimizing over M_0 and T_1 . 500 and 500 were selected as initial guesses in both cases because it was found that using a small number such as 1 or 0 resulted in a poor fit. M_0 was found to be 972 J/s and 359 J/s for tissue 1 and 2 respectively. the T_1 was found to be 609 msec 409 msec for tissue 1 and 2 respectively. The resulting fits (Figure 4) show good visual agreement.

%Q5

```
TI = [50 100 200 400 800 1600];
tissue1 = [-889 -684 -461 99.4 385 780];
tissue2 = [-261 -217 -108 118 254 339];

objFunc_tiss1 = @(param) LongMagIntensity(TI,tissue1,param);
optParams_tiss1 = fminsearch(objFunc_tiss1,[500,500]);
figure(1);clf();hold on;
plot(TI,tissue1,'r','linewidth',2);
plot([50:1600],optParams_tiss1(1).*(1-2*exp(-[50:1600]/
    optParams_tiss1(2))), 'b','linewidth',2);
xlabel('TI (msec)');ylabel('intensity');set(gca,'fontsize',18);

objFunc_tiss2 = @(param) LongMagIntensity(TI,tissue2,param);
optParams_tiss2 = fminsearch(objFunc_tiss2,[500,500]);
figure(2);clf();hold on;
plot(TI,tissue2,'r','linewidth',2);
```



```

plot([50:1600],optParams_tiss2(1).*(1-2*exp(-[50:1600]/
    optParams_tiss2(2))), 'b', 'linewidth',2);
xlabel('TI (msec)');ylabel('intensity');set(gca,'fontsize',18);

function cost = LongMagIntensity(TI,m,param)
M0 = param(1);
T1 = param(2);
cost = sum((m - M0 * (1 - 2*exp(-TI/T1))).^2);

end

```

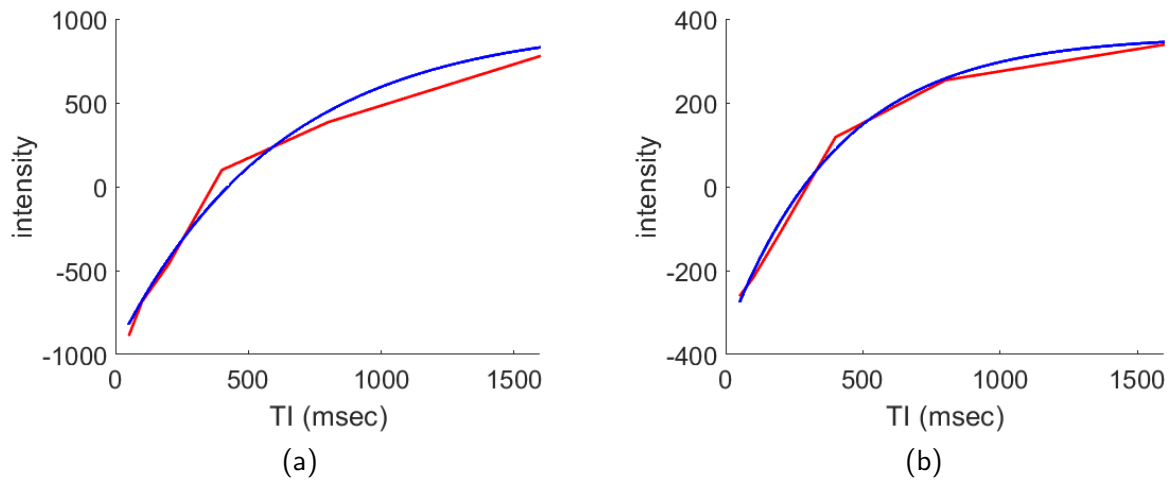


Figure 4: Results of fitting the data for each tissue (red) and plotting using the fit parameters (blue) and Eq 29. (a) shows the data and fit for tissue 1, while (b) shows the data and fit for tissue 2.