import Libraries

```
In [1]: import tensorflow as tf
        import keras
        from keras.layers import Conv2D,MaxPooling2D,Dense,Dropout,Flatten
        import numpy as np
        import matplotlib.pyplot as plt
        import pandas as pd
        from keras.utils import to_categorical
        import os
        from tqdm import tqdm
        import cv2
        import random
        import pickle as pk
        from sklearn.model_selection import train_test_split
        import numpy as np
        import pandas as pd
        from keras.models import load_model
        from keras.layers import BatchNormalization
        from sklearn.model_selection import GridSearchCV
        from keras.wrappers.scikit_learn import KerasClassifier
        from keras.callbacks import EarlyStopping
        from keras import regularizers
        from sklearn.metrics import roc_curve, auc
        import matplotlib.pyplot as plt
        import keras
        from sklearn.metrics import classification_report
        from sklearn.metrics import confusion_matrix
In [2]: #define constants
        imges num=5000
        class_num=10
        FAST RUN = False
        IMAGE_WIDTH=220
        IMAGE HEIGHT=220
        IMAGE_SIZE=(IMAGE_WIDTH, IMAGE_HEIGHT)
        IMAGE CHANNELS=3
In [3]: # training
        trainingPath=r"C:\Users\kirti\OneDrive\Desktop\MLPA\chest xray\chest_xray\train"
        categories=os.listdir(trainingPath)
        categories
Out[3]: ['NORMAL', 'PNEUMONIA']
In [4]: #save class names in dictionary
        classnames={}
        for i in range(2):
          classnames[i]=categories[i]
        classnames
```

```
Out[4]: {0: 'NORMAL', 1: 'PNEUMONIA'}
In [5]: #load the images and read it and save eacah image with folder name
        training =[]
        x=0
        for category in categories:
            folder=os.path.join(trainingPath, category)
            # print(folder)
            x2=0
             for file in tqdm(os.listdir(folder)):
                 # print(file)
                 try:
                     img=cv2.imread(os.path.join(folder,file))
                     img=cv2.cvtColor(img,cv2.COLOR_BGR2RGB)
                     img=cv2.resize(img,IMAGE_SIZE)# size image for CNN input
                     classNum=categories.index(category) #detrmine class
                     training.append([img,classNum])
                     # plt.imshow(img)
                     # plt.show()
                 except Exception as e:
                     pass
                 if x2>=imges_num:
                     break
                 x2+=1
             if x>=class_num-1:
                 break
             x+=1
                      | 1341/1341 [00:45<00:00, 29.36it/s]
                        3875/3875 [00:35<00:00, 110.06it/s]
In [6]:
        len(training)
Out[6]: 5216
In [7]:
        #shuffle my data to avoid overfitting
        training[0]
        random.shuffle(training)
        for trainSample in training[:10]:
             print(trainSample[1])
       1
       0
       0
       1
       1
       1
       1
       1
       1
       1
In [8]: #split features and labels
        x_train_data=[]
        y_train_data=[]
```

```
for feature,label in training:
    x_train_data.append(feature)
    y_train_data.append(label)

In [9]: #convert my data to nunpy array
    x_train_data=np.array(x_train_data)
    y_train_data=np.array(y_train_data)

In [10]: #reshape data to RGB form #3 channels with 220*220

    x_train_data=x_train_data.reshape(-1,IMAGE_HEIGHT,IMAGE_WIDTH,IMAGE_CHANNELS)
    print(x_train_data.shape)
    print(y_train_data.shape)

(5216, 220, 220, 3)
(5216,)
```

test data

```
In [11]: testPath=r"C:\Users\kirti\OneDrive\Desktop\MLPA\chest xray\chest_xray\test"
         categories=os.listdir(trainingPath)
         categories
Out[11]: ['NORMAL', 'PNEUMONIA']
In [12]: #load the images and read it and save eacah image with folder name [class name]
         test =[]
         x=0
         for category in categories:
              folder=os.path.join(testPath, category)
              # print(folder)
             x2=0
              for file in tqdm(os.listdir(folder)):
                  # print(file)
                  try:
                      img=cv2.imread(os.path.join(folder,file))
                      img=cv2.cvtColor(img,cv2.COLOR_BGR2RGB)
                      img=cv2.resize(img,IMAGE SIZE)# size image for CNN input
                      classNum=categories.index(category) #detrmine class
                      test.append([img,classNum])
                      # plt.imshow(img)
                      # plt.show()
                  except Exception as e:
                      pass
                  if x2>imges num:
                      break
                  x2+=1
              if x>=class num-1:
                  break
              x+=1
              # print(x)
```

```
| 234/234 [00:05<00:00, 44.30it/s]
        100%
                       | 390/390 [00:03<00:00, 107.70it/s]
In [13]: #shuffle my data to avoid overfitting
         random.shuffle(test)
         for trainSample in test[:10]:
             print(trainSample[1])
        1
        1
        1
        0
        1
        1
        0
        1
        1
In [14]: #split features and labels
         x_test=[]
         y_test=[]
         for feature,label in test:
             x_test.append(feature)
             y_test.append(label)
In [15]: #convert my data to nunpy array
         x_test=np.array(x_test)
         y_test=np.array(y_test)
In [16]: #reshape data to RGB form 3 channels with 220*220
         x_test=x_test.reshape(-1,IMAGE_HEIGHT,IMAGE_WIDTH,IMAGE_CHANNELS)
         print(x_test.shape)
         print(y_test.shape)
        (624, 220, 220, 3)
        (624,)
```

split data

```
In [17]: #split the train data into train and validation

# x_train=x_train_data
# y_train=y_train_data

x_train,x_validation_data,y_train,y_validation_data=train_test_split(x_train_dat print(x_train.shape)

print(y_train.shape)

print(x_validation_data.shape)

print(y_validation_data.shape)
```

```
print(x_test.shape)
print(y_test.shape)

(4433, 220, 220, 3)
(4433,)
(783, 220, 220, 3)
(783,)
(624, 220, 220, 3)
(624,)
```

shapes of data

```
In [18]: print(f'x_train shape :
                                         {x train.shape}')
         print(f'y_train shape :
                                         {y_train.shape}\n')
         print(f'x_validation shape :
                                        {x_validation_data.shape}')
         print(f'y_validation shape :
                                         {y_validation_data.shape}\n')
         print(f'x_test shape :
                                        {x_test.shape}')
         print(f'x_test shape :
                                         {y_test.shape}')
        x_train shape :
                              (4433, 220, 220, 3)
        y_train shape :
                               (4433,)
                             (783, 220, 220, 3)
        x_validation shape :
       y_validation shape :
                               (783,)
                               (624, 220, 220, 3)
       x_test shape :
        x_test shape :
                                (624,)
```

Build Model

```
In [19]: model=keras.models.Sequential()
         model.add(Conv2D(16,(3,3),input_shape=(IMAGE_WIDTH,IMAGE_HEIGHT,IMAGE_CHANNELS),
         model.add(BatchNormalization())
         model.add(MaxPooling2D(pool_size=(2,2)))
         model.add(Dropout(0.2))
         model.add(Conv2D(64,(3,3), activation='relu'))
         model.add(BatchNormalization())
         model.add(MaxPooling2D(pool_size=(2,2)))
         model.add(Dropout(0.4))
         model.add(Flatten())
         model.add(Dense(256, activation='relu'))
         model.add(BatchNormalization())
         model.add(Dropout(0.5))
         model.add(Dense(1,activation='sigmoid'))
In [20]: model.compile(optimizer='adam',loss='binary_crossentropy',metrics=['accuracy'])
In [21]: model.summary()
```

Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 218, 218, 16)	448
<pre>batch_normalization (BatchN ormalization)</pre>	(None, 218, 218, 16)	64
<pre>max_pooling2d (MaxPooling2D)</pre>	(None, 109, 109, 16)	0
dropout (Dropout)	(None, 109, 109, 16)	0
conv2d_1 (Conv2D)	(None, 107, 107, 64)	9280
<pre>batch_normalization_1 (Batc hNormalization)</pre>	(None, 107, 107, 64)	256
<pre>max_pooling2d_1 (MaxPooling 2D)</pre>	(None, 53, 53, 64)	0
dropout_1 (Dropout)	(None, 53, 53, 64)	0
flatten (Flatten)	(None, 179776)	0
dense (Dense)	(None, 256)	46022912
<pre>batch_normalization_2 (Batc hNormalization)</pre>	(None, 256)	1024
dropout_2 (Dropout)	(None, 256)	0
dense_1 (Dense)	(None, 1)	257
Total params: 46,034,241 Trainable params: 46.033.569		

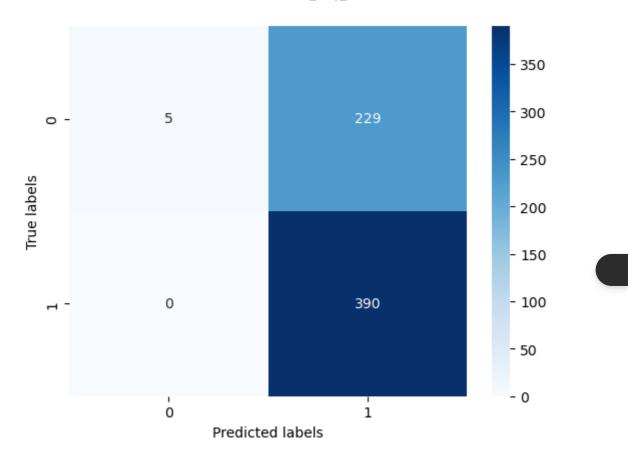
Total params: 46,034,241 Trainable params: 46,033,569 Non-trainable params: 672

In [22]: history=model.fit(x_train,y_train,epochs=10,validation_data=(x_validation_data,

```
Epoch 1/10
   cy: 0.9238 - val_loss: 0.3720 - val_accuracy: 0.8442
   Epoch 2/10
   cy: 0.9707 - val_loss: 0.1871 - val_accuracy: 0.9361
   Epoch 3/10
   cy: 0.9765 - val_loss: 0.1281 - val_accuracy: 0.9553
   Epoch 4/10
   cy: 0.9817 - val loss: 2.3034 - val accuracy: 0.6833
   Epoch 5/10
   cy: 0.9860 - val_loss: 11.1669 - val_accuracy: 0.7356
   Epoch 6/10
   cy: 0.9880 - val_loss: 0.2373 - val_accuracy: 0.9208
   Epoch 7/10
   cy: 0.9953 - val_loss: 0.3655 - val_accuracy: 0.8902
   Epoch 8/10
   cy: 0.9950 - val_loss: 561.4347 - val_accuracy: 0.7331
   Epoch 9/10
   cy: 0.9939 - val_loss: 0.3893 - val_accuracy: 0.8966
   Epoch 10/10
   cy: 0.9937 - val loss: 0.9791 - val accuracy: 0.8046
In [23]: # Saving the Model
    model.save('chest_xray_model.h5')
In [24]: model = load model('chest xray model.h5')
```

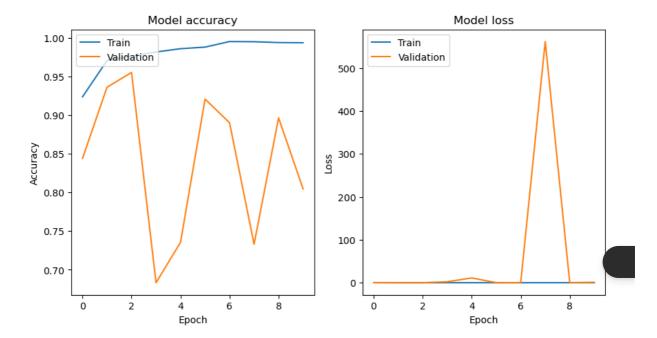
Confusion Matrix

```
In [25]: # Generate predictions on test data
        y_pred = model.predict(x_test)
        y_pred_binary = np.squeeze(np.round(y_pred))
        # Generate confusion matrix
        confusion_Matrix = confusion_matrix(y_test, y_pred_binary)
        confusion Matrix
       Out[25]: array([[ 5, 229],
               [ 0, 390]], dtype=int64)
In [26]: import matplotlib.pyplot as plt
        import seaborn as sns
        # Plot confusion matrix as heatmap
        sns.heatmap(confusion Matrix, annot=True, cmap='Blues',fmt='.3g')
        plt.xlabel('Predicted labels')
        plt.ylabel('True labels')
        plt.show()
```



graph between the training and validation accuracy and loss

```
In [27]: history
Out[27]: <keras.callbacks.History at 0x1c600262d90>
In [28]: fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(10, 5))
         # Plot the training and validation accuracy
         ax1.plot(history.history['accuracy'])
         ax1.plot(history.history['val_accuracy'])
         ax1.set_title('Model accuracy')
         ax1.set_ylabel('Accuracy')
         ax1.set xlabel('Epoch')
         ax1.legend(['Train', 'Validation'], loc='upper left')
         # Plot the training and validation loss
         ax2.plot(history.history['loss'])
         ax2.plot(history.history['val_loss'])
         ax2.set_title('Model loss')
         ax2.set_ylabel('Loss')
         ax2.set_xlabel('Epoch')
         ax2.legend(['Train', 'Validation'], loc='upper left')
         # Display the plots
         plt.show()
```



Classification Report

```
y_pred_binary = np.squeeze(np.round(y_pred))
 print(classification_report(y_test, y_pred_binary))
              precision
                            recall f1-score
                                                support
                              0.02
           0
                    1.00
                                         0.04
                                                    234
           1
                    0.63
                              1.00
                                         0.77
                                                    390
                                                    624
                                         0.63
    accuracy
   macro avg
                    0.82
                              0.51
                                         0.41
                                                    624
                                         0.50
weighted avg
                    0.77
                              0.63
                                                    624
```

AUC curve for binary classification

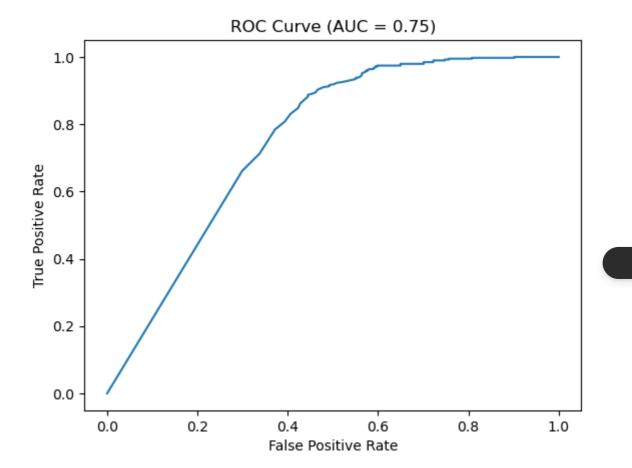
```
In [30]: from sklearn.metrics import roc_curve, roc_auc_score

# Get the predicted probabilities for the positive class
y_prob = model.predict(x_test)

# Compute the FPR, TPR, and thresholds for various classification thresholds
fpr, tpr, thresholds = roc_curve(y_test, y_prob)

# Compute the AUC score
auc_score = roc_auc_score(y_test, y_prob)

# Plot the ROC curve
import matplotlib.pyplot as plt
plt.plot(fpr, tpr)
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title(f'ROC Curve (AUC = {auc_score:.2f})')
plt.show()
```



Save the model

```
In [31]: model.save('chest_xray_pneumonia_model.h5')
```

Load Model and Test

```
In [32]:
        modelLoaded = load_model('chest_xray_pneumonia_model.h5')
In [33]:
         def predictImg():
             fig, axs = plt.subplots(2, 5, figsize=(10, 4))
             axs = axs.flatten()
             for i in range(10):
                 testimgnum = i+30
                  predicted = modelLoaded.predict(np.array([x_test[testimgnum]]))
                  predicted = int(np.round(predicted))
                  predictedClass = int(predicted >= 0.5)
                  actual = y_test[testimgnum]
                  axs[i].imshow(x test[testimgnum], cmap='gray')
                  axs[i].set_title(f'Predicted: {classnames[predictedClass]} \nActual: {cl
                  axs[i].axis('off')
             plt.tight_layout()
             plt.show()
In [34]:
         predictImg()
```

```
1/1 [=======] - 0s 203ms/step
1/1 [=======] - 0s 47ms/step
```

C:\Users\kirti\AppData\Local\Temp\ipykernel_6668\2954564863.py:8: DeprecationWarn ing: Conversion of an array with ndim > 0 to a scalar is deprecated, and will err or in future. Ensure you extract a single element from your array before performing this operation. (Deprecated NumPy 1.25.)

predicted = int(np.round(predicted))

Predicted: PNEUMONIA Predicted: PNEUMONIA Predicted: PNEUMONIA Predicted: PNEUMONIA Predicted: PNEUMONIA Actual: PNEUMONIA Actual: PNEUMONIA Actual: NORMAL Actual: NORMAL Actual: NORMAL











Predicted: PNEUMONIA Predicted: PNEUMONIA Predicted: PNEUMONIA Predicted: PNEUMONIA Predicted: PNEUMONIA Actual: PNEUMONIA











In []: