

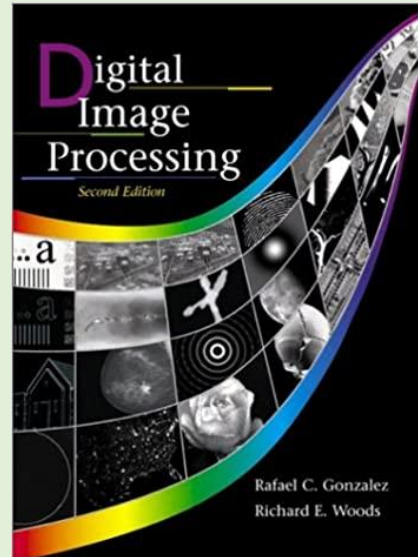
# Digital Image Processing

**From**

Image processing

**To**

Deep-learning using CNN



**IMAGE → PROCESSING → ANALYSIS → RECOGNIZNG**



**AASTMT**  
**Smart Village**

# Acknowledgement

I would like to thank our advisor Assoc. Prof. Dr Aliaa Youssif who really directed me to produce this project. I would like to give special thanks to our Professor for:

- ❖ Providing me with the needed technical information throughout the project.
- ❖ Not only the technical information but also for the soft skills; by giving me the chance to present every step of project in a formal way.
- ❖ Giving me the chance to search for the needed information in design.

Nonetheless, I would also thank Engineer Engy Ehab who gives me much knowledge in

- ❖ CNN (Convolutional neural network).
- ❖ Deep-learning as all
- ❖ Python
- ❖ Techniques

# **Our Project Split to 3 Phases 2 of them Image Processing and Last one using CNN Deep-Learning.**

No worry everything will be present clearly in Table of Contents

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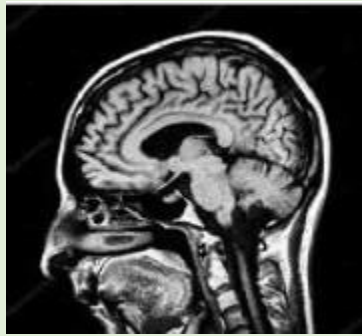
# Chapter One

## Analysis of MRI Images Using Image Processing Technique

### Example ONE

So, here's our image and we going to use some process on it to get it out

Original Image



Target Image



**Let's go ahead and see what need to happen**

### Process & Pipeline:

#### Step 1 Contrast:

```
imgcon=cv2.addWeighted(imgabs,alpha,np.zeros(imgabs.shape, imgabs.dtype),0,beta)
```

Note (Alpha control in Degree of Contrast)



#### Step 2 Blurring:

```
imggau2 = cv2.GaussianBlur(imgcon,(3,3),sigmaX=1,borderType=cv2.BORDER_CONSTANT)
```



### Step 3 Edge Detection using Canny:

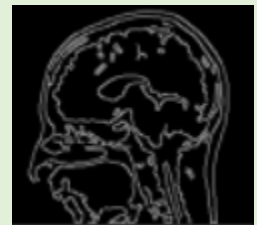
```
imgcanny= cv2.Canny(imggau2,300,800)
```



### Step 4 Repeat Step 3 with different values:

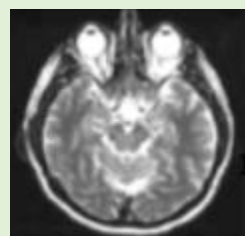
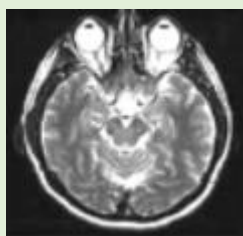
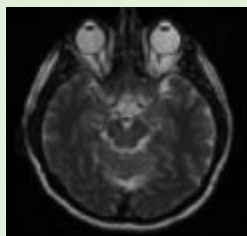
After applying many of canny but with different values we got our Target Finally

As you can See →



### Example Two

We know What happens from Example one so we will move Quickly on this one 😊



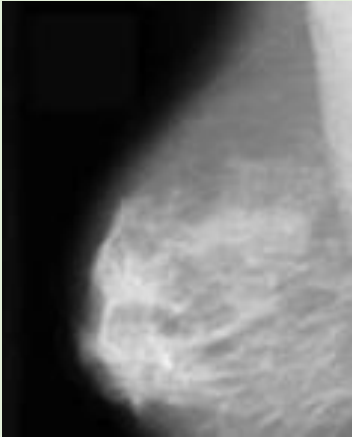
### Final Output



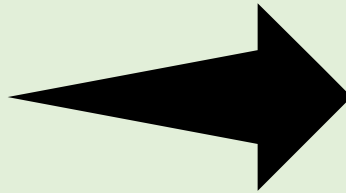
## Chapter Two

### Analysis of Breast Cancer Images Using Image Processing Technique

Original Image



Target Image

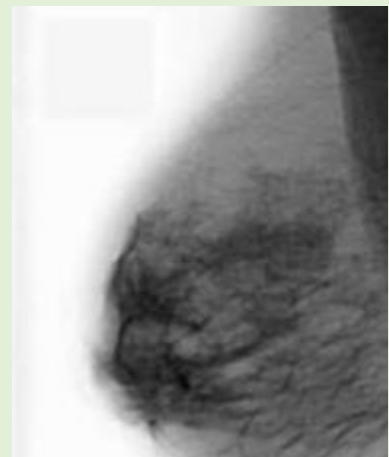


At Target Image you can see the cancer clearly after Applying some Image Processing Techniques

### Process & Pipeline:

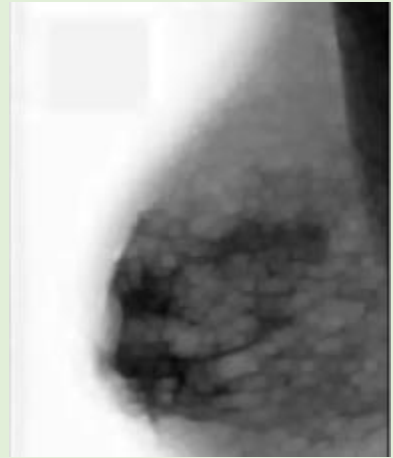
Step 1 Negative:

```
imgneg = 255-img
```



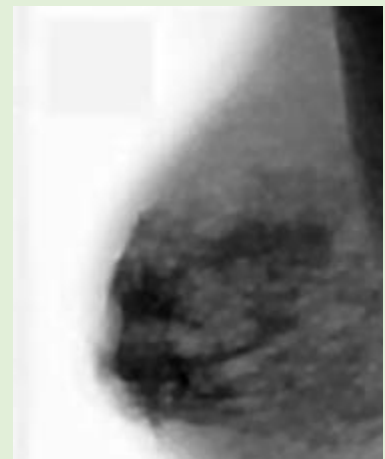
**Step 2 Dilation:**

```
imgdil = cv2.dilate(imgneg, kernel, iterations=1)
```



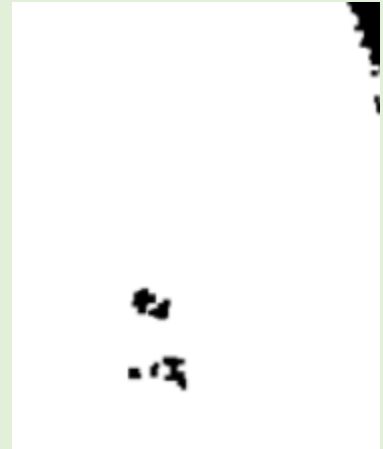
**Step 3 erosion:**

```
imgero = cv2.erode(imgdil, kernel, iterations=1)
```



#### Step 4 Threshold:

```
imgthreshold = cv2.threshold(imgero,50,90,cv2.THRESH_BINARY)
```



**Here at Threshold, you can see Cancer cells clearly in the breast**



## Chapter Three & Last

White Blood cells Classification using CNN Deep learning & TensorFlow & Keras & Real-life Dataset

**First, Let's Define our Dataset**

**It's 9956 Real images for white Blood Cells  
(EOSINOPHIL, LYMPHOCYTE, MONOCYTE, NEUTROPHIL)**

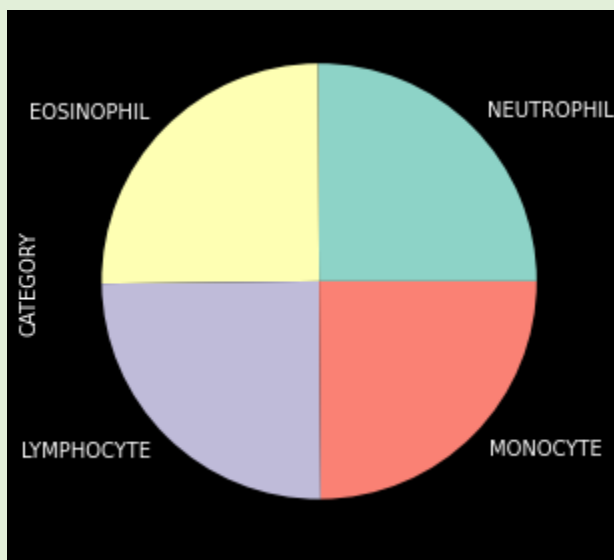
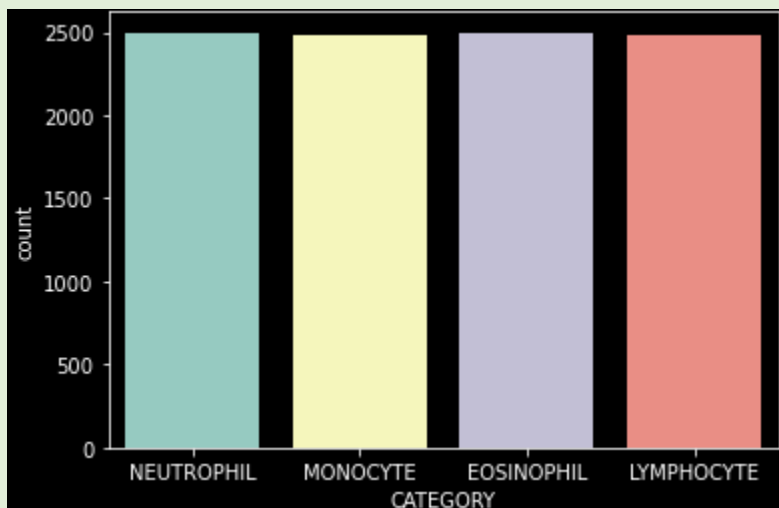
**We now will pick some of it to train and other to test and also a few as validation**

# Train

```
Train_Data_Path = Path("C:/Users/moham/Downloads/dataset2-master/dataset2-master/images/TRAIN")
```

```
print("EOSINOPHIL: ", Train_JPG_Labels.count("EOSINOPHIL"))  
print("LYMPHOCYTE: ", Train_JPG_Labels.count("LYMPHOCYTE"))  
print("MONOCYTE: ", Train_JPG_Labels.count("MONOCYTE"))  
print("NEUTROPHIL: ", Train_JPG_Labels.count("NEUTROPHIL"))
```

```
EOSINOPHIL: 2497  
LYMPHOCYTE: 2483  
MONOCYTE: 2478  
NEUTROPHIL: 2499
```

