

Medical project: Lung Cancer Prediction

Problem statement:

Predict whether the patient have lung cancer or not using the below features.

	Name	Surname	Age	Smokes	AreaQ	Alkhol	Result
0	John	Wick	35	3	5	4	1
1	John	Constantine	27	20	2	5	1
2	Camela	Anderson	30	0	5	2	0
3	Alex	Telles	28	0	8	1	0
4	Diego	Maradona	68	4	5	6	1

Result column is target variable which says whether the patient have cancer or not.

Exploratory Data Analysis – EDA:

Check for null values and data type in dataset.

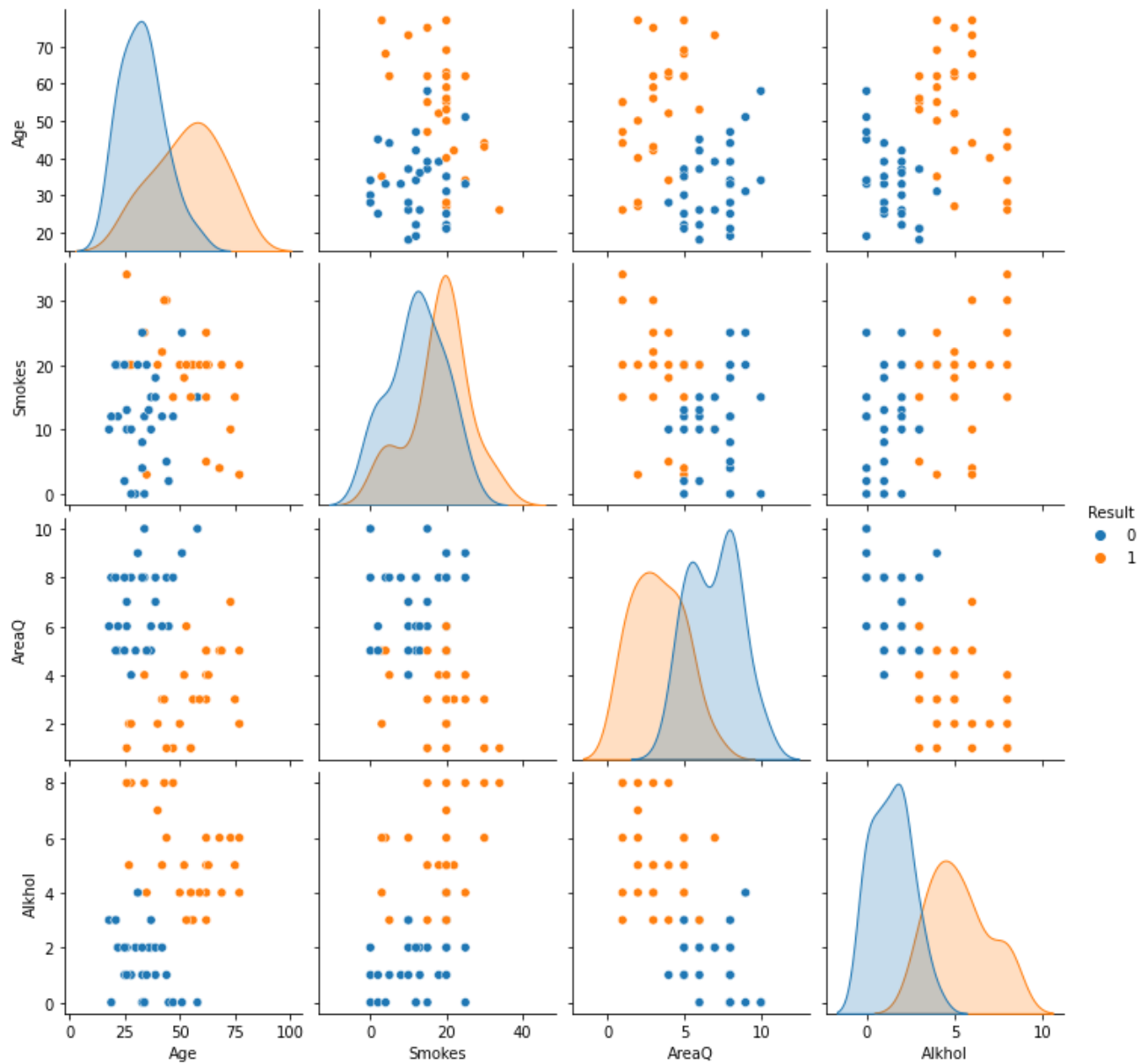
```
] data.isnull().sum()
```

```
Name      0
Surname    0
Age        0
Smokes     0
AreaQ      0
Alkhol     0
Result     0
dtype: int64
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 59 entries, 0 to 58
Data columns (total 7 columns):
#   Column      Non-Null Count  Dtype
---  ---
0   Name        59 non-null    object
1   Surname     59 non-null    object
2   Age         59 non-null    int64
3   Smokes      59 non-null    int64
4   AreaQ       59 non-null    int64
5   Alkhol      59 non-null    int64
6   Result      59 non-null    int64
dtypes: int64(5), object(2)
memory usage: 3.4+ KB
```

- No null values
- All necessary variables are numeric
- Hence no action required in data pre-processing

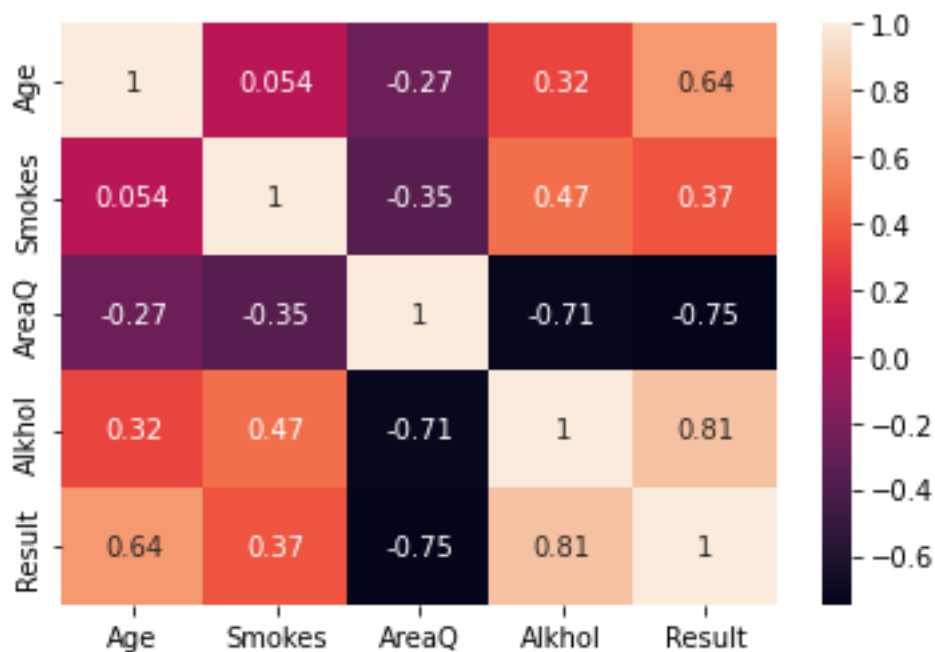
Pair plot among all numeric features:



Observations:

- Lower the age and smoke, less chance of lung cancer
- Higher the alkhol and lower the areaQ, more chance of lung cancer.
- This clear colour segregation in each plot shows that each variable have significant effect on lung cancer.

Correlation heatmap:



Observations:

- Features – Age and Alkhol have strong positive correlation
- This indicates that if age and alkhol level increases, more possibility of lung cancer.
- Feature: AreaQ have strong negative correlation, this indicates that how small the values – AreaQ that much high possibility of getting lung cancer.

ANN - Model Summary:

Model: "sequential_5"

Layer (type)	Output Shape	Param #
dense_16 (Dense)	(None, 128)	640
dense_17 (Dense)	(None, 64)	8256
dense_18 (Dense)	(None, 1)	65
Total params: 8,961		
Trainable params: 8,961		
Non-trainable params: 0		

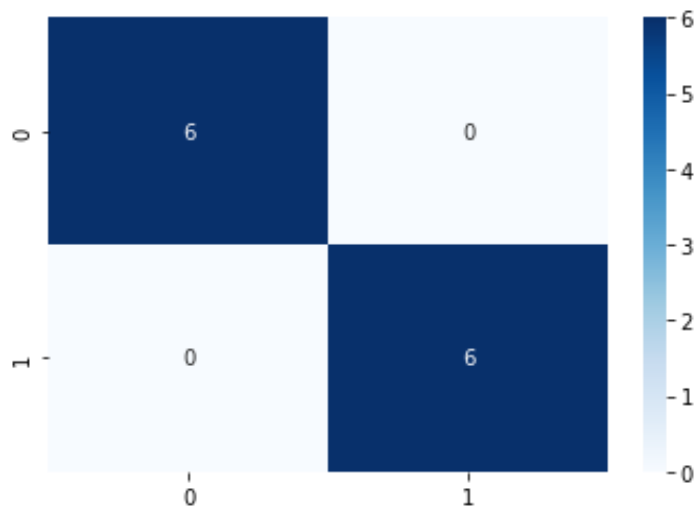
Hidden layer 1 – 128 neurons

Hidden layer 2 – 64 neurons

Output layer – 1 neuron with sigmoid activation

Performance:

Confusion matrix



No false positive and false negative which means accuracy is 100%.

Classification report:

```
!3] print(classification_report(y_test,y_pred))
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	6
1	1.00	1.00	1.00	6
accuracy			1.00	12
macro avg	1.00	1.00	1.00	12
weighted avg	1.00	1.00	1.00	12

All metrics are 100%, our model works in best way.

Hyper-parameter tuning:

```
Accuracy of the model with 4, 64 neurons in each hidden layers: 0.4166666666666667
Accuracy of the model with 16, 64 neurons in each hidden layers: 0.8333333333333334
Accuracy of the model with 32, 64 neurons in each hidden layers: 0.9166666666666666
Accuracy of the model with 64, 64 neurons in each hidden layers: 1.0
Accuracy of the model with 128, 64 neurons in each hidden layers: 1.0
```

As we increase the no of neurons in the first hidden layer, we obtain the best accuracy.

Different optimizers with 20 epochs:

```
Accuracy of the model with <keras.optimizer_v2.adam.Adam object at 0x7faf8810fed0>: 1.0
Accuracy of the model with <keras.optimizer_v2.gradient_descent.SGD object at 0x7faf8810fc10>: 0.8333333333333334
Accuracy of the model with <keras.optimizer_v2.rmsprop.RMSprop object at 0x7faf8810f950>: 0.9166666666666666
Accuracy of the model with <keras.optimizer_v2.adadelta.Adadelta object at 0x7faf8810fad0>: 0.5
Accuracy of the model with <keras.optimizer_v2.adagrad.Adagrad object at 0x7faf8810fe50>: 1.0
```

Adam and Adagrad is the best optimizer for our model which gives 100% accuracy.

New input is given - predict the output of your model:

```
▶ age = int(input("Age: "))
  smoke = int(input("Smokes: "))
  areaq = int(input("AreaQ: "))
  alkhol = int(input("Alkhol: "))

  xnew = [[age,smoke,areaq,alkhol]]
  pred = model.predict(xnew)
  y_pred=(pred>0.5)
  if y_pred==True:
      print("The person has Lung cancer")
  else:
      print("Free from lung cancer")

Age: 77
Smokes: 20
AreaQ: 5
Alkhol: 4
The person has Lung cancer
```

Based on the independent variables our model predicted that the patient has lung cancer.

Conclusion:

Our model performed well with below parameters

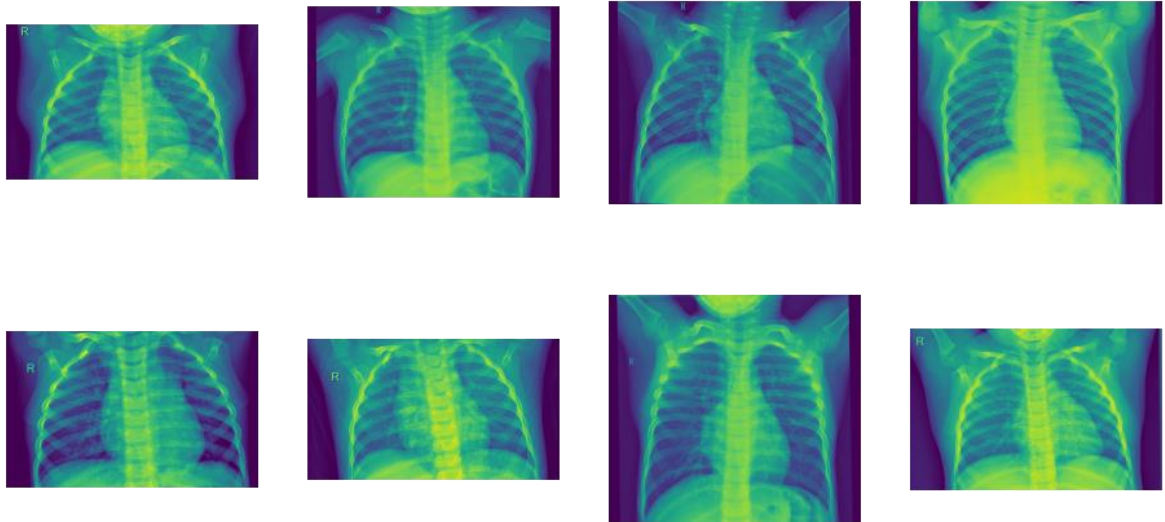
- 20 epochs
- Two hidden layers
- First hidden layers: 128 neurons
- Second hidden layers: 64 neurons.
- Optimizer: Adam or Adagrad

Covid X-Ray Classification

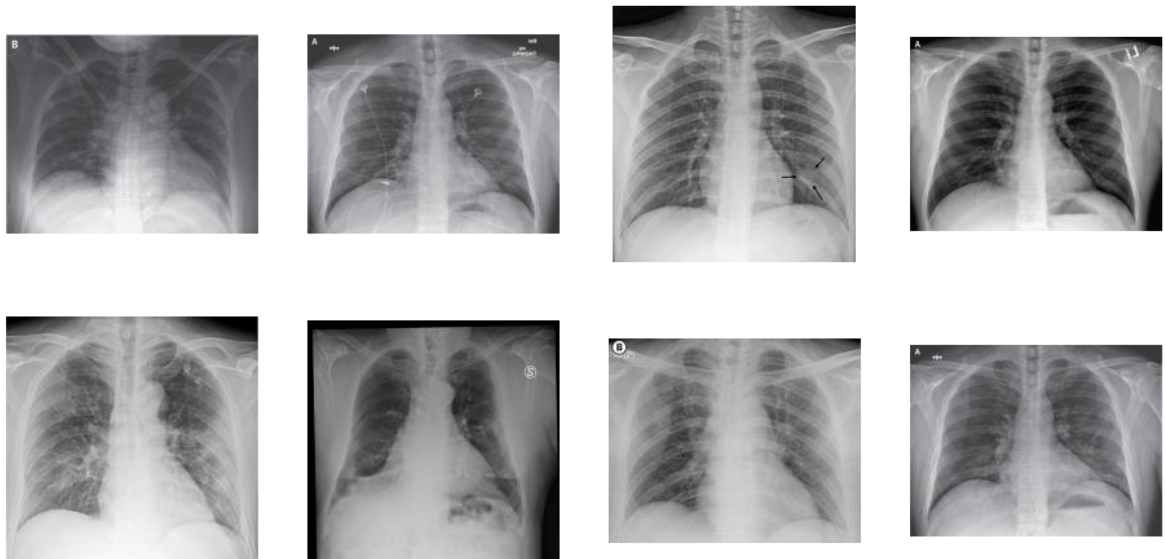
Problem statement:

Given with the X-Ray images of the covid-19 patients, predict whether the person has Covid or not.

Normal person X-ray images

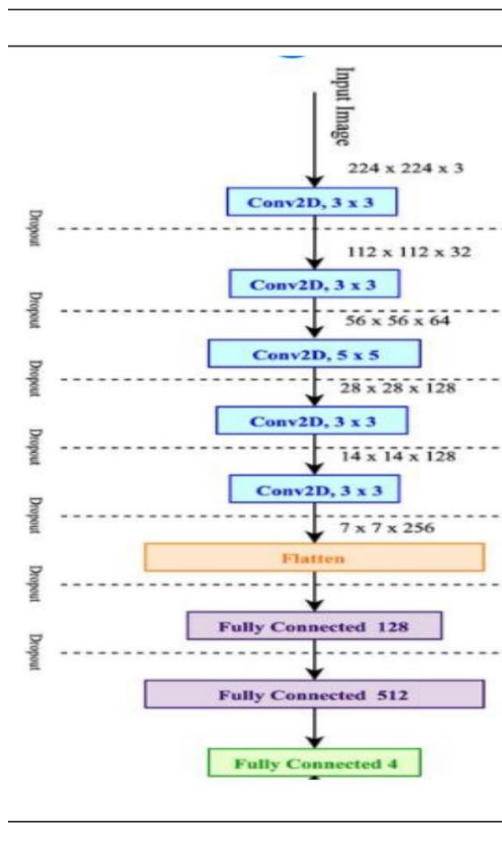


Covid patient - X-ray images



Building the Models:

Model 1 - CNN model given in the below figure.



We implemented the above architecture.

Model summary:

Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 112, 112, 32)	896
dropout (Dropout)	(None, 112, 112, 32)	0
max_pooling2d (MaxPooling2D)	(None, 56, 56, 32)	0
conv2d_1 (Conv2D)	(None, 56, 56, 64)	18496
dropout_1 (Dropout)	(None, 56, 56, 64)	0
max_pooling2d_1 (MaxPooling2D)	(None, 28, 28, 64)	0
conv2d_2 (Conv2D)	(None, 28, 28, 128)	204928
dropout_2 (Dropout)	(None, 28, 28, 128)	0
max_pooling2d_2 (MaxPooling2D)	(None, 14, 14, 128)	0
conv2d_3 (Conv2D)	(None, 14, 14, 128)	147584
dropout_3 (Dropout)	(None, 14, 14, 128)	0
max_pooling2d_3 (MaxPooling2D)	(None, 7, 7, 128)	0

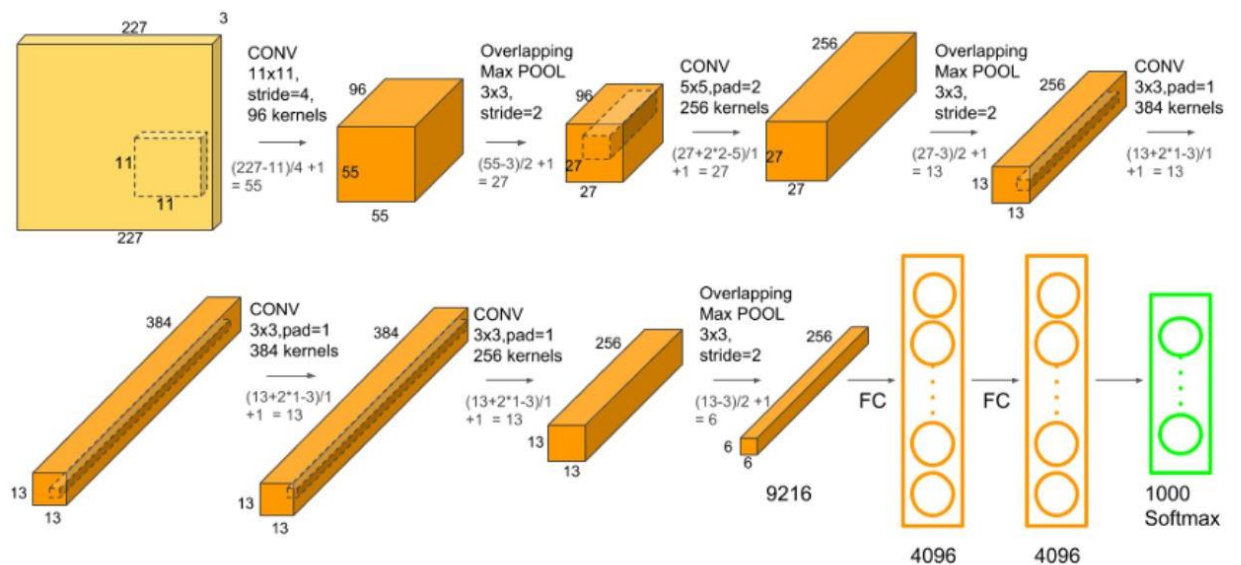
conv2d_4 (Conv2D)	(None, 7, 7, 256)	295168
dropout_4 (Dropout)	(None, 7, 7, 256)	0
max_pooling2d_4 (MaxPooling 2D)	(None, 3, 3, 256)	0
flatten (Flatten)	(None, 2304)	0
dropout_5 (Dropout)	(None, 2304)	0
dense (Dense)	(None, 128)	295040
dropout_6 (Dropout)	(None, 128)	0
dense_1 (Dense)	(None, 512)	66048
dense_2 (Dense)	(None, 4)	2052
dense_3 (Dense)	(None, 1)	5

=====

Total params: 1,030,217
Trainable params: 1,030,217
Non-trainable params: 0

Model – 2 – AlexNet architecture:

We implemented the below structure.



Changes in Output layer:

1 neuron and sigmoid – Activation function.

Model summary:

model: "sequential_1"

Layer (type)	Output Shape	Param #
conv2d_5 (Conv2D)	(None, 54, 54, 96)	34944
max_pooling2d_5 (MaxPooling 2D)	(None, 26, 26, 96)	0
conv2d_6 (Conv2D)	(None, 26, 26, 256)	614656
max_pooling2d_6 (MaxPooling 2D)	(None, 12, 12, 256)	0
conv2d_7 (Conv2D)	(None, 12, 12, 384)	885120
conv2d_8 (Conv2D)	(None, 12, 12, 384)	1327488
conv2d_9 (Conv2D)	(None, 12, 12, 256)	884992
max_pooling2d_7 (MaxPooling 2D)	(None, 5, 5, 256)	0
flatten_1 (Flatten)	(None, 6400)	0
dense_4 (Dense)	(None, 4096)	26218496
dropout_7 (Dropout)	(None, 4096)	0
dense_5 (Dense)	(None, 4096)	16781312
dropout_8 (Dropout)	(None, 4096)	0
dense_6 (Dense)	(None, 1)	4097

=====
Total params: 46,751,105
Trainable params: 46,751,105
Non-trainable params: 0
=====

Performance:

4/4 [=====] - 19s 35/step - loss: 0.2680 - accuracy: 0.8917 - val_loss

```
[ ] cnn_model.evaluate(test_dataset)
```

2/2 [=====] - 4s 3s/step - loss: 0.5372 - accuracy: 0.9750
[0.5371925234794617, 0.9750000238418579]

```
[ ] alex_model.evaluate(test_dataset)
```

2/2 [=====] - 2s 273ms/step - loss: 0.0719 - accuracy: 0.9750
[0.07185763120651245, 0.9750000238418579]

Figure based CNN model – Model 1:

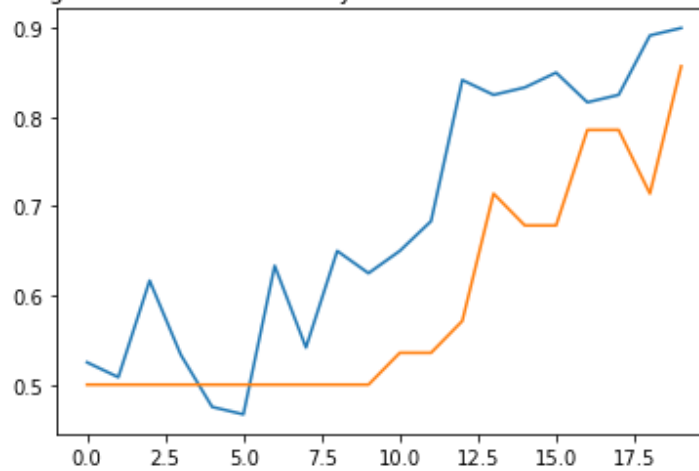
- Accuracy: 97.5%
- Loss: 0.53

AlexNet Model -2 :

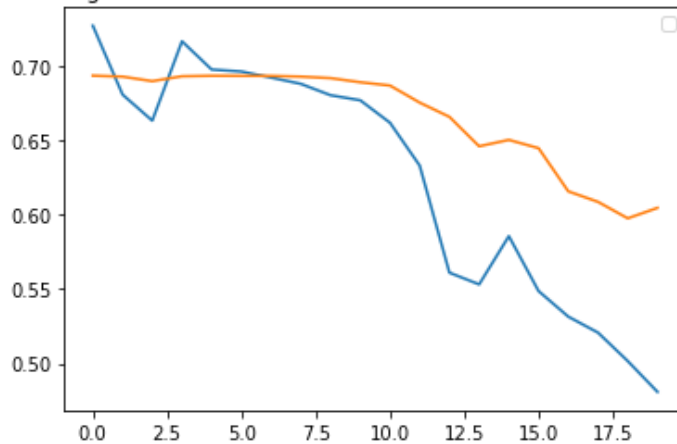
- Accuracy: 97.5%
- Loss: 0.07

Accuracy and Loss plots:

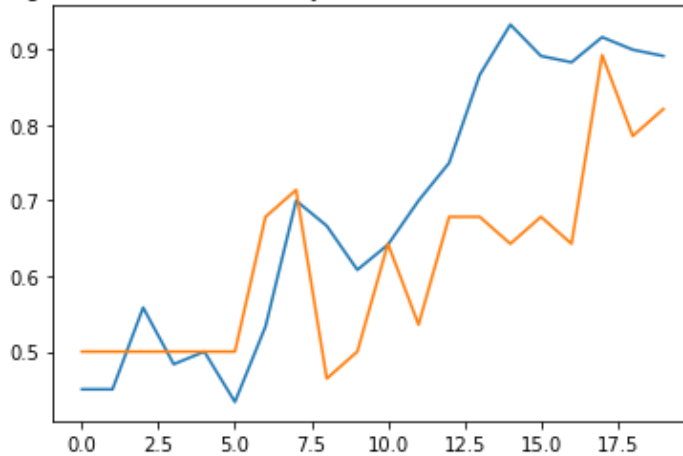
Training and validation accuracy of CNN Model in Covid X-RAY Dataset



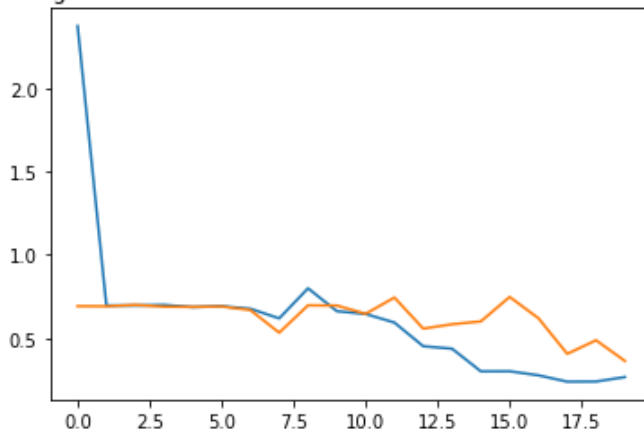
Training and validation loss of CNN Model in Covid X-RAY Dataset



Training and validation accuracy of Alexnet Model in Covid X-RAY Dataset



Training and validation loss of Alexnet Model in Covid X-RAY Dataset



Conclusion:

- In both models, training and validation – loss is decreasing as epoch increases
- Training and validation – accuracy is increasing as epoch increases
- This indicates the model is best fit, there's no underfit and overfit.
- Both a models performed well equally with same high accuracy (97.5%)
- However, model 2 is elected as best one because it has lesser loss ($0.07 < 0.53$)
- Therefore AlexNet model is best.

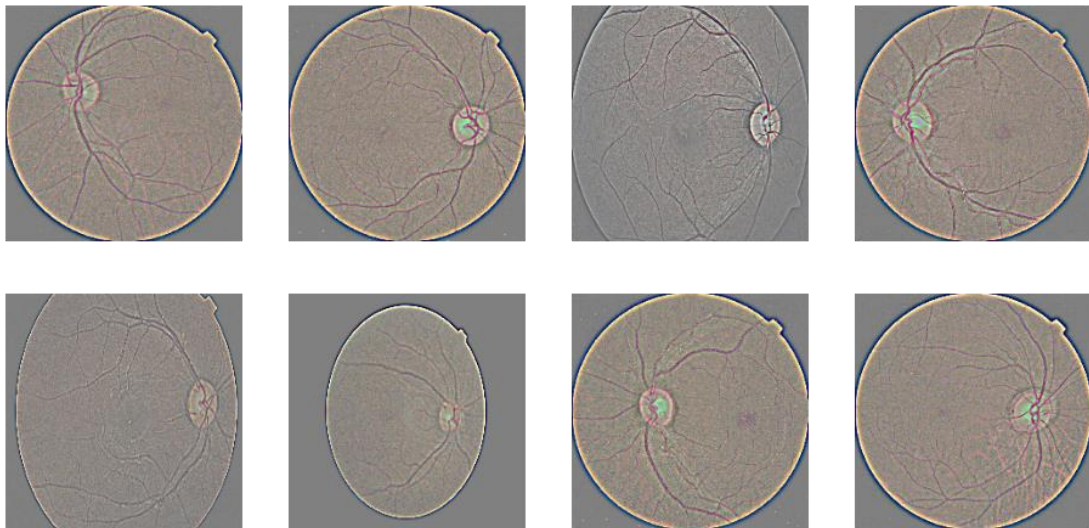
Diabetic Retinopathy – Retina images classification

Problem statement:

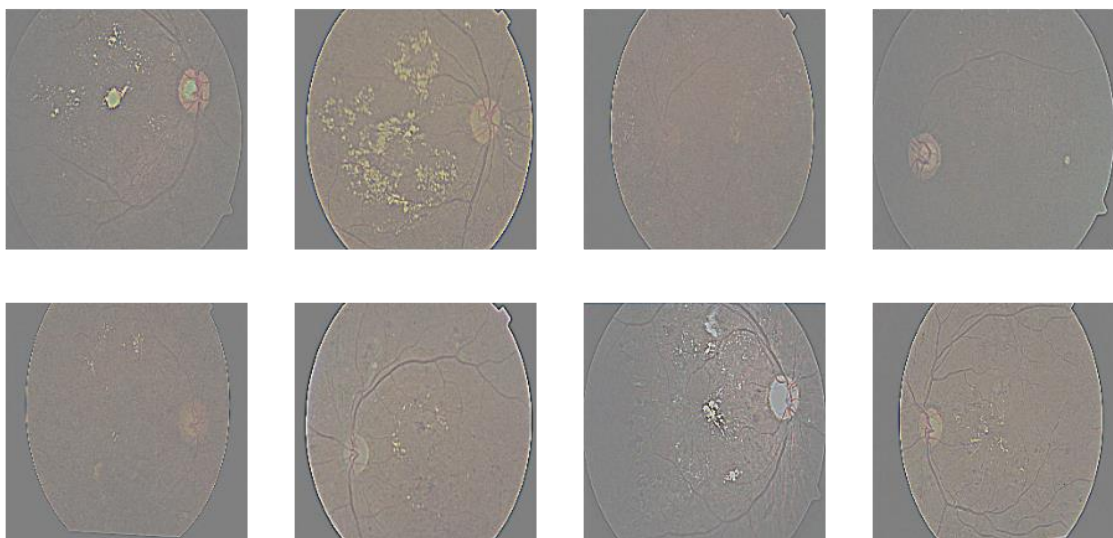
Given with the retina images of patients, classify the patients based on the below 5 conditions:

- **No DR** : normal person
- **Severe**: Having extreme diabetic retinopathy
- **Mild**: starting stage of DR
- **Moderate**: next stage of mild
- **Proliferate**: rapidly growing DR.

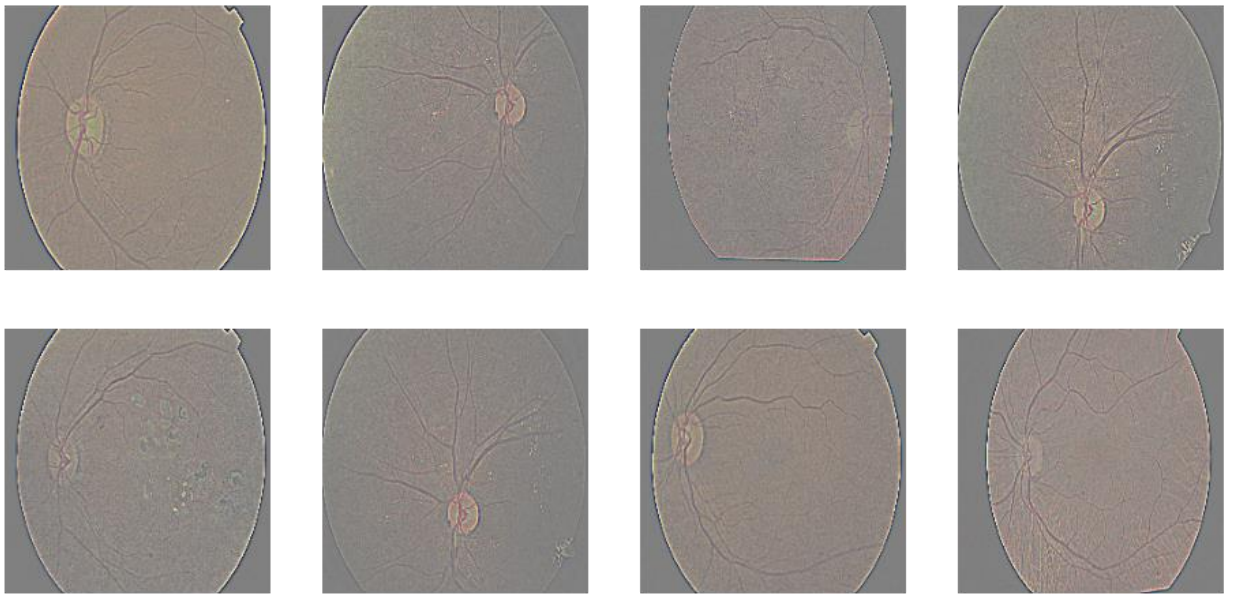
Normal



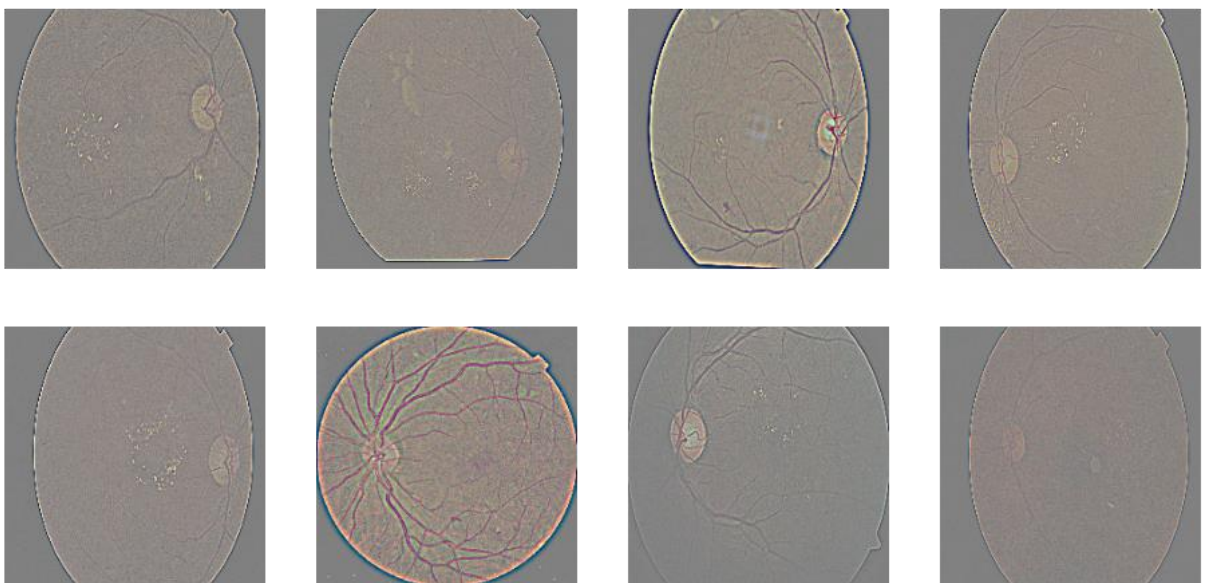
Severe



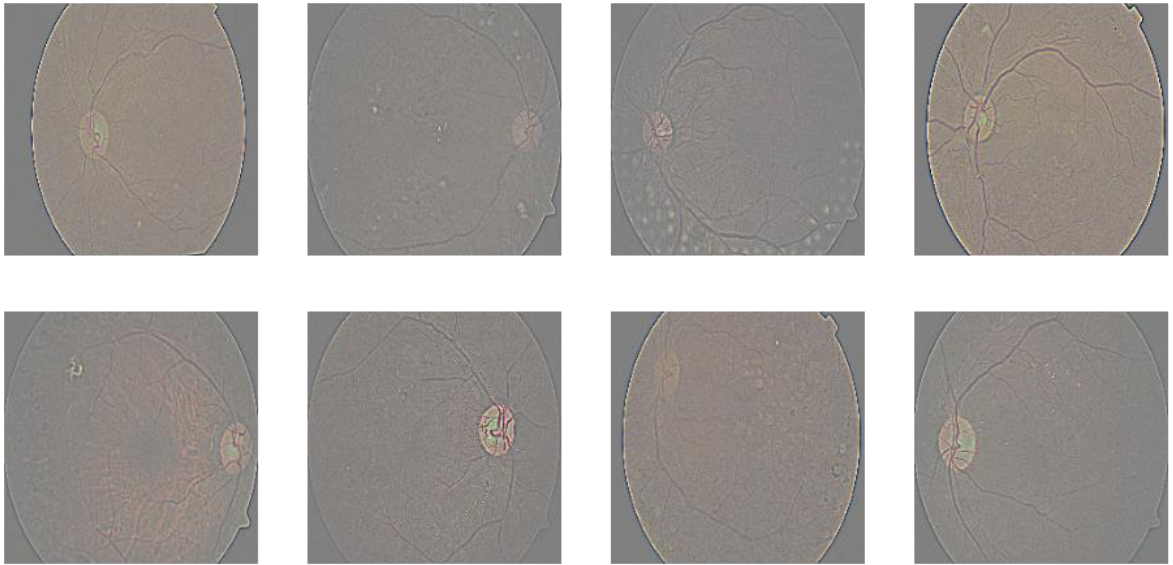
Mild



Moderate



Proliferate



Models:

We implemented the same figure based CNN and AlexNet model for this dataset.

However, the output layer neurons will be 5 because we have 5 class variables and activation function is Softmax.

Performance:

```
cnn_model.evaluate(test_dataset)
```

```
6/6 [=====] - 2s 363ms/step - loss: 1.0436 - accuracy: 0.6222  
[1.04362154006958, 0.622222447395325]
```

```
alex_model.evaluate(test_dataset)
```

```
6/6 [=====] - 6s 945ms/step - loss: 1.2136 - accuracy: 0.5000  
[1.2135668992996216, 0.5]
```

CNN model:

Accuracy: 62%

Loss: 1

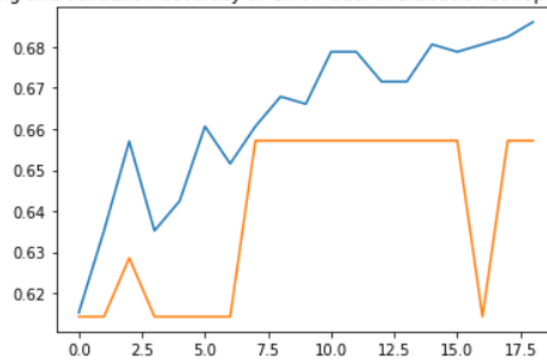
AlexNet model:

Accuracy: 50%

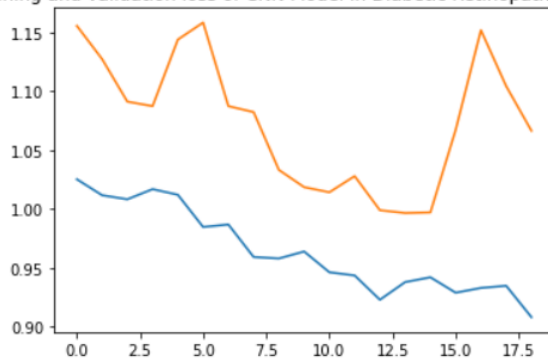
Loss: 1.21

Training and Validation – Accuracy and loss plot:

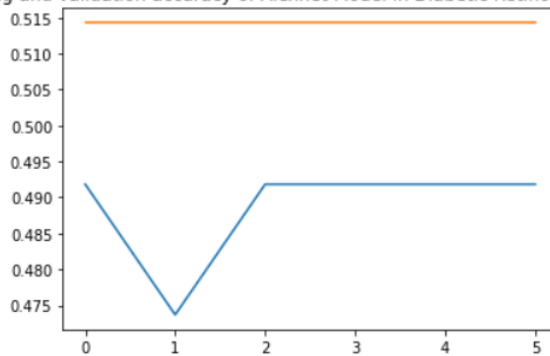
Training and validation accuracy of CNN Model in Diabetic Retinopathy Dataset



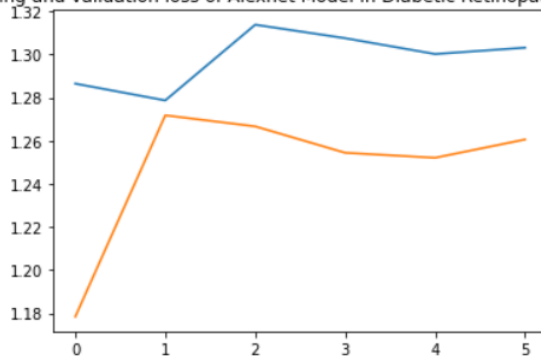
Training and validation loss of CNN Model in Diabetic Retinopathy Dataset



Training and validation accuracy of Alexnet Model in Diabetic Retinopathy Dataset



Training and validation loss of Alexnet Model in Diabetic Retinopathy Dataset



Conclusion:

- Training and test - Accuracy is increasing better in CNN model.
- Training and test – Loss are decreasing well in CNN model.
- Flat line in accuracy plot of AlexNet model shows that accuracy haven't increase though the epochs are increase.
- This indicates the weakness of the model, the training itself insufficient.
- CNN model have higher accuracy and lower loss in comparison of AlexNet.
- Therefore, we can conclude that CNN model performed well in DR dataset.

Saving the weights of the best model

```
[ ] model_final.save("/content/MyDrive/models/diabeticretino.h5")  
    print("Saved model to disk")  
    model_final.save_weights("/content/MyDrive/models/diabeticretino.h5")  
    print("Saved weights to disk")
```