```
import os
import numpy as np
import pandas as pd
import random
import cv2
import matplotlib.pyplot as plt
%matplotlib inline
import keras as K
from keras.models import Model, Sequential
from keras.layers import Input, Dense, Flatten, Dropout, BatchNormalization
from keras.layers import Conv2D, SeparableConv2D, MaxPool2D, LeakyReLU, Activation
"from keras.optimizers import Adam"
from keras.preprocessing.image import ImageDataGenerator
from keras.callbacks import ModelCheckpoint, ReduceLROnPlateau, EarlyStopping
import tensorflow as tf
from mlxtend.plotting import plot confusion matrix
seed =232
np.random.seed(seed)
tf.random.set seed(seed)
from google.colab import drive
drive.mount('/content/drive')
     Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.
input_path = "/content/drive/MyDrive/AI-Project/chest_xray/"
fig, ax = plt.subplots(2,3, figsize=(15, 7))
ax = ax.ravel()
plt.tight_layout()
for i, _set in enumerate(['train', 'val', 'test']):
    set_path =input_path+_set
    ax[i].imshow(plt.imread(set path+'/NORMAL/'+os.listdir(set path+'/NORMAL')[0]),cmap='g
    ax[i].set title('Set: {}, Condition: Normal'.format( set))
    ax[i+3].imshow(plt.imread(set_path+'/PNEUMONIA/'+os.listdir(set_path+'/PNEUMONIA')[0])
    ax[i+3].set_title('Set: {}, Condition: Pneumonia'.format(_set))
```

Set: test, Condition: Normal

Set: val, Condition: Normal

Set: train, Condition: Normal

```
200
                                         200
        500
                                                                      400
                                         400
       1000
                                                                      600
                                         600
                                                                      800
                                         800
                                                                      1000
                                        1000
                                                                      1200
                                              200 400 600 800 1000 1200
Set: val, Condition: Pneumonia
                                                                              500 750 1000 1250 1500 1750
Set: test, Condition: Pneumonia
            500 1000 1500 2000 2500
Set: train, Condition: Pneumonia
                                         0
                                                                       100
       200
                                       200
                                                                       200
       400
                                       400
                                                                       400
       600
                                                                       500
for _set in ['train', 'val', 'test']:
    n_normal = len(os.listdir(input_path + _set + '/NORMAL'))
    n infect = len(os.listdir(input path + set + '/PNEUMONIA'))
    print('Set: {}, normal images: {}, pneumonia images: {}'.format(_set, n_normal, n_infe
     Set: train, normal images: 1341, pneumonia images: 3875
     Set: val, normal images: 8, pneumonia images: 8
     Set: test, normal images: 234, pneumonia images: 390
def process_data(img_dims, batch_size):
    # Data generation objects
    train_datagen = ImageDataGenerator(rescale=1./255, zoom_range=0.3, vertical_flip=True)
    test_val_datagen = ImageDataGenerator(rescale=1./255)
    # This is fed to the network in the specified batch sizes and image dimensions