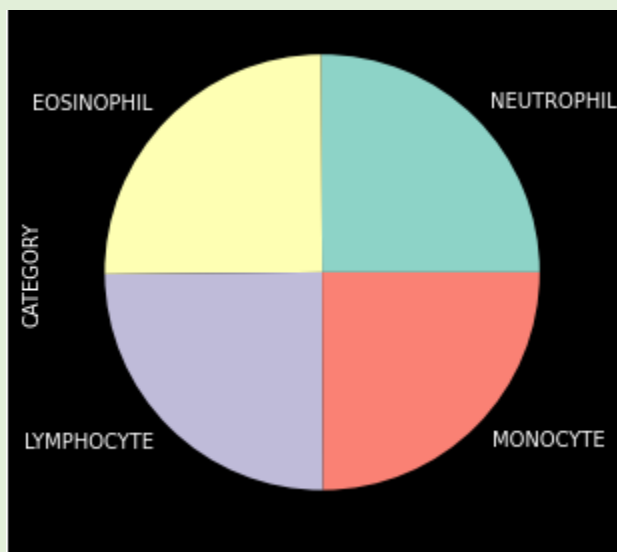
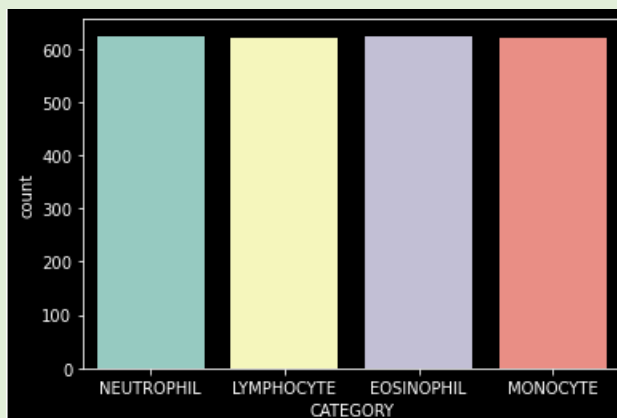


## Test

```
Test_Data_Path = Path("C:/Users/moham/Downloads/dataset2-master/dataset2-master/images/TEST")
```

```
print("EOSINOPHIL: ", Test_JPG_Labels.count("EOSINOPHIL"))  
print("LYMPHOCYTE: ", Test_JPG_Labels.count("LYMPHOCYTE"))  
print("MONOCYTE: ", Test_JPG_Labels.count("MONOCYTE"))  
print("NEUTROPHIL: ", Test_JPG_Labels.count("NEUTROPHIL"))
```

```
EOSINOPHIL: 623  
LYMPHOCYTE: 620  
MONOCYTE: 620  
NEUTROPHIL: 624
```

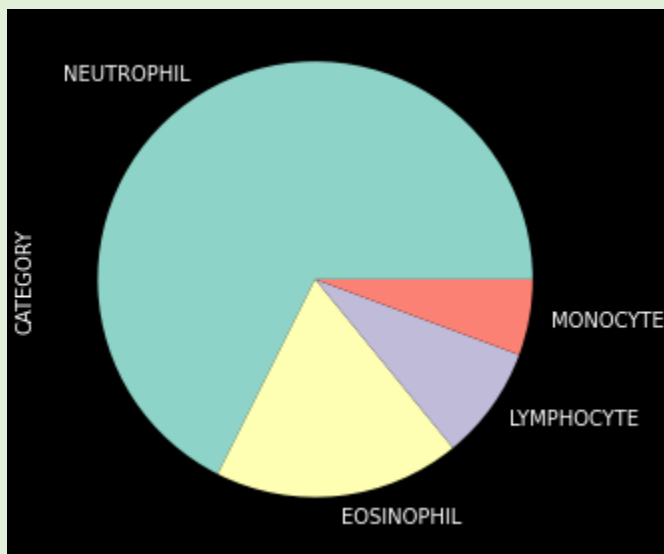
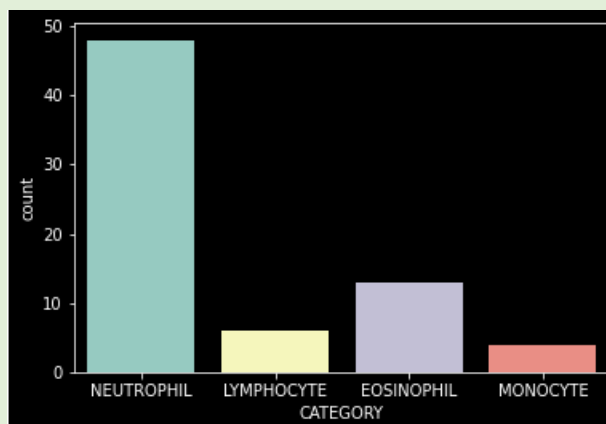


# Validation

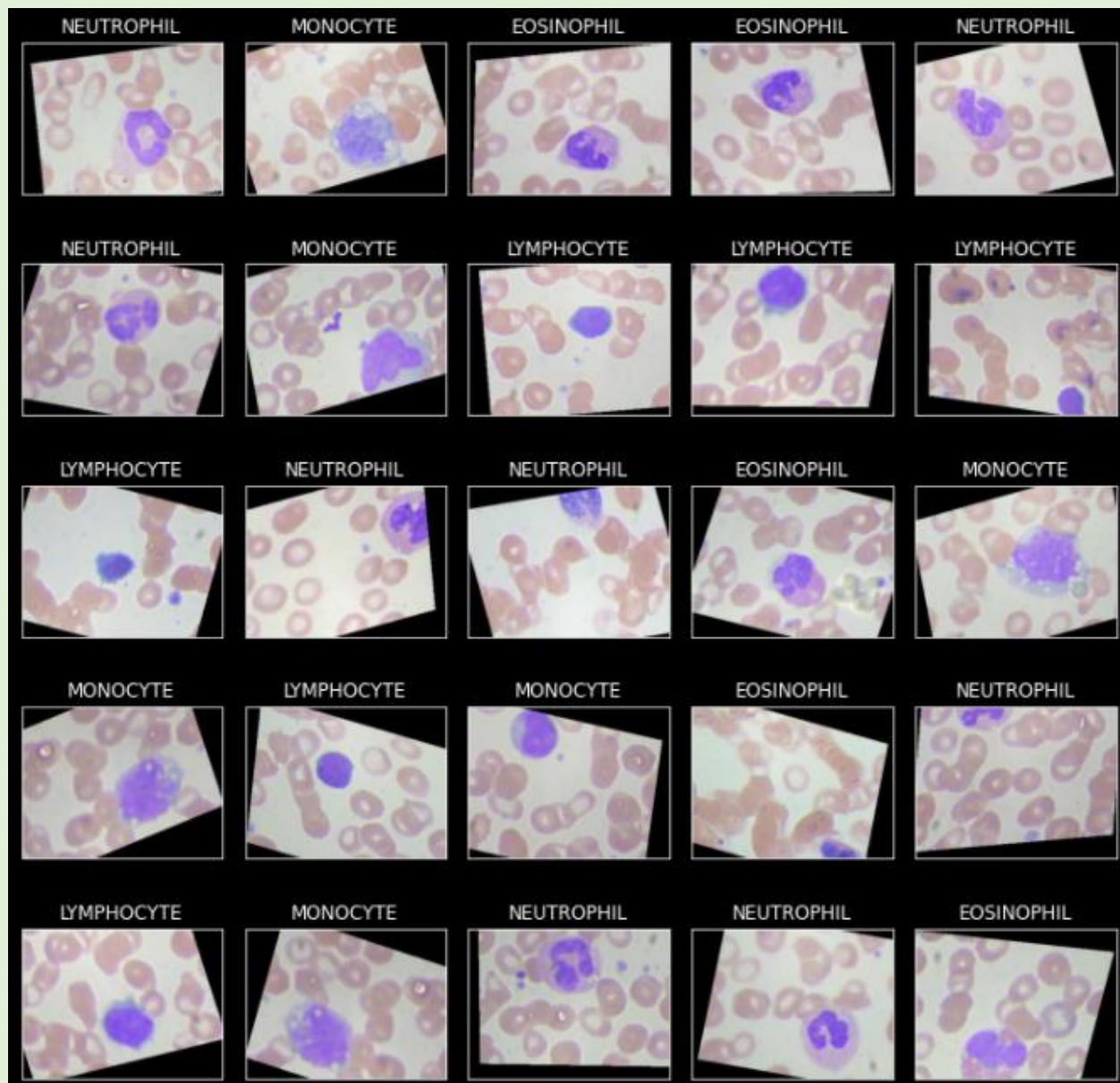
```
Validation_Data_Path = Path("C:/Users/moham/Downloads/dataset2-master/dataset2-master/images/TEST_SIMPLE")
```

```
print("EOSINOPHIL: ", Validation_JPG_Labels.count("EOSINOPHIL"))  
print("LYMPHOCYTE: ", Validation_JPG_Labels.count("LYMPHOCYTE"))  
print("MONOCYTE: ", Validation_JPG_Labels.count("MONOCYTE"))  
print("NEUTROPHIL: ", Validation_JPG_Labels.count("NEUTROPHIL"))
```

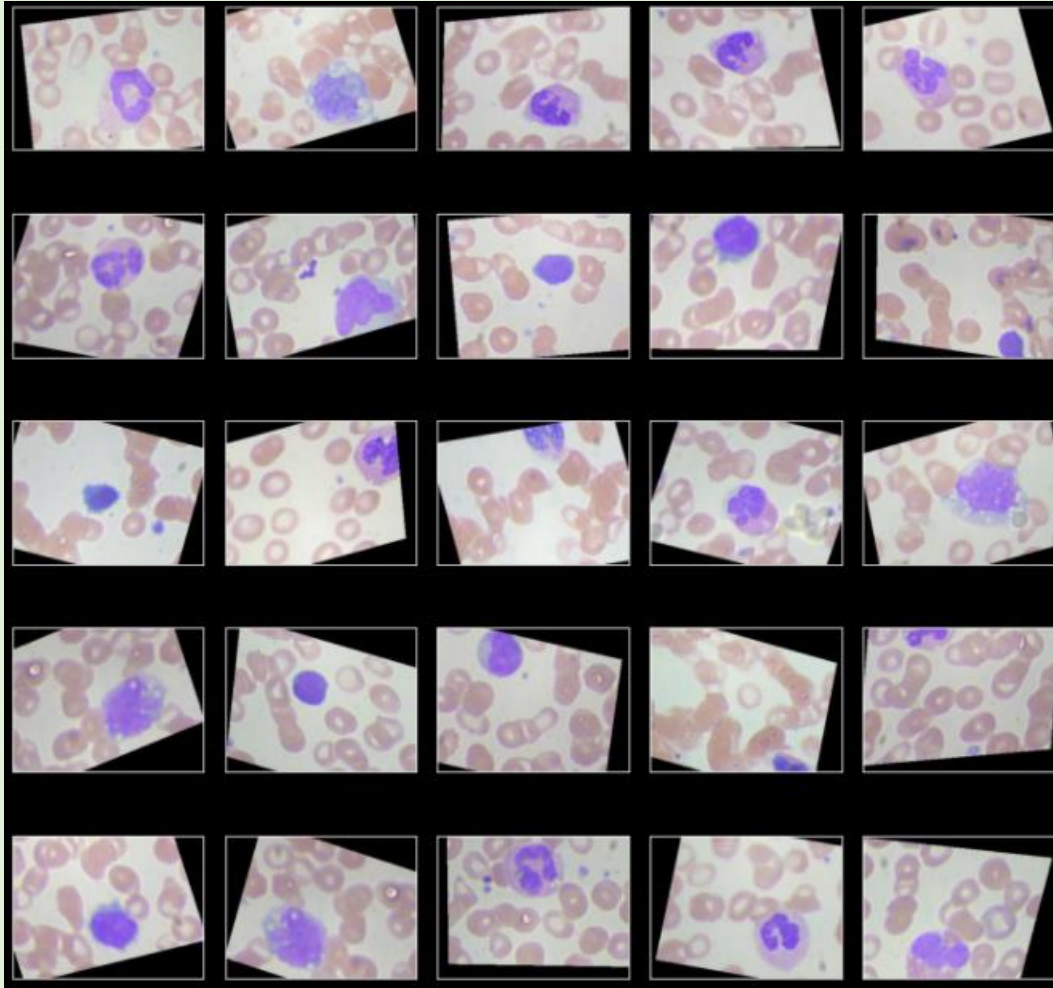
```
EOSINOPHIL: 13  
LYMPHOCYTE: 6  
MONOCYTE: 4  
NEUTROPHIL: 48
```



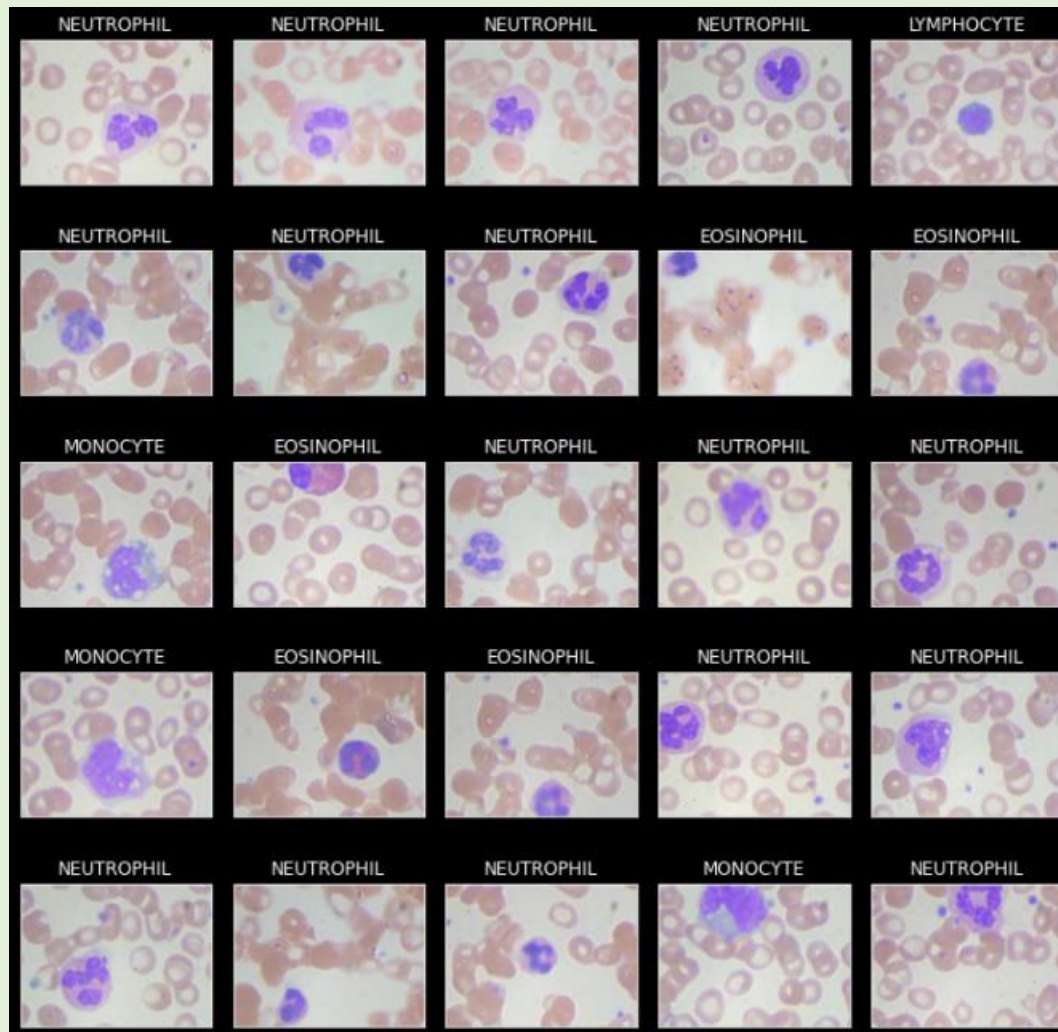
**Let's take a look on Our images that we will train**



## And Our images that we will test



## Here also Our Validation images





## Now we will class Our Data as numbers

```
TRAIN:
{'EOSINOPHIL': 0, 'LYMPHOCYTE': 1, 'MONOCYTE': 2, 'NEUTROPHIL': 3}
[3, 2, 0, 0, 3]
(220, 220, 3)
-----
VALIDATION:
{'EOSINOPHIL': 0, 'LYMPHOCYTE': 1, 'MONOCYTE': 2, 'NEUTROPHIL': 3}
[3, 3, 3, 3, 1]
(220, 200, 3)
-----
TEST:
{'EOSINOPHIL': 0, 'LYMPHOCYTE': 1, 'MONOCYTE': 2, 'NEUTROPHIL': 3}
[3, 1, 1, 1, 0]
(220, 220, 3)
```

**So, we can train it and get result as Predictions of (0,1,2,3)**

**Though...**

**Things will Happens at next pages**



# Let's the Deep-learning Start

## Activation of CNN

### \*CNN STRUCTURE WITH SeparableConv2D\*

```
Model = Sequential()

Model.add(SeparableConv2D(32,3,
                        activation="relu",
                        input_shape=(220,220,3)))
Model.add(BatchNormalization())
Model.add(MaxPooling2D((2)))

#
Model.add(SeparableConv2D(64,3,
                        activation="relu"))
Model.add(SeparableConv2D(128,(3,3),
                        activation="relu"))
Model.add(Dropout(0.5))
Model.add(MaxPooling2D((2)))

#
Model.add(SeparableConv2D(64,3,
                        activation="relu"))
Model.add(SeparableConv2D(128,3,
                        activation="relu"))
Model.add(Dropout(0.5))
Model.add(GlobalAveragePooling2D())

#
Model.add(Flatten())
Model.add(Dense(256,
                activation="relu"))
Model.add(Dropout(0.5))
Model.add(Dense(4,
                activation="softmax"))
```

### Then we add our TensorFlow Keras

```
Call_Back = tf.keras.callbacks.EarlyStopping(monitor="val_accuracy",patience=5,mode="max")
```

### And here just to define loss & accuracy print in our cmd

```
Model.compile(optimizer="rmsprop",loss="categorical_crossentropy",metrics=["accuracy"])
```

**So, Let's train it up**

**Normal we use between 10 epochs to 28 epochs**

**But to get best result & accuracy we will go for 50 epochs**

```
CNN_Model = Model.fit(Train_IMG_Set,  
                        validation_data=Validation_IMG_Set,  
                        callbacks=Call_Back,  
                        epochs=50)
```

**Small look on training...**

```
Epoch 1/50  
312/312 [=====] - 240s 760ms/step - loss: 1.3538 - accuracy: 0.2874 - val_loss: 1.3300 - val_accuracy: 0.1714  
Epoch 2/50  
312/312 [=====] - 687s 2s/step - loss: 1.0500 - accuracy: 0.5460 - val_loss: 1.4686 - val_accuracy: 0.1429  
Epoch 3/50  
312/312 [=====] - 645s 2s/step - loss: 0.6806 - accuracy: 0.7265 - val_loss: 0.7950 - val_accuracy: 0.6000  
Epoch 4/50  
312/312 [=====] - 645s 2s/step - loss: 0.5100 - accuracy: 0.8006 - val_loss: 0.7801 - val_accuracy: 0.7429  
Epoch 5/50  
312/312 [=====] - 664s 2s/step - loss: 0.4175 - accuracy: 0.8404 - val_loss: 0.3906 - val_accuracy: 0.8571  
Epoch 6/50  
312/312 [=====] - 616s 2s/step - loss: 0.3592 - accuracy: 0.8651 - val_loss: 0.5349 - val_accuracy: 0.8286
```

**So, after 8 hours of training, we finally back.....**

**Let's check accuracy of our model now**

```
Model_Results = Model.evaluate(Test_IMG_Set)  
print("LOSS: " + "%.4f" % Model_Results[0])  
print("ACCURACY: " + "%.2f" % Model_Results[1])  
  
78/78 [=====] - 8s 104ms/step - loss: 0.3448 - accuracy: 0.8697  
LOSS: 0.3448  
ACCURACY: 0.87
```

## And here's model summary if anyone interest

```
print(Model.summary())
```

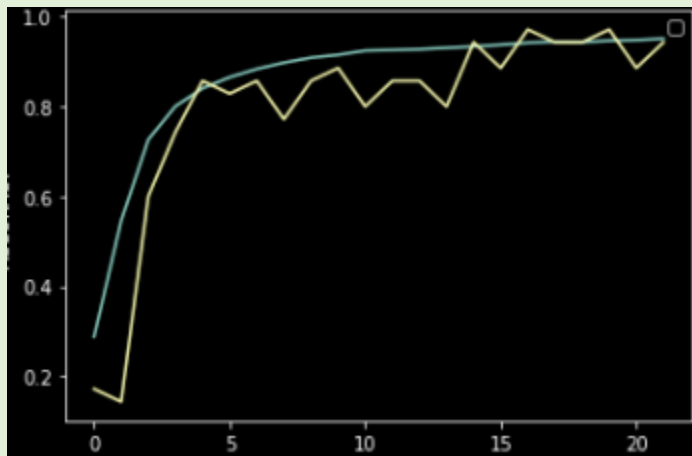
Model: "sequential"

Layer (type)	Output Shape	Param #
separable_conv2d (Separable Conv2D)	(None, 218, 218, 32)	155
batch_normalization (Batch Normalization)	(None, 218, 218, 32)	128
max_pooling2d (MaxPooling2D)	(None, 109, 109, 32)	0
separable_conv2d_1 (Separable Conv2D)	(None, 107, 107, 64)	2400
separable_conv2d_2 (Separable Conv2D)	(None, 105, 105, 128)	8896
dropout (Dropout)	(None, 105, 105, 128)	0
max_pooling2d_1 (MaxPooling2D)	(None, 52, 52, 128)	0
separable_conv2d_3 (Separable Conv2D)	(None, 50, 50, 64)	9408
separable_conv2d_4 (Separable Conv2D)	(None, 48, 48, 128)	8896
dropout_1 (Dropout)	(None, 48, 48, 128)	0
global_average_pooling2d (GlobalAveragePooling2D)	(None, 128)	0
flatten (Flatten)	(None, 128)	0
dense (Dense)	(None, 256)	33024
dropout_2 (Dropout)	(None, 256)	0
dense_1 (Dense)	(None, 4)	1028
Total params: 63,935		
Trainable params: 63,871		
Non-trainable params: 64		

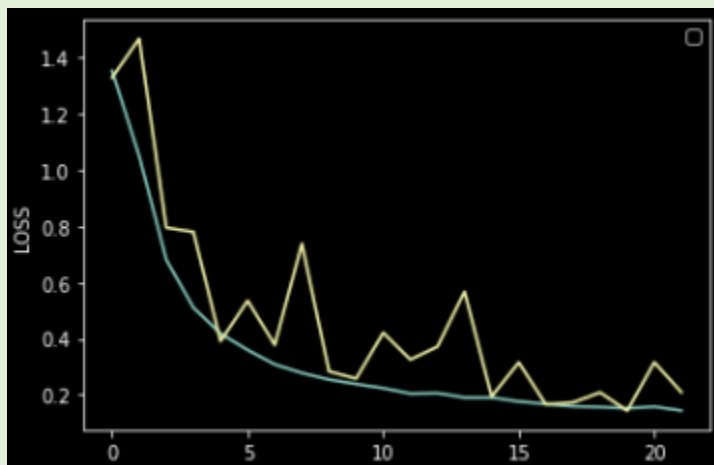
## So, let's start now with some of CNN Benefits

Note (we got best result at 23 epochs so it didn't continue)

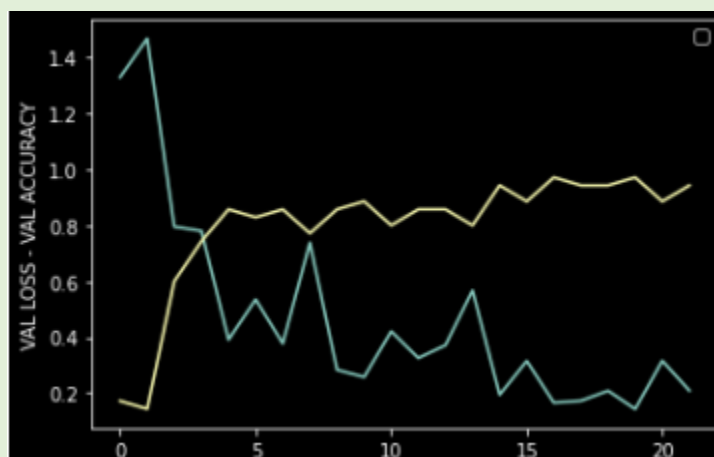
## Accuracy Plot



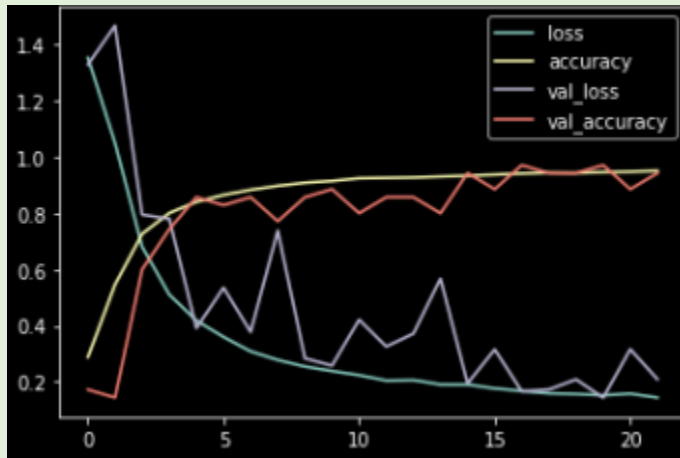
## Loss Plot



## Val Acc vs Val Loss



**And here's everything**

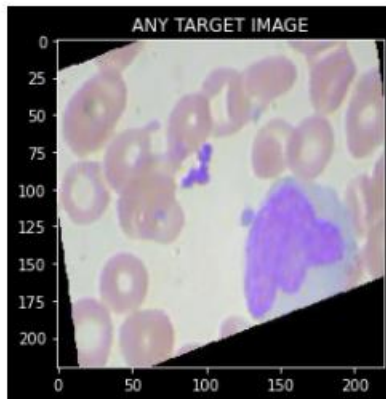


**So, let's go ahead and have some Image Processing**

## Let's Pick a random Image and Play with it

```
Any_IMG = Main_Train_Data["JPG"][6]
IMG = image.load_img(Any_IMG,target_size=(220,220))
Array_IMG = image.img_to_array(IMG)
Array_IMG = np.expand_dims(Array_IMG,axis=0)
Array_IMG /= 255

plt.imshow(Array_IMG[0])
plt.title("ANY TARGET IMAGE")
plt.show()
```



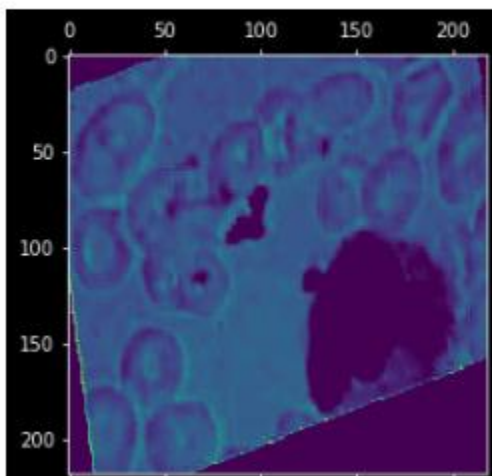
## Apply some layers?

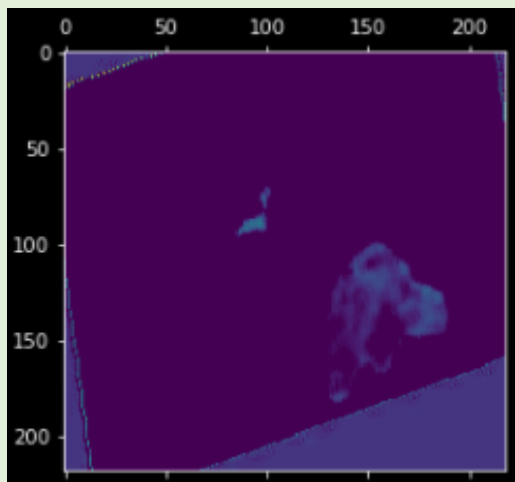
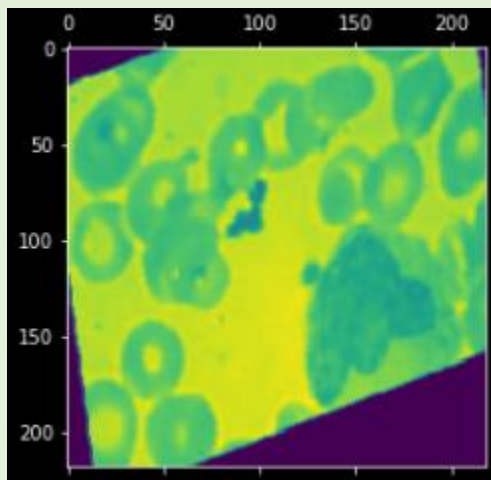
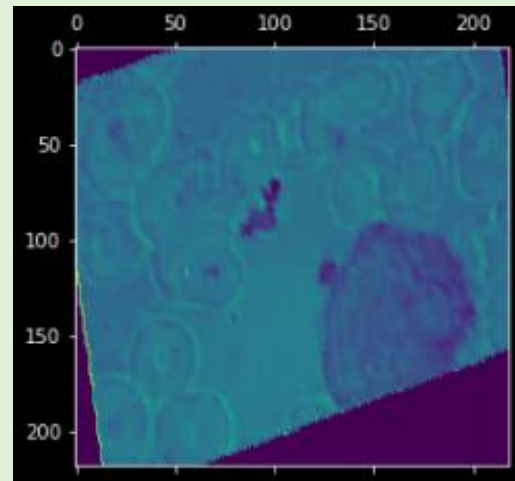
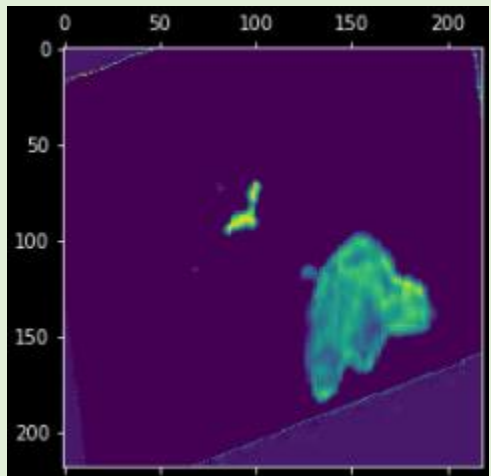
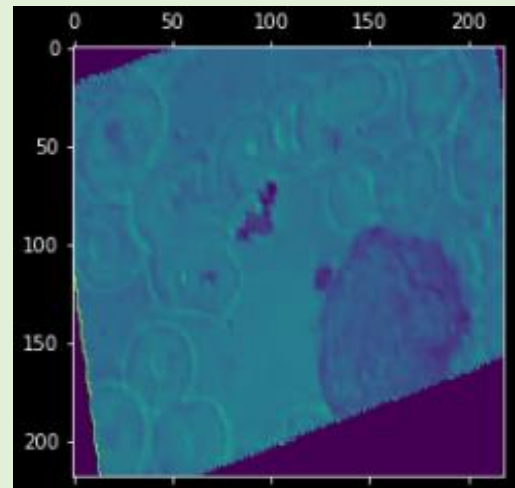
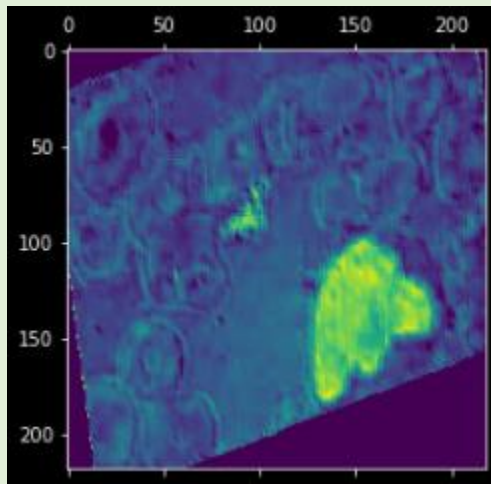
We can apply 0-15 different layer

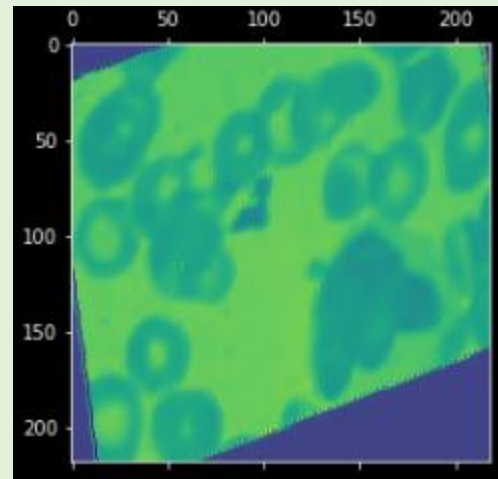
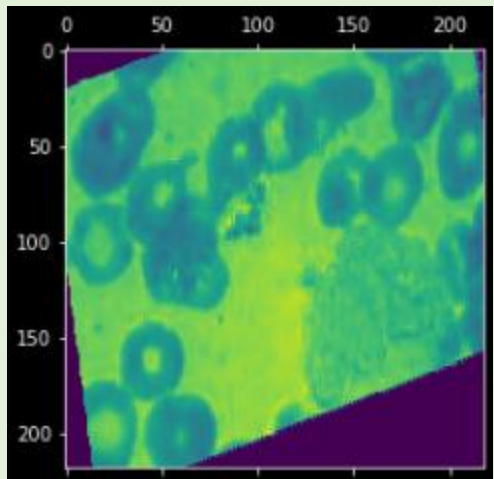
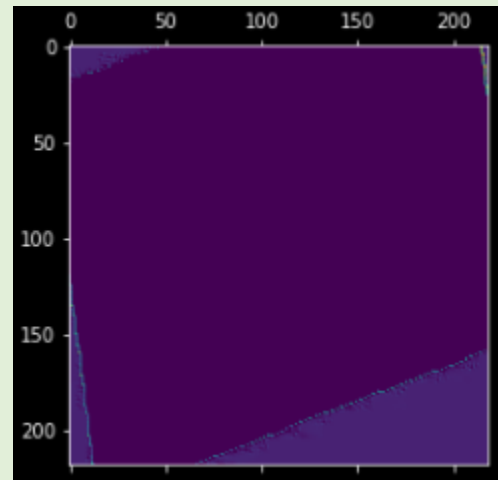
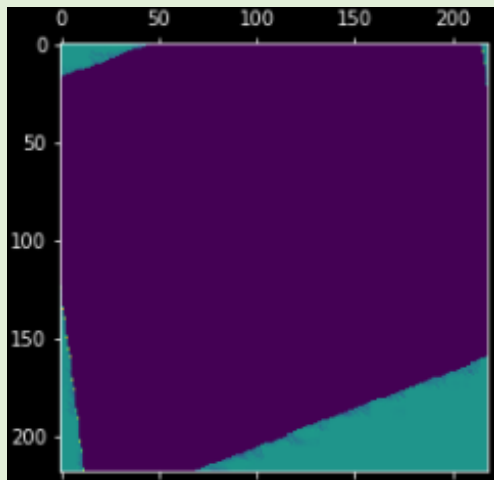
Here's some

```
plt.matshow(first_layer_act[0,:,:,:4],cmap="viridis")
```

<matplotlib.image.AxesImage at 0x2a2deb0a580>







**So, we Almost at the End 😞**



**If you remember we Classed our White Blood Cells  
as 0,1,2,3**

**So, here's the Predication & Classification now  
(EOSINOPHIL 0, LYMPHOCYTE 1, MONOCYTE 2, NEUTROPHIL 3)**

