BMEN 509 W2019 - Laboratory 1 - Name

January 25, 2019

1 BMEN 509 - Laboratory 1 - Image Domain

The purpose of this laboratory is to become familiar with digital images and how to work with images. In this exercise, you will demonstrate 1) a basic understanding of digital image representation, 2) an ability to perform simple operations on images, and 3) an understanding of fundamental image processing techniques.

Your rubric is as follows:

Please use this notebook to complete your assignment. Sections to be completed are blank and are marked by TODO. Please upload the notebook (.ipynn) and as a PDF (.pdf) by using File > Download as > PDF via LaTeX.

TODO: Bryce A. Besler, November 19th, 2018

1.0.1 Objective

TODO

1.1 Notebook Setup

```
In [27]: # Library Imports
         import os
         import math
         import numpy as np
         import matplotlib.pyplot as plt
         from skimage import io
         from scipy import signal, ndimage
         # Setup Jupyter notebook specifics
         %matplotlib inline
         # Numpy printing help
         np.set_printoptions(precision=2)
In [3]: # Constants
        data_directory = os.path.join('.', 'Data')
        file_names = [
            'BreastUltrasound5.tif', 'T1Weighted8.jpg',
            'Thorax_PD.png', 'calcs_grids.tiff',
```

Found all image files!

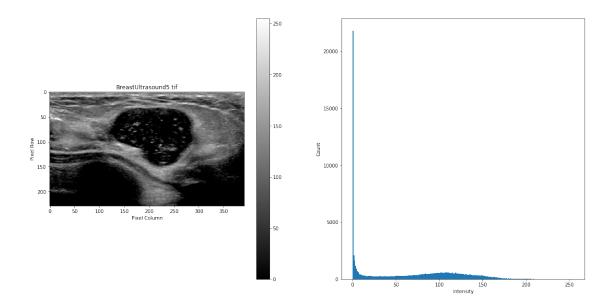
1.2 Question 1 - Image Basics

Using the image BreastUltrasound5.tif do the following: 1. Load the image 2. Display the image 3. Give the image a title, x-axis label, y-axis label, and add a colorbar 4. What is the size of the image in the X and Y dimensions? 5. How many pixels are in the image? 6. Which corner of the image is the pixel (0,0)? 7. Plot the histogram of the image with 256 bins

```
In [8]: # TODO: Implement
    filename = os.path.join(data_directory, 'BreastUltrasound5.tif')
    I = io.imread(filename)

# plot image with labels and color bar, and histogram with 256 bins
    plt.subplots(1, 2, figsize=(20,10))
    plt.subplot(1, 2, 1); plt.imshow(I, cmap='gray'); plt.title('BreastUltrasound5.tif'); plt.subplot(1, 2, 2); plt.hist(I.ravel(), bins=256); plt.xlabel('Intensity'); plt.ylabel.show()

    print('4. The Image has: {0} pixels in the X dimension and {1} pixels in the Y dimension print('5. The image has {} pixels.'.format(np.prod(I.shape)))
    print('6. The top-left corner of the image is the pixel (0, 0), as seen by the image pixels.'.
```



- 4. The Image has: 229 pixels in the X dimension and 392 pixels in the Y dimension
- 5. The image has 89768 pixels.
- 6. The top-left corner of the image is the pixel (0, 0), as seen by the image plot

1.3 Question 2 - Transfer Functions

Window and Level is a common technique for visualizing biomedical images. You can visualize specific tissues by expanding the contrast in a narrow range of gray values while ignore all other gray values. Window and Level is easy to implement as a simple transfer function, given below.

The linear transform is given by:

$$I(x,y) = m * I(x,y) + b$$

Determine how to calculate m and b. If you get stuck, refer to the laboratory notes. Write a function window_level_function that takes as input an image, a window, and a level and returns as output an image that has been windowed and levelled. The output image should be an 8-bit unsigned integer. The input image will need to be converted to a double and you will have to convert the output image to an unsigned 8-bit integer. Remember that the maximum value for an 8-bit unsigned integer is 255.

The code below loads the image mri_T2.tif. Using your function, it displays the image and its histogram at the following windows and levels. The corresponding tissue for each window and level is given on the right.

Tissue	Window	Level
Whole dynamic range	255	128
High contrast between white and gray matter	50	50
Background	10	20
Tumour	50	100

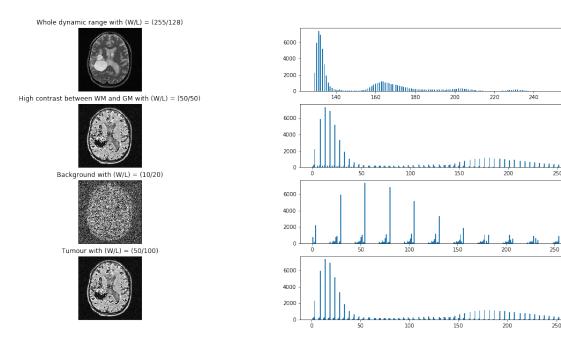
After writing your function and running the given code, answer the following questions:

```
1) Compare the histogram of the second window level transformation (W=50, L=50) to the histogram
2) In your own words, why does windowing and levelling allow us to see different tissues?
In [15]: def window_level_function(image, window, level):
             # TODO: Implement this function
             float image = image.astype(np.float32)
             min = 0
             max = 255
             # calculate m and b
             m = (max - min) / window
             b = max - (((max - min) / window) * (level + (window / 0.5)))
             # apply window/level function to image
             wl_image = m * float_image + b
             return wl_image.astype(np.uint8)
In [17]: image = io.imread(os.path.join(data_directory, 'mri_T2.tif'))
         pairs = {
             'Whole dynamic range':
                                                 (255, 128),
             'High contrast between WM and GM': (50, 50),
             'Background':
                                                 (10, 20),
             'Tumour':
                                                 (50, 100)
         }
         plt.subplots(len(pairs), 2, figsize=(20,10))
         for key in pairs:
             window, level = pairs[key]
             new_image = window_level_function(image, window, level)
             plt.subplot(len(pairs), 2, 2*i-1)
             plt.imshow(new_image, cmap='gray')
             plt.title('\{\} with (\{\}/\{\})'.format(key, window, level))
             plt.xticks([])
             plt.yticks([])
             plt.subplot(len(pairs), 2, 2*i)
             plt.hist(new_image.ravel(), bins=256)
             i += 1
         plt.show()
         print('''Question 1: Compare the histogram of the second window level transformation
         TODO: Answer
```

The pixels with intensities ranging from 128-140 are truncated in the second window level, and are drastically reduced in the second window level.

print('''Question 2: In your own words, why does windowing and levelling allow us to a
TODO: Answer

Window and levelling allows for pixels that fall within a specific range of intensity by a certain intensity range in images will be visualized better (or worse) with vary



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Window and levelling allows for pixels that fall within a specific range of intensity values to by a certain intensity range in images will be visualized better (or worse) with varying windown

1.4 Question 3 - Exploring a Gaussian Function

The Gaussian filter is one of the most powerful filters in image processing. Here, we are going to explore the effects of changing the kernel size and the response size of a Gaussian filter to better understand the parameters.

The code below loads the image knee_ct2.tif and displays the image. Next, it filters the image with a sigma of 2 and at five different kernel sizes: 3, 5, 7, 21 and 51. Next, it filters the image with a kernel size of 21 and at five different sigmas: 0.5, 1, 2, 3, 5. Plot the filtered image for all 11 responses.

Your objective is to write the function gaussian_kernel. This function creates a Gaussian kernel using the equation:

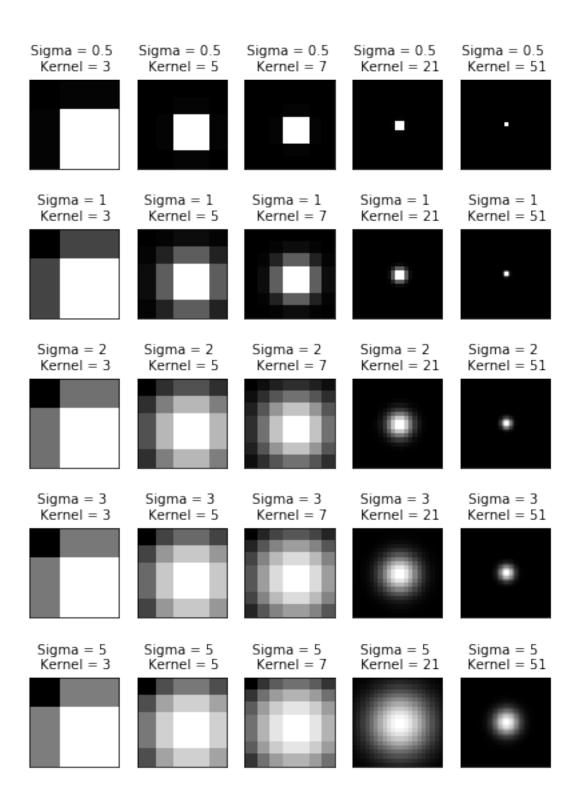
$$J(x,y) = \frac{1}{\sqrt{2\pi}\sigma} exp\left(-\frac{(x-\mu_x)^2 + (x-\mu_y)^2}{2\sigma^2}\right)$$

where σ determines the frequency response of the fitler and (μ_x, μ_y) are the centers of the kernel. Note that the kernel size is also a parameter!

If you need help conceptualizing image convolution, I would recommend Image Kernels Explained Visually.

In your own words, answer the following questions: 1. What is the effect of increasing the kernel size but keeping sigma constant? 2. What is the effect of increasing the sigma but keeping the kernel size constant? 3. Often a Gaussian filter is used to filter noise in an image. How should the parameters (sigma, kernel size) be chosen for filtering noise? 4. What is the disadvantage of making the kernel extremely large?

```
In [52]: def gaussian_kernel(sigma, kernel_size):
             # TODO: Implement this function
             kernel = np.zeros((kernel_size, kernel_size))
             centerx = kernel_size / 2
             centery = kernel_size / 2
             for i in range(kernel_size):
                 for j in range(kernel_size):
                     denom = 1 / ((math.sqrt(2 * math.pi)) * sigma)
                     expon = math.exp(-((i - centerx)**2 + (j - centery)**2) / (2 * sigma**2))
                     value = denom * expon
                     kernel[i, j] = value
             return kernel
In [53]: # Load image
         image = io.imread(os.path.join(data_directory, 'knee_ct2.tif'))
In [54]: # Create kernels
         sigmas = [0.5, 1, 2, 3, 5]
         kernels = [3, 5, 7, 21, 51]
         plt.subplots(len(sigmas), len(kernels), figsize=(7, 7))
         for i, sigma in enumerate(sigmas):
```



In [55]: sigma = []
 kernel = []

```
# Kernel sizes
sigma = 1
kernels = [3, 5, 7, 21, 51]
plt.subplots(2, (len(kernels)+1)//2, figsize=(30, 15))
plt.subplot(2, (len(kernels)+1)//2, 1)
plt.imshow(image, cmap='gray')
plt.title('Original')
plt.xticks([])
plt.yticks([])
i = 2
for kernel in kernels:
    gauss = gaussian_kernel(sigma, kernel)
    smooth_image = signal.convolve2d(image, gauss, boundary='pad', mode='same')
    plt.subplot(2, (len(kernels)+1)//2, i)
    plt.imshow(smooth_image, cmap='gray')
    plt.title('Sigma = {}, Kernel = {}'.format(sigma, kernel))
    plt.xticks([])
    plt.yticks([])
    i += 1
plt.show()
```

Sigmas

```
sigmas = [0.5, 1, 2, 3, 5]
kernel = 21
plt.subplots(2, (len(kernels)+1)//2, figsize=(30, 15))
plt.subplot(2, (len(kernels)+1)//2, 1)
plt.imshow(image, cmap='gray')
plt.title('Original')
plt.xticks([])
plt.yticks([])
i = 2
for sigma in sigmas:
    gauss = gaussian_kernel(sigma, kernel)
    smooth = signal.convolve2d(image, gauss, boundary='pad', mode='same')
    plt.subplot(2, (len(kernels)+1)//2, i)
    plt.imshow(smooth, cmap='gray')
    plt.title('Sigma = {}, Kernel = {}'.format(sigma, kernel))
    plt.xticks([])
    plt.yticks([])
    i += 1
plt.show()
```

In [2]: print('''Question 1. What is the effect of increasing the kernel size but keeping sigma
TODO: Answer

There is little qualitative difference, so it is difficult to tell by visual inspection

111)

print('''Question 2. What is the effect of increasing the sigma but keeping the kernel
TODO: Answer

AS the sigma value is increased with a constant kernel size, there is an increasing black.