    train gen = train datagen.flow from directory(
    directory=input path+'train',
    target_size=(img_dims, img_dims),
    batch_size=batch_size,
    class_mode='binary',
    shuffle=True)
    test gen = test val datagen.flow from directory(
    directory=input path+'test',
    target_size=(img_dims, img_dims),
    batch_size=batch_size,
    class mode='binary',
    shuffle=True)
    # I will be making predictions off of the test set in one batch size
    # This is useful to be able to get the confusion matrix
    test_data = []
    test_labels = []
    for cond in ['/NORMAL/', '/PNEUMONIA/']:
         for img in (os.listdir(input_path + 'test' + cond)):
             img = plt.imread(input_path+'test'+cond+img)
```

```
img = cv2.resize(img, (img_dims, img_dims))
            img = np.dstack([img, img, img])
            img = img.astype('float') / 255
            if cond=='/NORMAL/':
                label = 0;
            elif cond=='/PNEUMONIA/':
                label = 1;
            test_data.append(img)
            test_labels.append(label)
    test_data = np.array(test_data)
    test_labels = np.array(test_labels)
    return train_gen, test_gen, test_data, test_labels
img dims = 150
epochs = 10
batch size = 32
train_gen, test_gen, test_data, test_labels = process_data(img_dims, batch_size)
     Found 5216 images belonging to 2 classes.
     Found 624 images belonging to 2 classes.
inputs = Input(shape=(img_dims, img_dims, 3))
#First conv block
x = Conv2D(filters=16, kernel_size=(3,3), activation='relu', padding='same')(inputs)
x = Conv2D(filters=16, kernel_size=(3,3), activation='relu', padding='same')(x)
x = MaxPool2D(pool\_size=(2,2))(x)
#Second conv block
x = SeparableConv2D(filters=32, kernel_size=(3,3), activation='relu', padding='same')(x)
x = SeparableConv2D(filters=32, kernel_size=(3,3), activation='relu', padding='same')(x)
x = BatchNormalization()(x)
x = MaxPool2D(pool size=(2,2))(x)
#Third conv block
x = SeparableConv2D(filters=64, kernel size=(3,3), activation='relu', padding='same')(x)
x = SeparableConv2D(filters=64, kernel_size=(3,3), activation='relu', padding='same')(x)
x = BatchNormalization()(x)
x = MaxPool2D(pool\_size=(2,2))(x)
#Fourth conv block
x = SeparableConv2D(filters=128, kernel_size=(3,3), activation='relu', padding='same')(x)
x = SeparableConv2D(filters=128, kernel size=(3,3), activation='relu', padding='same')(x)
x = BatchNormalization()(x)
x = MaxPool2D(pool_size=(2,2))(x)
x = Dropout(rate=0.2)(x)
#Fifth conv block
x = SeparableConv2D(filters=256, kernel_size=(3,3), activation='relu', padding='same')(x)
x = SeparableConv2D(filters=256, kernel_size=(3,3), activation='relu', padding='same')(x)
x = BatchNormalization()(x)
```

```
x = MaxPool2D(pool\_size=(2,2))(x)
x = Dropout(rate=0.2)(x)
#FC layer
x = Flatten()(x)
x = Dense(units=512, activation='relu')(x)
x = Dropout(rate=0.7)(x)
x = Dense(units=128, activation='relu')(x)
x = Dropout(rate=0.5)(x)
x = Dense(units=64, activation='relu')(x)
x = Dropout(rate=0.3)(x)
#Output Layer
output = Dense(units = 1, activation='sigmoid')(x)
#Creating model and compiling
model = Model(inputs=inputs, outputs=output)
model.compile(optimizer='adam', loss='binary_crossentropy', metrics=['accuracy'])
#Callbacks
checkpoint = ModelCheckpoint(filepath='best_weights.hdf5', save_best_only=True, save_weight
lr_reduce = ReduceLROnPlateau(monitor='val_loss', factor=0.3, patience=2, verbose=2, mode=
early_stop = EarlyStopping(monitor='val_loss', min_delta=0.1, patience=1, mode='min')
```

Model: "model"

print(model.summary())

Layer (type)	Output Shape	Param #
input_2 (InputLayer)	[(None, 150, 150, 3)]	0
conv2d_2 (Conv2D)	(None, 150, 150, 16)	448
conv2d_3 (Conv2D)	(None, 150, 150, 16)	2320
<pre>max_pooling2d_5 (MaxPooling 2D)</pre>	(None, 75, 75, 16)	0
<pre>separable_conv2d_8 (Separab leConv2D)</pre>	(None, 75, 75, 32)	688
<pre>separable_conv2d_9 (Separab leConv2D)</pre>	(None, 75, 75, 32)	1344
<pre>batch_normalization_4 (Batc hNormalization)</pre>	(None, 75, 75, 32)	128
<pre>max_pooling2d_6 (MaxPooling 2D)</pre>	(None, 37, 37, 32)	0
<pre>separable_conv2d_10 (Separa bleConv2D)</pre>	(None, 37, 37, 64)	2400
<pre>separable_conv2d_11 (Separa bleConv2D)</pre>	(None, 37, 37, 64)	4736

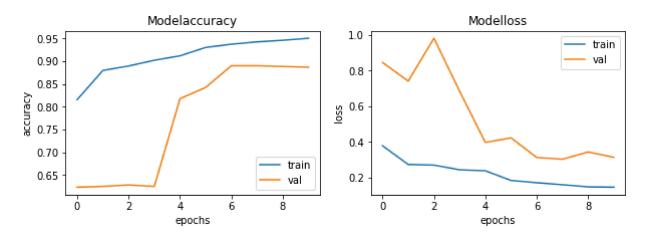
256

batch_normalization_5 (Batc (None, 37, 37, 64)

```
hNormalization)
   max_pooling2d_7 (MaxPooling (None, 18, 18, 64)
                                  a
   2D)
   separable_conv2d_12 (Separa (None, 18, 18, 128)
                                  8896
   bleConv2D)
   separable_conv2d_13 (Separa (None, 18, 18, 128)
                                  17664
   bleConv2D)
   batch_normalization_6 (Batc (None, 18, 18, 128)
                                  512
   hNormalization)
   max_pooling2d_8 (MaxPooling (None, 9, 9, 128)
                                  0
   2D)
   dropout 5 (Dropout)
                   (None, 9, 9, 128)
   separable conv2d 14 (Separa (None, 9, 9, 256)
                                  34176
   bleConv2D)
   separable conv2d 15 (Separa (None, 9, 9, 256)
                                  68096
   bleConv2D)
   L _L L
hist = model.fit_generator(
      train_gen, steps_per_epoch = train_gen.samples //batch_size,
      epochs=epochs, validation_data=test_gen,
      validation_steps=test_gen.samples //batch_size, callbacks=[checkpoint, lr_redu
  /usr/local/lib/python3.7/dist-packages/ipykernel launcher.py:4: UserWarning: `Model.
    after removing the cwd from sys.path.
  Epoch 1/10
  163/163 [========================== ] - 2710s 17s/step - loss: 0.3792 - accuracy:
  Epoch 2/10
  Epoch 3/10
  Epoch 4/10
  Epoch 5/10
  Epoch 00005: ReduceLROnPlateau reducing learning rate to 0.0003000000142492354.
  Epoch 6/10
  Epoch 7/10
  Epoch 00007: ReduceLROnPlateau reducing learning rate to 9.000000427477062e-05.
  Epoch 8/10
  Epoch 9/10
  Epoch 00009: ReduceLROnPlateau reducing learning rate to 2.700000040931627e-05.
```

```
Epoch 10/10
  fig, ax = plt.subplots(1,2, figsize=(10,3))
```

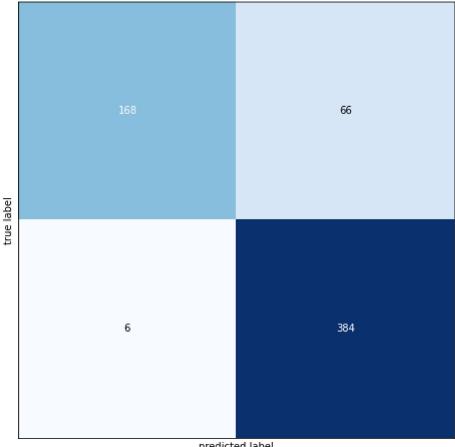
```
ax = ax.ravel()
for i, met in enumerate(['accuracy', 'loss']):
    ax[i].plot(hist.history[met])
    ax[i].plot(hist.history['val_' + met])
    ax[i].set_title('Model{}'.format(met))
    ax[i].set_xlabel('epochs')
    ax[i].set_ylabel(met)
    ax[i].legend(['train', 'val'])
```



from sklearn.metrics import accuracy_score, confusion_matrix

```
preds = model.predict(test_data)
accuracy = accuracy_score(test_labels, np.round(preds))*100
cm = confusion_matrix(test_labels, np.round(preds))
tn, fp, fn, tp = cm.ravel()
print('-----')
fig, ax = plot confusion matrix(conf mat=cm, figsize=(10,8), hide ticks=True, cmap=plt.cm.
plt.show()
print('\n-----')
print('Train accuracy: {}'.format(np.round((hist.history['accuracy'][-1])*100,2)))
print('\n-----')
precision = tp/(tp+fp)*100
recall = tp/(tp+fn)*100
print('Accuracy: {}%'.format(accuracy))
print('Precision: {}%'.format(precision))
print('Recall: {}%'.format(recall))
print('F1-score: {}'.format(2*precision*recall/(precision+recall)))
```

-----CONFUSION MATRIX-----



predicted label

-----TRAIN METRIC-----

Train accuracy: 95.0

-----TEST METRIC-----

Accuracy: 88.46153846153845% Precision: 85.3333333333334% Recall: 98.46153846153847% F1-score: 91.42857142857143

√ 7s completed at 5:23 PM