# C1\_W1\_Lab\_1\_data\_exploration\_and\_image\_preprocessing

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## 1 Data Exploration & Image Pre-Processing

In the first assignment of this course, you will work with chest x-ray images taken from the public ChestX-ray8 dataset. In this notebook, you'll get a chance to explore this dataset and familiarize yourself with some of the techniques you'll use in the first graded assignment.

The first step before jumping into writing code for any machine learning project is to explore your data. A standard Python package for analyzing and manipulating data is pandas.

With the next two code cells, you'll import pandas and a package called numpy for numerical manipulation, then use pandas to read a csv file into a dataframe and print out the first few rows of data.

```
In [1]: # Import necessary packages
    import pandas as pd
    import numpy as np
    import matplotlib.pyplot as plt
    %matplotlib inline
    import os
    import seaborn as sns
    sns.set()
```

## 1.1 1. Exploration

Read the data from csv files.

There are 1000 rows and 16 columns in this data frame

3	00012359_002.png		0	0		0	0		
4	00017951_001.png		0		0		0	0	
	Effusion	Emphysema	Fibrosis	Hernia	Infilt	cation	Mass	Nodule	\
0	0	0	0	0		0	0	0	
1	1	0	0	0		1	0	0	
2	0	0	0	0		0	0	0	
3	0	0	0	0		0	0	0	
4	0	0	0	0		1	0	0	
	PatientId	Pleural_Thickening		Pneumon	ia Pneı	mothor	ax		
0	8270		0		0		0		
1	29855		0		0		0		
2	1297		1		0		0		
3	12359		0		0		0		
4	17951		0		0		0		

Have a look at the various columns in this csv file. The file contains the names of chest x-ray images ("Image" column) and the columns filled with ones and zeros identify which diagnoses were given based on each x-ray image.

### 1.1.1 1.1 Data Types and Null Values Check

Run the next cell to explore the data types present in each column and whether any null values exist in the data.

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1000 entries, 0 to 999
Data columns (total 16 columns):
Image
                      1000 non-null object
                      1000 non-null int64
Atelectasis
Cardiomegaly
                      1000 non-null int64
Consolidation
                      1000 non-null int64
Edema
                      1000 non-null int64
                      1000 non-null int64
Effusion
                      1000 non-null int64
Emphysema
                      1000 non-null int64
Fibrosis
                      1000 non-null int64
Hernia
                      1000 non-null int64
Infiltration
                      1000 non-null int64
Mass
Nodule
                      1000 non-null int64
PatientId
                      1000 non-null int64
                      1000 non-null int64
Pleural_Thickening
Pneumonia
                      1000 non-null int64
Pneumothorax
                      1000 non-null int64
dtypes: int64(15), object(1)
```

memory usage: 125.1+ KB

### 1.1.2 1.2 Unique IDs Check

"PatientId" has an identification number for each patient. One thing you'd like to know about a medical dataset like this is if you're looking at repeated data for certain patients or whether each image represents a different person.

```
In [4]: print(f"The total patient ids are {train_df['PatientId'].count()}, from those the uniq
```

As you can see, the number of unique patients in the dataset is less than the total number so there must be some overlap. For patients with multiple records, you'll want to make sure they do not show up in both training and test sets in order to avoid data leakage (covered later in this week's lectures).

#### 1.1.3 1.3 Data Labels

Run the next two code cells to create a list of the names of each patient condition or disease.

The total patient ids are 1000, from those the unique ids are 928

There are 14 columns of labels for these conditions: ['Atelectasis', 'Cardiomegaly', 'Consolidations', 'Cardiomegaly', 'Consolidations', 'Cardiomegaly', 'Consolidations', 'Cardiomegaly', 'Consolidations', 'Cardiomegaly', 'Consolidations', 'Cardiomegaly', 'Cardiomegaly',

Run the next cell to print out the number of positive labels (1's) for each condition

```
The class Emphysema has 13 samples
The class Fibrosis has 14 samples
The class Hernia has 2 samples
The class Infiltration has 175 samples
The class Mass has 45 samples
The class Nodule has 54 samples
The class Pleural_Thickening has 21 samples
The class Pneumonia has 10 samples
The class Pneumonia has 38 samples
```

Have a look at the counts for the labels in each class above. Does this look like a balanced dataset?

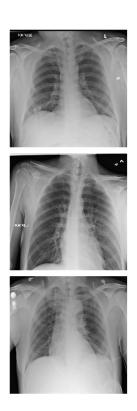
#### 1.1.4 1.4 Data Visualization

Display Random Images

Using the image names listed in the csv file, you can retrieve the image associated with each row of data in your dataframe.

Run the cell below to visualize a random selection of images from the dataset.

```
In [8]: # Extract numpy values from Image column in data frame
        images = train_df['Image'].values
        # Extract 9 random images from it
        random_images = [np.random.choice(images) for i in range(9)]
        # Location of the image dir
        img_dir = 'data/nih/images-small/'
        print('Display Random Images')
        # Adjust the size of your images
        plt.figure(figsize=(20,10))
        # Iterate and plot random images
        for i in range(9):
            plt.subplot(3, 3, i + 1)
            img = plt.imread(os.path.join(img_dir, random_images[i]))
            plt.imshow(img, cmap='gray')
            plt.axis('off')
        # Adjust subplot parameters to give specified padding
        plt.tight_layout()
```



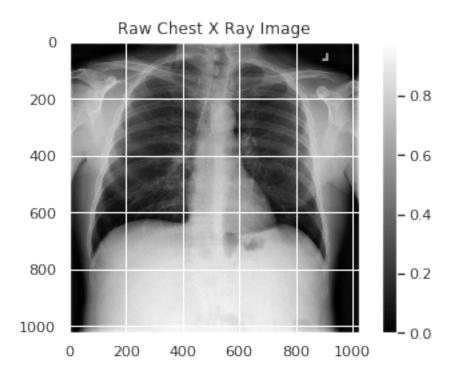




## 1.1.5 1.5 Investigating a Single Image

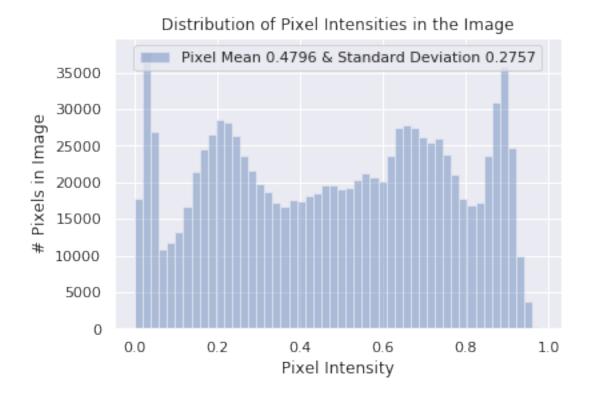
Run the cell below to look at the first image in the dataset and print out some details of the image contents.

The dimensions of the image are 1024 pixels width and 1024 pixels height, one single color character maximum pixel value is 0.9804 and the minimum is 0.0000 The mean value of the pixels is 0.4796 and the standard deviation is 0.2757



## 1.1.6 1.6 Investigating Pixel Value Distribution

Run the cell below to plot up the distribution of pixel values in the image shown above.



## 1.2 2. Image Preprocessing in Keras

Before training, you'll first modify your images to be better suited for training a convolutional neural network. For this task you'll use the Keras ImageDataGenerator function to perform data preprocessing and data augmentation.

Run the next two cells to import this function and create an image generator for preprocessing.

Using TensorFlow backend.

```
In [12]: # Normalize images
    image_generator = ImageDataGenerator(
        samplewise_center=True, #Set each sample mean to 0.
        samplewise_std_normalization= True # Divide each input by its standard deviation
)
```

#### 1.2.1 2.1 Standardization

The image\_generator you created above will act to adjust your image data such that the new mean of the data will be zero, and the standard deviation of the data will be 1.

In other words, the generator will replace each pixel value in the image with a new value calculated by subtracting the mean and dividing by the standard deviation.

$$\frac{x_i - \mu}{\sigma}$$

Run the next cell to pre-process your data using the image\_generator. In this step you will also be reducing the image size down to 320x320 pixels.

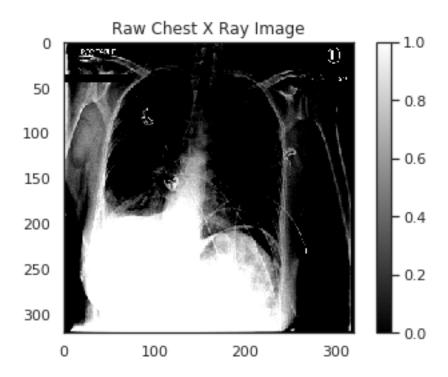
Found 1000 validated image filenames.

In above function class\_mode is set to raw because the data of Mass is continous and it is a regression problem. If it is a classification problem then it would be set to 'binary'

Run the next cell to plot up an example of a pre-processed image

Clipping input data to the valid range for imshow with RGB data ([0..1] for floats or [0..255]

```
The dimensions of the image are 320 pixels width and 320 pixels height. The maximum pixel value is 2.1455 and the minimum is -1.8822. The mean value of the pixels is -0.0000 and the standard deviation is 1.0000.
```

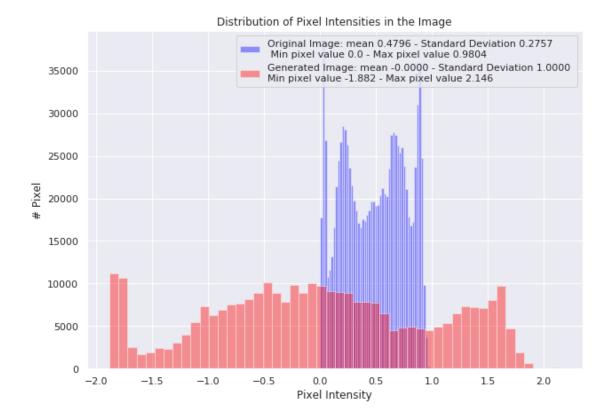


Run the cell below to see a comparison of the distribution of pixel values in the new preprocessed image versus the raw image.

```
In [17]: # Include a histogram of the distribution of the pixels
         sns.set()
         plt.figure(figsize=(10, 7))
         # Plot histogram for original iamge
         sns.distplot(raw_image.ravel(),
                       label=f'Original Image: mean {np.mean(raw_image):.4f} - Standard Deviation
                       f'Min pixel value {np.min(raw_image):.4} - Max pixel value {np.max(raw_image):.4}
                       color='blue',
                       kde=False)
         # Plot histogram for generated image
         sns.distplot(generated_image[0].ravel(),
                       label=f'Generated Image: mean {np.mean(generated_image[0]):.4f} - Standar
                       f'Min pixel value {np.min(generated_image[0]):.4} - Max pixel value {np.min(generated_image[0]):.4}
                       color='red',
                       kde=False)
         # Place legends
         plt.legend()
         plt.title('Distribution of Pixel Intensities in the Image')
         plt.xlabel('Pixel Intensity')
```

plt.ylabel('# Pixel')

Out[17]: Text(0, 0.5, '# Pixel')



That's it for this exercise, you should now be a bit more familiar with the dataset you'll be using in this week's assignment!