print('''Question 3. Often a Gaussian filter is used to filter noise in an image. How a
TODO: Answer

The kernel size should be chosen according to the value of sigma, since sigma defines in a Gaussian distribution, the kernel size defines the width of the distribution. As a very

print('''Question 4. What is the disadvantage of making the kernel extremely large?
TODO: Answer

This can quickly result in high computational requirements. This is because the number

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1.5 Question 4 - Understanding Signal, Noise, and Contrast

Qualitative description and quantitative metrics are complementary tools for assessing image processing algorithms. Metrics are free from observer variability, but can fail to capture nuance. In

this section, we are going to repeat the Gaussian and use the median filtering (see Lab01Ex05) but include metrics for signal-to-noise ratio (SNR) and contrast-to-noise ratio (CNR). Using the image mri_pd.tif, the code below does the following: 1. Loads the image and add salt and pepper noise (with noise density 0.1) and Gaussian noise (with mean 0, variance 0.01) 2. Filters the resulting noisy images with a median filter and a Gaussian filter with sigma 3 3. Selects two points on the image to compare contrast. 4. Computes the SNR for the image, the CNR for the two points, and plots the points and the images together. 5. Produces an image similar to the following, where the red and green points are the indices passed to the function.

- 6. Plots the original image, the image with salt and pepper noise, and the image with Gaussian noise
- 7. Plots the salt and pepper noise image after median filtering and after Gaussian filtering
- 8. Plots the Gaussian noise image after median filtering and after Gaussian filtering

Your task is to write the functions SNR and CNR functions. Additionally, answer the following questions: 1. Qualitatively compare the original image to the Gaussian and salt and pepper noise images. What is the difference between Gaussian noise and salt and pepper noise? 2. Compare the median and Gaussian filtered salt and pepper noise images. Which filter produced the best CNR? Which filter produced the best SNR? From the perspective of a radiologist, which filter produced a qualitatively better image? 3. Compare the median and Gaussian filtered Gaussian noise images. Which filter produced the best CNR? Which filter produced the best SNR? From the perspective of a radiologist, which filter produced a qualitatively better image? 4. Does the best qualitative result always agree with the best numerical result? Why is it wise to visualize your images and not just rely on metrics?

```
In [57]: def SNR(image):
             # TODO: Implement
             N = np.prod(image.shape)
             mean = np.mean(image)
             unbiased_std = math.sqrt(N / (N - 1)) * np.std(image)
             return mean / unbiased_std
         def CNR(image, first_point, second_point):
             # TODO: Implement
             I1 = image[first_point[0], first_point[1]]
             I2 = image[second_point[0], second_point[1]]
             contrast = np.abs(I1 - I2)
             N = np.prod(image.shape)
             unbiased_std = math.sqrt(N / (N - 1)) * np.std(image)
             return contrast / unbiased_std
In [58]: def add_gaussian_noise(image, mean, var):
             # Express the variance as a percentage of the data type
             std = np.sqrt(var)*(np.iinfo(image.dtype).max - np.iinfo(image.dtype).min)
             noise = std * np.random.randn(*image.shape) + mean
             return image + noise
```

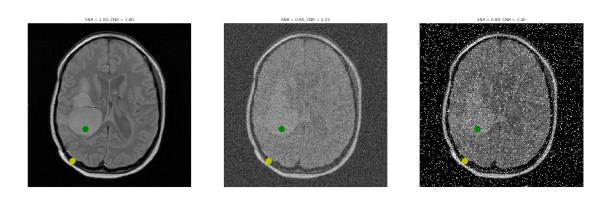
```
def add_salt_and_pepper_noise(image, density):
             # Draw random var. With probability p, ceil at bottom. With probability p, floor
             probability = density / 2.0
             rand_image = np.random.rand(*image.shape)
             output = image.copy()
             output[ rand_image < probability ] = np.iinfo(image.dtype).min</pre>
             output[ np.logical_and(rand_image >= probability, rand_image < density) ] = np.ii:
             return output
In [59]: def plot_image_with_point(image, first_point, second_point, show=True):
             plt.imshow(image, cmap='gray')
             plt.scatter(first_point[0], first_point[1], c='g', s=20*4**2)
             plt.scatter(second_point[0], second_point[1], c='y', s=20*4**2)
             plt.title('SNR = {0:0.2f}, CNR = {1:0.2f}'.format(
                 SNR(image),
                 CNR(image, first_point, second_point)
             ))
             plt.xticks([])
             plt.yticks([])
             if show:
                 plt.show()
In [60]: brain_point = (90, 165)
         csf_point = (70, 215)
         image = io.imread(os.path.join(data_directory, 'mri_pd.tif'))
         gauss_noise = add_gaussian_noise(image, 0, 0.01)
         med_noise = add_salt_and_pepper_noise(image, 0.1)
         print('Image and noise')
         plt.subplots(1, 3, figsize=(30, 10))
         for i, this_image in enumerate([image, gauss_noise, med_noise]):
             plt.subplot(1, 3, i+1)
             plot_image_with_point(this_image, brain_point, csf_point, False)
         plt.show()
         print('Salt & Pepper Noise')
         plt.subplots(1, 3, figsize=(30, 10))
         plt.subplot(1, 3, 1)
         plot_image_with_point(med_noise, brain_point, csf_point, False)
         plt.subplot(1, 3, 2)
         plot_image_with_point(ndimage.filters.gaussian_filter(med_noise, 1), brain_point, csf
```

```
plt.subplot(1, 3, 3)
plot_image_with_point(ndimage.filters.median_filter(med_noise, 3), brain_point, csf_point.show()

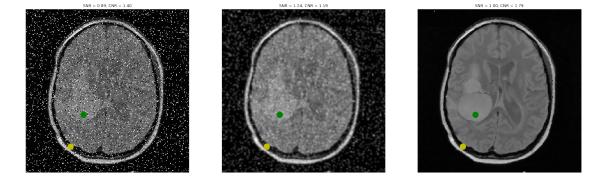
print('Gaussian Noise')
plt.subplots(1, 3, figsize=(30, 10))

plt.subplot(1, 3, 1)
plot_image_with_point(gauss_noise, brain_point, csf_point, False)
plt.subplot(1, 3, 2)
plot_image_with_point(ndimage.filters.gaussian_filter(gauss_noise, 1), brain_point, csplt.subplot(1, 3, 3)
plot_image_with_point(ndimage.filters.median_filter(gauss_noise, 3), brain_point, csf_plt.show()
```

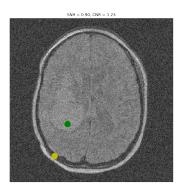
Image and noise

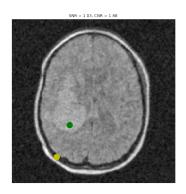


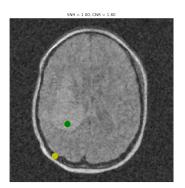
Salt & Pepper Noise



Gaussian Noise







The Gaussian noise is distributed more evenly throughout the image, whilst the salt and the image with the Gaussian noise has a lower contrast to noise ratio.

print('''Question 2. Compare the median and Gaussian filtered salt and pepper noise im-TODO: Answer

On the salt and pepper noise images, the Gaussian filter produced a better CNR, although filter image is of much greater quality as its features can be distinguished to a consit almost impossible to inspect. Thus, the Gaussian filter produced a better image from the constant of the con

print('''Question 3. Compare the median and Gaussian filtered Gaussian noise images. W.
TODO: Answer

Quantitatively, the median filter produced a better image than the Gaussian filter for both images have low contrasts and it is difficult to distinguish between features. The better in the perspective of a radiologist.

print('''Question 4. Does the best qualitative result always agree with the best numer
TODO: Answer

No, an image can be superior quantitatively, but may not be better qualitatively. This evenly throughout the image. In the example of the salt and pepper images, the median (1.00), yet is a much poorer image than the Gaussian filter image for diagnosis due to ''')

Question 1. Qualitatively compare the original image to the Gaussian and salt and pepper noise TODO: Answer

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Question 4. Does the best qualitative result always agree with the best numerical result? Why TODO: Answer

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1.5.1 Conclusion

TODO: Write your conclusion here

In [2]: print('''Quantitative metrics alone cannot be used to determine the diagnostic appropr
 For example, an image may have a better SNR than another image, but may be too poor in
 interpreted for diagnosis.''')

Quantitative metrics alone cannot be used to determine the diagnostic appropriateness of medic. For example, an image may have a better SNR than another image, but may be too poor in contrastinterpreted for diagnosis.

In []: