```
In [1]:
```

#Import into Colab

In [1]:

from google.colab import drive
drive.mount('/content/drive')

Drive already mounted at /content/drive; to attempt to forcibly remount, cal l drive.mount("/content/drive", force_remount=True).

In [2]:

#Imporing libraries

In [3]:

```
import matplotlib.pyplot as plt
import tensorflow as tf
from tensorflow import keras
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Dense, Activation, Conv2D, MaxPooling2D, Flatten, Dropo
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.callbacks import EarlyStopping
from tensorflow.keras.preprocessing.image import ImageDataGenerator
from sklearn.metrics import precision recall curve, roc curve, accuracy score, confusion ma
from sklearn.decomposition import PCA
from sklearn.model selection import train test split
import matplotlib.pyplot as plt
import seaborn as sns
plt.style.use('fivethirtyeight')
import pickle
import os
import numpy as np
import cv2
%matplotlib inline
```

In [3]:

```
#Labeling and sizing
```

```
In [4]:
```

In [4]:

```
#Training data
```

In [5]:

```
train = get_training_data('/content/drive/MyDrive/Colab Notebooks/Kaggle/chest_xray/train')
test = get_training_data('/content/drive/MyDrive/Colab Notebooks/Kaggle/chest_xray/test')
val = get_training_data('/content/drive/MyDrive/Colab Notebooks/Kaggle/chest_xray/val')
```

/usr/local/lib/python3.7/dist-packages/ipykernel_launcher.py:15: VisibleDepr ecationWarning: Creating an ndarray from ragged nested sequences (which is a list-or-tuple of lists-or-tuples-or ndarrays with different lengths or shape s) is deprecated. If you meant to do this, you must specify 'dtype=object' w hen creating the ndarray

from ipykernel import kernelapp as app

```
In [7]:
```

```
pnenumonia = 0
normal = 0

for i, j in train:
    if j == 0:
        pnenumonia+=1
    else:
        normal+=1
print('Pneumonia:', pnenumonia)
print('Normal:', normal)
print('Pneumonia - Normal:', pnenumonia-normal)
```

Pneumonia: 3875 Normal: 1341

Pneumonia - Normal: 2534

In [8]:

```
plt.imshow(train[1][0], cmap='gray')
plt.axis('off')
print(labels[train[1][1]])
```

PNEUMONIA



In [5]:

#Labelling feature

```
In [9]:
```

```
X = []
y = []
for feature, label in train:
    X.append(feature)
    y.append(label)
for feature, label in test:
    X.append(feature)
    y.append(label)
for feature, label in val:
    X.append(feature)
   y.append(label)
# resize data for deep learning
X = np.array(X).reshape(-1, img_size, img_size, 1)
y = np.array(y)
X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=32)
X train, X val, y train, y val = train test split(X train, y train, test size=0.20, random
```

In [10]:

```
X_train = X_train / 255
X_test = X_test / 255
X_val = X_val / 255
```

In [6]:

#ImageDataGenerator

In [11]:

In [7]:

```
#Optimizers for model
```

In [12]:

```
model = Sequential()
model.add(Conv2D(256, (3, 3), input shape=X train.shape[1:], padding='same'))
model.add(Activation('relu'))
model.add(MaxPooling2D(pool size=(2, 2), padding='same'))
model.add(BatchNormalization(axis=1))
model.add(Conv2D(64, (3, 3), padding='same'))
model.add(Activation('relu'))
model.add(MaxPooling2D(pool size=(2, 2), padding='same'))
model.add(BatchNormalization(axis=1))
model.add(Conv2D(16, (3, 3), padding='same'))
model.add(Activation('relu'))
model.add(MaxPooling2D(pool size=(2, 2), padding='same'))
model.add(BatchNormalization(axis=1))
model.add(Flatten()) # this converts our 3D feature maps to 1D feature vectors
model.add(Dropout(0.5))
model.add(Dense(64))
model.add(Activation('relu'))
model.add(Dropout(0.5))
model.add(Dense(1))
```

```
model.add(Activation('sigmoid'))
early_stop = EarlyStopping(patience=3, monitor='val_loss', restore_best_weights=True)
adam = Adam(learning_rate=0.0001)
model.compile(loss='binary_crossentropy',optimizer=adam,metrics=['acc'])
```

In [13]:

model.summary()

Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 200, 200, 256)	2560
activation (Activation)	(None, 200, 200, 256)	0
<pre>max_pooling2d (MaxPooling2D)</pre>	(None, 100, 100, 256)	0
<pre>batch_normalization (BatchN ormalization)</pre>	(None, 100, 100, 256)	400
conv2d_1 (Conv2D)	(None, 100, 100, 64)	147520
<pre>activation_1 (Activation)</pre>	(None, 100, 100, 64)	0
<pre>max_pooling2d_1 (MaxPooling 2D)</pre>	(None, 50, 50, 64)	0
<pre>batch_normalization_1 (Batc hNormalization)</pre>	(None, 50, 50, 64)	200

conv2d_2 (Conv2D)	(None, 50, 50, 16)	9232
<pre>activation_2 (Activation)</pre>	(None, 50, 50, 16)	0
<pre>max_pooling2d_2 (MaxPooling 2D)</pre>	(None, 25, 25, 16)	0
<pre>batch_normalization_2 (Batc hNormalization)</pre>	(None, 25, 25, 16)	100
flatten (Flatten)	(None, 10000)	0
dropout (Dropout)	(None, 10000)	0
dense (Dense)	(None, 64)	640064
activation_3 (Activation)	(None, 64)	0
dropout_1 (Dropout)	(None, 64)	0
dense_1 (Dense)	(None, 1)	65
activation_4 (Activation)	(None, 1)	0

Total params: 800,141 Trainable params: 799,791 Non-trainable params: 350

_

```
In [ ]:
```

```
y = model.fit(datagen.flow(X_train, y_train, batch_size=10), callbacks=[early_stop], valida
Epoch 1/15
acc: 0.7128 - val loss: 0.6822 - val acc: 0.7407
Epoch 2/15
375/375 [============ ] - 1735s 5s/step - loss: 0.4881 -
acc: 0.7427 - val loss: 0.4381 - val acc: 0.7449
Epoch 3/15
acc: 0.7966 - val loss: 0.4585 - val acc: 0.7150
Epoch 4/15
acc: 0.8217 - val loss: 0.4891 - val acc: 0.6894
Epoch 5/15
375/375 [============= ] - 1718s 5s/step - loss: 0.3396 -
acc: 0.8548 - val loss: 0.2462 - val acc: 0.9007
Epoch 6/15
acc: 0.8551 - val loss: 0.2492 - val_acc: 0.8986
Epoch 7/15
375/375 [============= ] - 1736s 5s/step - loss: 0.3127 -
acc: 0.8620 - val loss: 0.2851 - val_acc: 0.8762
Epoch 8/15
```

```
265/375 [==========>:....] - ETA: 7:57 - loss: 0.3056 - acc: 0.8751
```

In [8]:

#Evaluation

In [15]:

model.evaluate(X_test, y_test)

Out[15]:

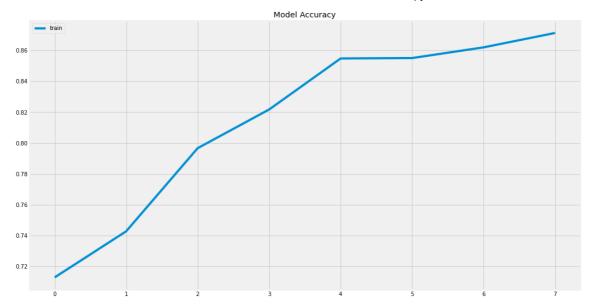
[0.24025382101535797, 0.9035836458206177]

In [9]:

#Graph showing accuracy and loss

In [16]:

```
plt.figure(figsize=(16, 9))
plt.plot(history.epoch, history.history['acc'])
plt.title('Model Accuracy')
plt.legend(['train'], loc='upper left')
plt.show()
plt.figure(figsize=(16, 9))
plt.plot(history.epoch, history.history['loss'])
plt.title('Model Loss')
plt.legend(['train'], loc='upper left')
plt.show()
plt.figure(figsize=(16, 9))
plt.plot(history.epoch, history.history['val acc'])
plt.title('Model Validation Accuracy')
plt.legend(['train'], loc='upper left')
plt.show()
plt.figure(figsize=(16, 9))
plt.plot(history.epoch, history.history['val_loss'])
plt.title('Model Validation Loss')
plt.legend(['train'], loc='upper left')
plt.show()
```

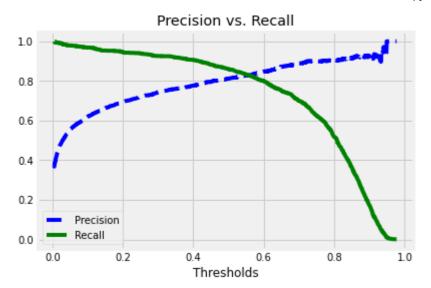


In [17]:

```
pred = model.predict(X_train)
precisions, recalls, thresholds = precision_recall_curve(y_train, pred)
fpr, tpr, thresholds2 = roc_curve(y_train, pred)
```

