

# HW 2: Medical Imaging Systems

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## Q1

a)

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  $1.6\text{mm}^3$  per voxel. 85% of this volume is occupied by water in our tumor. Water has a density of  $1\text{g}/\text{cm}^3 = 0.001\text{g}/\text{mm}^3$  giving us  $1.6\text{mm}^3 * 85\% * 0.001\text{g}/\text{mm}^3 = 0.0014\text{g}$  of water. At an atomic mass of 14 g/mole, at a mass of 0.0014 g this gives us  $\frac{0.0014\text{g}}{14\text{g}/\text{mole}} = 0.0001\text{mole}$  of water molecules. At 2 hydrogen per water molecule this gives us a total of  $0.0002\text{mole}$  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  $\gamma/2 * \pi = 42.58\text{MHz}/\text{T}$  for hydrogen nuclei, with a 3 T magnet:

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

%%

%Problem 5

%kernel a

averaging\_kernel = ones(3,3)/9;

%kernel b

%orientation flipped due to matlab conventions

vertical\_edge\_detector = zeros(3,3);

vertical\_edge\_detector(:,3) = 1;

vertical\_edge\_detector(:,1) = -1;

%kernel c

%orientation flipped due to matlab conventions

horizontal\_edge\_detector = zeros(3,3);

horizontal\_edge\_detector(1,:) = 1;

horizontal\_edge\_detector(3,:) = -1;

conv1\_results = ...

conv2(MRI\_Image, averaging\_kernel, 'same');

%I use the same argument to not get increased image size

%It does however leave in zero padded edges in the calculation  
but the

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%edges are all background and I do not mind as much there
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conv2_results = ...  
conv2(MRI_Image,vertical_edge_detector,'same');  
  
conv3_results = ...  
conv2(MRI_Image,horizontal_edge_detector,'same');  
  
figure(1);  
subplot(2,2,1);  
imagesc(MRI_Image')  
axis('equal')  
axis('tight')  
colormap(gray);  
title('Original')  
subplot(222);  
imagesc(conv1_results')  
axis('equal')  
axis('tight')  
colormap(gray);  
title('3x3 Average')  
subplot(223);  
imagesc(conv2_results')  
axis('equal')  
axis('tight')  
colormap(gray);  
title('Vertical edge detector')  
subplot(224);  
imagesc(conv3_results')  
axis('equal')  
axis('tight')  
colormap(gray);  
title('Horizontal edge detector')
```