In [18]:

```
def plot precision recall(precisions, recalls, thresholds):
    plt.plot(thresholds, precisions[:-1], 'b--')
    plt.plot(thresholds, recalls[:-1], 'g-')
    plt.title('Precision vs. Recall')
    plt.xlabel('Thresholds')
    plt.legend(['Precision', 'Recall'], loc='best')
   plt.show()
def plot roc(fpr, tpr):
    plt.plot(fpr, tpr)
    plt.plot([0, 1], [0, 1], 'k--')
    plt.title('FPR (False Positive rate) vs TPR (True Positive Rate)')
    plt.xlabel('False Positive Rate')
    plt.ylabel('True Positive Rate (Recall)')
    plt.show()
plot_precision_recall(precisions, recalls, thresholds)
plot roc(fpr, tpr)
```





~

In [10]:

#Predictions

In [19]:

```
predictions = model.predict(X_test)
```

In [20]:

```
binary_predictions = []
threshold = thresholds[np.argmax(precisions >= 0.80)]
for i in predictions:
    if i >= threshold:
        binary_predictions.append(1)
    else:
        binary_predictions.append(0)
```

```
In [21]:
```

```
binary_predictions = []
threshold = thresholds[np.argmax(precisions >= 0.80)]
for i in predictions:
    if i >= threshold:
        binary_predictions.append(1)
    else:
        binary_predictions.append(0)
```

In [22]:

```
print('Accuracy on testing set:', accuracy_score(binary_predictions, y_test))
print('Precision on testing set:', precision_score(binary_predictions, y_test))
print('Recall on testing set:', recall_score(binary_predictions, y_test))
```

Accuracy on testing set: 0.9001706484641638 Precision on testing set: 0.8693009118541033 Recall on testing set: 0.794444444444444

In [11]:

```
#Confusion matrix
```

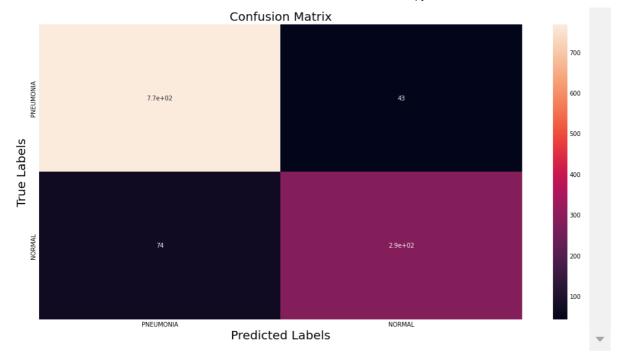
```
In [23]:
```

```
matrix = confusion_matrix(binary_predictions, y_test)
plt.figure(figsize=(16, 9))
ax= plt.subplot()
sns.heatmap(matrix, annot=True, ax = ax)

# labels, title and ticks
ax.set_xlabel('Predicted Labels', size=20)
ax.set_ylabel('True Labels', size=20)
ax.set_title('Confusion Matrix', size=20)
ax.xaxis.set_ticklabels(labels)
ax.yaxis.set_ticklabels(labels)
```

Out[23]:

```
[Text(0, 0.5, 'PNEUMONIA'), Text(0, 1.5, 'NORMAL')]
```



In [12]:

#Results

In [24]:

```
plt.figure(figsize=(10,10))
for i in range(25):
    plt.subplot(5,5,i+1)
    plt.xticks([])
    plt.yticks([])
    plt.grid(False)
    plt.imshow(X_train.reshape(-1, img_size, img_size)[i], cmap='gray')
    if(binary predictions[i]==y test[i]):
        plt.xlabel(labels[binary predictions[i]], color='blue')
    else:
        plt.xlabel(labels[binary_predictions[i]], color='red')
plt.show()
```



PNEUMONIA



NORMAL



PNEUMONIA



NORMAL



NORMAL



NORMAL



PNEUMONIA



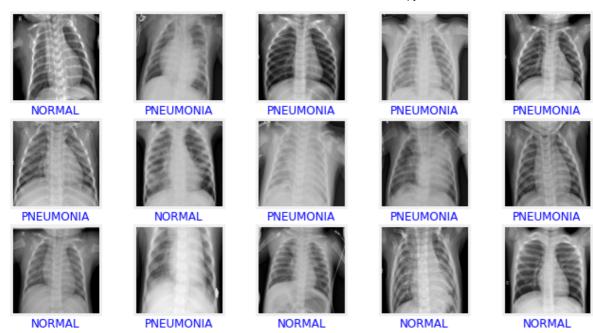
PNEUMONIA



PNEUMONIA



NORMAL



In []: