KMeans

Import necessary packages

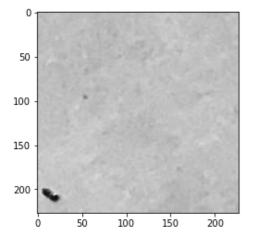
```
In [2]:
```

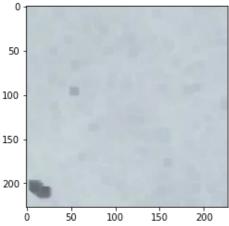
```
from sklearn.cluster import KMeans
import matplotlib.pyplot as plt
from matplotlib.image import imread
import numpy as np
import cv2
import utils
import os
from tqdm import tqdm
import PIL
from PIL import Image
```

View image and histogram

```
In [4]:
```

```
img_pos = cv2.imread("data/Negative/07146.jpg")
gray_pos = cv2.cvtColor(img_pos, cv2.COLOR_BGR2GRAY)
kernel = np.ones((7,7), np.uint8)
img_dilation = cv2.erode(img_pos, kernel, iterations=1)
plt.imshow(gray_pos, cmap="gray")
plt.show()
plt.imshow(img_dilation, cmap="gray")
plt.show()
```

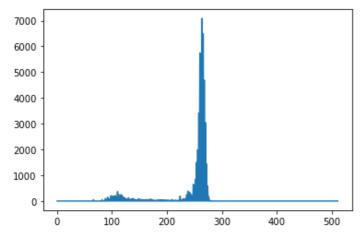




In [34]:

```
# img pos = cv2.imread("data/Positive/00005.jpg")
```

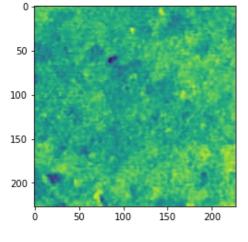
```
# gray_pos = cv2.cvtColor(img_pos, cv2.COLOR_BGR2GRAY)
# plt.imshow(gray_pos)
# plt.show()
hist_pos = cv2.calcHist([img_dilation],[0],None,[512],[0,256])
plt.plot(hist_pos)
plt.show()
```

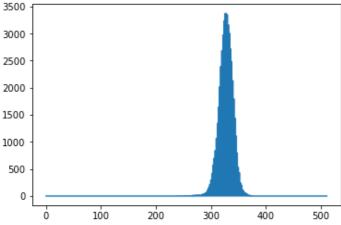


Convert to grayscale

In [22]:

```
img_neg = cv2.imread("data/Negative/00006.jpg")
gray_neg = cv2.cvtColor(img_neg, cv2.COLOR_BGR2GRAY)
plt.imshow(gray_neg)
plt.show()
hist_neg = cv2.calcHist([gray_neg],[0],None,[512],[0,256])
plt.plot(hist_neg)
plt.show()
```





Test OpenCV Kmeans

```
import numpy as np
import cv2
img = cv2.imread("data/Positive/00001.jpg")
Z = np.float32(img.reshape((-1,3)))
criteria = (cv2.TERM CRITERIA EPS + cv2.TERM CRITERIA MAX ITER, 10, 1.0)
K = 4
, labels, centers = cv2.kmeans(Z, K, None, criteria, 10, cv2.KMEANS RANDOM CENTERS)
labels = labels.reshape((img.shape[:-1]))
reduced = np.uint8(centers)[labels]
result = [np.hstack([img, reduced])]
for i, c in enumerate(centers):
   mask = cv2.inRange(labels, i, i)
   mask = np.dstack([mask]*3) # Make it 3 channel
   ex img = cv2.bitwise and(img, mask)
   ex reduced = cv2.bitwise and(reduced, mask)
   result.append(np.hstack([ex img, ex reduced]))
cv2.imwrite('out.jpg', np.vstack(result))
```

Out[2]:

True

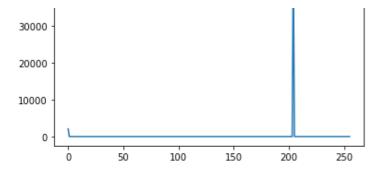
Plot histogram of KMeans result

In [2]:

```
import numpy as np
import cv2
img = cv2.imread("data/Positive/01051.jpg")
Z = np.float32(img.reshape((-1,3)))
criteria = (cv2.TERM CRITERIA EPS + cv2.TERM CRITERIA MAX ITER, 10, 1.0)
K = 2
, labels, centers = cv2.kmeans(Z, K, None, criteria, 10, cv2.KMEANS RANDOM CENTERS)
labels = labels.reshape((img.shape[:-1]))
reduced = np.uint8(centers)[labels]
result = [np.hstack([img, reduced])]
for i, c in enumerate(centers):
    if i ==1:
       mask = cv2.inRange(labels, i, i)
       mask = np.dstack([mask]*3) # Make it 3 channel
        ex img = cv2.bitwise and(img, mask)
       ex reduced = cv2.bitwise and(reduced, mask)
        result.append(np.hstack([ex img, ex_reduced]))
hist neg = cv2.calcHist([ex reduced],[0], None, [256], [0,256])
flat = ex reduced.flatten()
cnt crack=0
cnt=0
for x in range(0, len(flat), 3):
    # print(x, flat[x:x+3])
   if (flat[x:x+3] == [0,0,0]).all():
        cnt crack+=1
    else:
        cnt+=1
print(cnt crack, cnt)
plt.plot(hist neg)
plt.show()
cv2.imwrite('watermelon out.jpg', np.vstack(result))
```

2070 49459

50000



Out[2]:

True

Wrap to function

```
In [3]:
```

```
def test(img path):
    img = Image.open(img path)
    img = np.asarray(img)
    Z = np.float32(img.reshape((-1,3)))
    criteria = (cv2.TERM_CRITERIA_EPS + cv2.TERM_CRITERIA_MAX_ITER, 10, 1.0)
    , labels, centers = cv2.kmeans(Z, K, None, criteria, 10, cv2.KMEANS RANDOM CENTERS)
    labels = labels.reshape((img.shape[:-1]))
    reduced = np.uint8(centers)[labels]
    result = [np.hstack([img, reduced])]
    for i, c in enumerate(centers):
        if i==0:
            mask = cv2.inRange(labels, i, i)
            mask = np.dstack([mask]*3) # Make it 3 channel
            ex img = cv2.bitwise and(img, mask)
            ex reduced = cv2.bitwise and(reduced, mask)
            result.append(np.hstack([ex img, ex reduced]))
   hist neg = cv2.calcHist([ex reduced],[0],None,[256],[0,256])
    flat = ex reduced.flatten()
    cnt crack=0
    cnt=0
    for x in range(0, len(flat), 3):
        # print(x, flat[x:x+3])
        if (flat[x:x+3] == [0,0,0]).all():
            cnt crack+=1
        else:
            cnt+=1
    return [cnt crack/(cnt+cnt crack), 1- cnt crack/(cnt+cnt crack)]
```

Test with Positive images

```
In [4]:
```

```
import pandas as pd
fs = os.listdir("data/Positive/")
res_list = []
f_list =[]
for f in tqdm(fs, desc="fs"):
    res = test("data/Positive/"+f)
    res_list.append(res)
    f_list.append(f)
dic = {"res":res_list,"img":f_list}
df =pd.DataFrame(dic)
df.to_csv("positive_opencv_kmeans.csv")

fs: 100%| 20000/20000 [1:51:20<00:00, 2.99it/s]</pre>
```

Test with Negative images

```
import pandas as pd
fs = os.listdir("data/Negative/")
res_list = []
f_list =[]
for f in tqdm(fs, desc="fs"):
    res = test("data/Negative/"+f)
    res_list.append(res)
    f_list.append(f)
dic = {"res":res_list,"img":f_list}
df =pd.DataFrame(dic)
df.to_csv("negative_opencv_kmeans.csv")
```

Try KMeans with sklearn and kmeans++

```
In [ ]:
```

```
from sklearn.cluster import KMeans
img = cv2.imread("data/Negative/16471.jpg")
img=cv2.cvtColor(img,cv2.COLOR BGR2HSV)
# print(img)
img=img.reshape((img.shape[1]*img.shape[0],3))
kmeans=KMeans(n clusters=2,init="k-means++")
s=kmeans.fit(img)
centroid=kmeans.cluster centers
print(centroid)
labels=list(kmeans.labels )
percent=[]
for i in range(len(centroid)):
  j=labels.count(i)
  j=j/(len(labels))
  percent.append(j)
print(percent)
plt.pie(percent,colors=np.array(centroid/255),labels=np.arange(len(centroid)))
plt.show()
```

Wrap to function

```
In [ ]:
```

```
def test_sklearn(img_path):
    img = cv2.imread(img_path)
    img=cv2.cvtColor(img,cv2.COLOR_BGR2RGB)
    img=img.reshape((img.shape[1]*img.shape[0],3))
    kmeans=KMeans(n_clusters=2,init="k-means++")
    s=kmeans.fit(img)
    centroid=kmeans.cluster_centers_
    labels=list(kmeans.labels_)
    percent=[]
    for i in range(len(centroid)):
        j=labels.count(i)
        j=j/(len(labels))
        percent.append(j)
    return max(percent)
```

Test with Positive images and save result to csv file since it takes long time for each run (20k images)

```
In [ ]:
```

```
import pandas as pd
fs = os.listdir("data/Positive/")
res_list = []
f_list =[]
for f in tqdm(fs, desc="fs"):
    res = test_sklearn("data/Positive/"+f)
    res_list.append(res)
    f_list.append(f)
dic = {"res":res_list,"img":f_list}
```

```
df =pd.DataFrame(dic)
df.to_csv("positive.csv")
```

Test with Negative images and save result to csv file since it takes long time for each run (20k images)

```
In [ ]:
```

```
import pandas as pd
fs = os.listdir("data/Negative/")
res_list = []
f_list =[]
for f in tqdm(fs, desc="fs"):
    res = test_sklearn("data/Negative/"+f)
    res_list.append(res)
    f_list.append(f)
dic = {"res":res_list,"img":f_list}
df =pd.DataFrame(dic)
df.to_csv("negative.csv")
```

Evaluate

```
In [ ]:
```

```
import pandas as pd
import numpy as np
from sklearn.metrics import classification report, confusion matrix
threshold=0.75
df_pos = pd.read_csv('positive_opencv_kmeans.csv')
res pos = df pos.iloc[:,1]
# print(res.tolist())
cnt=0
positive list = []
positive label = np.ones((20000,)).astype(int)
for r in res pos.tolist():
   # print(type(r))
   s = r[1:10]
    t=float(s)
    if t>=threshold:
       positive_list.append(1)
    else:
        positive list.append(0)
        cnt+=1
postive pred= np.array(positive list)
df neg = pd.read csv('negative opencv kmeans.csv')
res_neg = df_neg.iloc[:,1]
cnt neg=0
negative list=[]
negative label=np.zeros((20000,)).astype(int)
label = np.concatenate((positive label, negative label))
for r in res neg.tolist():
   s = r[1:10]
   t=float(s)
   if t<threshold:</pre>
       negative list.append(0)
    else:
        negative_list.append(1)
negative pred= np.array(negative list)
pred = np.concatenate((postive pred, negative pred))
print(classification report(label, pred, target names = ['Positive','Negative']))
print(confusion matrix(label, pred))
```

KMeans++

```
In [ ]:
```

```
from sklearn.cluster import KMeans
```

```
import matplotlib.pyplot as plt
from matplotlib.image import imread
import numpy as np
import cv2
import utils
import os
from tqdm import tqdm
import PIL
from PIL import Image
```

```
In []:
img_pos = cv2.imread("data/Negative/10465.jpg")

gray_pos = cv2.cvtColor(img_pos, cv2.COLOR_BGR2GRAY)
plt.imshow(gray_pos, cmap="gray")
plt.show()
hist_pos = cv2.calcHist([gray_pos],[0],None,[512],[0,256])
plt.plot(hist_pos)
plt.show()
```

Try KMeans with sklearn and kmeans++

```
In [ ]:
```

```
from sklearn.cluster import KMeans
img = cv2.imread("data/Negative/16471.jpg")
img=cv2.cvtColor(img,cv2.COLOR BGR2HSV)
# print(img)
img=img.reshape((img.shape[1]*img.shape[0],3))
kmeans=KMeans(n clusters=2,init="k-means++")
s=kmeans.fit(img)
centroid=kmeans.cluster_centers_
print(centroid)
labels=list(kmeans.labels )
percent=[]
for i in range(len(centroid)):
 j=labels.count(i)
 j=j/(len(labels))
 percent.append(j)
print(percent)
plt.pie(percent,colors=np.array(centroid/255),labels=np.arange(len(centroid)))
plt.show()
```

Wrap to function

```
In []:

def test_sklearn(img_path):
    img = cv2.imread(img_path)
    img=cv2.cvtColor(img,cv2.COLOR_BGR2RGB)
    img=img.reshape((img.shape[1]*img.shape[0],3))
    kmeans=KMeans(n_clusters=2,init="k-means++")
    s=kmeans.fit(img)
    centroid=kmeans.cluster_centers_
    labels=list(kmeans.labels_)
    percent=[]
    for i in range(len(centroid)):
        j=labels.count(i)
        j=j/(len(labels))
        percent.append(j)
    return max(percent)
```

Test with Positive images and save result to csv file since it takes long time for each run (20k images)

```
In []:
import pandas as pd
fs = os.listdir("data/Positive/")
```

```
res_list = []
f_list =[]
for f in tqdm(fs, desc="fs"):
    res = test_sklearn("data/Positive/"+f)
    res_list.append(res)
    f_list.append(f)
dic = {"res":res_list,"img":f_list}
df =pd.DataFrame(dic)
df.to_csv("positive.csv")
```

Test with Negative images and save result to csv file since it takes long time for each run (20k images)

```
import pandas as pd
fs = os.listdir("data/Negative/")
res_list = []
f_list =[]
for f in tqdm(fs, desc="fs"):
    res = test_sklearn("data/Negative/"+f)
    res_list.append(res)
    f_list.append(f)
dic = {"res":res_list,"img":f_list}
df =pd.DataFrame(dic)
df.to_csv("negative.csv")
```

Evaluate

```
In [ ]:
```

```
import pandas as pd
import numpy as np
from sklearn.metrics import classification report, confusion matrix
threshold=0.8
df pos = pd.read csv('positive.csv')
res pos = df pos.iloc[:,1]
# print(res.tolist())
cnt=0
positive_list = []
positive label = np.ones((20000,)).astype(int)
for r in res pos.tolist():
    if r>=threshold:
       positive_list.append(1)
    else:
       positive_list.append(0)
       cnt+=1
postive pred= np.array(positive list)
df_neg = pd.read csv('negative.csv')
res neg = df neg.iloc[:,1]
cnt neg=0
negative list=[]
negative label=np.zeros((20000,)).astype(int)
label = np.concatenate((positive label, negative label))
for r in res neg.tolist():
    if r<threshold:</pre>
       negative_list.append(0)
    else:
       negative list.append(1)
negative pred= np.array(negative list)
pred = np.concatenate((postive pred, negative pred))
print(classification_report(label, pred, target_names = ['Positive','Negative']))
print(confusion matrix(label, pred))
```

CNN

In []:

```
import matplotlib.pyplot as plt
import seaborn as sns
import keras
from keras.models import Sequential
from keras.layers import Dense, Conv2D , MaxPool2D , Flatten , Dropout
from keras.preprocessing.image import ImageDataGenerator
from keras.optimizers import Adam, RMSprop, Adagrad
from keras.layers import BatchNormalization
from sklearn.metrics import classification report, confusion matrix
# %tensorflow version 1.x
import tensorflow as tf
import cv2
import os
import numpy as np
import warnings
warnings.filterwarnings('ignore')
print(tf.__version__)
```

With generator

Split train test

```
import shutil
cnt=0
for f in os.listdir("/content/data/train/Positive"):
    if cnt<=1000:
        shutil.move("/content/data/train/Positive/"+f, "/content/data/test/Positive/"+f)
        cnt+=1
cnt=0
for f in os.listdir("/content/data/train/Negative"):
    if cnt<=1000:
        shutil.move("/content/data/train/Negative/"+f, "/content/data/test/Negative/"+f)
        cnt+=1</pre>
```

Create CNN

```
model = Sequential() model.add(Conv2D(64,3,padding="same", activation="relu", input_shape = (120,120,1))) model.add(MaxPool2D())
```

model.add(Conv2D(64, 3, padding="same", activation="relu")) model.add(MaxPool2D())

model.add(Conv2D(128, 3, padding="same", activation="relu")) model.add(MaxPool2D())

model.add(Flatten()) model.add(Dense(256,activation="relu")) model.add(Dropout(0.5)) model.add(BatchNormalization()) model.add(Dense(2, activation="softmax"))

model.summary()

Create image generator for train and valid

```
In [ ]:
```

```
from keras.preprocessing.image import ImageDataGenerator
train_datagen = ImageDataGenerator(
    rescale=1./255,
    shear_range=0.2,
    width_shift_range=0.2,
    height_shift_range=0.2,
    zoom_range=0.2,
    rotation_range=40,
    horizontal_flip=True)

test_datagen = ImageDataGenerator(rescale=1./255)
```

Load images to train and validation generator

In []: batch size = 128 train generator = train datagen.flow from directory('/content/data/train', target size=(120,120), batch size=batch size, color mode="grayscale", class mode='binary') # since we use binary crossentropy loss, we need binary lab els validation generator = test datagen.flow from directory('/content/data/valid', target size=(120,120), batch size=batch size, color mode="grayscale", class mode='binary') # import pandas as pd # pd.get dummies(pd.Series(validation generator.classes)) print(validation_generator.class indices)

Choose parameters, compiling model and train

```
In [ ]:
opt = Adam(lr=0.0005)
model.compile(loss="sparse categorical crossentropy", optimizer=opt, metrics=["accuracy"]
checkpoint path = "/content/drive/MyDrive/CVcourse/training/cp.ckpt"
# checkpoint dir = os.path.dirname(checkpoint path)
# Create a callback that saves the model's weights
model checkpoint callback = keras.callbacks.ModelCheckpoint(
    filepath=checkpoint path,
    save weights only=True,
   monitor='val accuracy',
   mode='max',
    save best only=True)
history = model.fit_generator(train_generator,
                              steps per epoch=len(train generator) // batch size,
                              validation data=validation generator,
                              verbose=1,
                              epochs = 200,
                              callbacks=[model checkpoint callback],
                              validation steps=len(validation generator) // batch size
```

Plot accurcay and loss graph

```
In [ ]:
```

```
plt.figure(figsize=(12, 12))
plt.style.use('ggplot')
plt.subplot(2,2,1)
plt.plot(history.history['accuracy'])
plt.plot(history.history['val_accuracy'])
plt.title('Accuracy of the Model')
plt.ylabel('Accuracy', fontsize=12)
plt.xlabel('Epoch', fontsize=12)
plt.legend(['train accuracy', 'validation accuracy'], loc='lower right', prop={'size': 1
2})
plt.subplot(2,2,2)
plt.plot(history.history['loss'])
plt.plot(history.history['val_loss'])
plt.title('Loss of the Model')
```

```
plt.ylabel('Loss', fontsize=12)
plt.xlabel('Epoch', fontsize=12)
plt.legend(['train loss', 'validation loss'], loc='best', prop={'size': 12})
```

Create image generator for test and evaluate model

print(confusion_matrix(test_generator.classes, predictions))

Without generator

ve','Positive']))

Load image

```
In [ ]:
labels = ['Negative', 'Positive']
img size = 120
def read images(data dir):
    data = []
    for label in labels:
        path = os.path.join(data dir, label)
        class num = labels.index(label)
        for img in os.listdir(path):
            try:
                img arr = cv2.imread(os.path.join(path, img), cv2.IMREAD GRAYSCALE)
                resized arr = cv2.resize(img arr, (img size, img size))
                data.append([resized arr, class num])
            except Exception as e:
                print(e)
    return np.array(data)
Dataset origin = read images("/content/data/train")
Dataset valid = read images("/content/data/valid")
Dataset test = read images("/content/data/test")
```

Images data to array x contains feature, y contains label

```
x_val = []
y_val = []

for feature, label in Dataset_valid:
    x_val.append(feature)
    y_val.append(label)

x_val = np.array(x_val).reshape(-1, img_size, img_size, 1)
x_val = x_val / 255
y_val = np.array(y_val)
```

Visualize data distribution

```
In []:

Im = []
for i in Dataset_origin:
    if(i[1] == 0):
        Im.append("Negative")
    elif(i[1] == 1):
        Im.append("Positive")

plt.figure(figsize=(10, 10))
plt.subplot(2, 2, 1)
sns.set_style('darkgrid')
axl = sns.countplot(Im)
axl.set_title("Number of Images")
```

Creating CNN

```
In []:
model_origin = Sequential()
model_origin.add(Conv2D(64,3,padding="same", activation="relu", input_shape = x.shape[1:
]))
model_origin.add(MaxPool2D())

model_origin.add(Conv2D(64, 3, padding="same", activation="relu"))
model_origin.add(MaxPool2D())

model_origin.add(Conv2D(128, 3, padding="same", activation="relu"))
model_origin.add(MaxPool2D())

model_origin.add(Flatten())
model_origin.add(Dense(256,activation="relu"))
model_origin.add(Dropout(0.5))
model_origin.add(BatchNormalization())
model_origin.add(Dense(2, activation="softmax"))

model_origin.summary()
```

Setting parameter, compile and train

```
In []:

opt = Adam(lr=1e-5)

model_origin.compile(loss="sparse_categorical_crossentropy", optimizer=opt, metrics=["acc uracy"])

history_origin = model_origin.fit(x, y, validation_data=(x_val, y_val), epochs = 15, bat ch_size = 128, verbose=1)
```

PLot accuracy and loss graph

```
In []:
plt.figure(figsize=(12, 12))
```

```
plt.style.use('ggplot')
plt.subplot(2,2,1)
plt.plot(history.history['accuracy'])
plt.plot(history.history['val accuracy'])
plt.title('Accuracy of the Model')
plt.ylabel('Accuracy', fontsize=12)
plt.xlabel('Epoch', fontsize=12)
plt.legend(['train accuracy', 'validation accuracy'], loc='lower right', prop={'size': 1
2 } )
plt.subplot(2,2,2)
plt.plot(history.history['loss'])
plt.plot(history.history['val loss'])
plt.title('Loss of the Model')
plt.ylabel('Loss', fontsize=12)
plt.xlabel('Epoch', fontsize=12)
plt.legend(['train loss', 'validation loss'], loc='best', prop={'size': 12})
```

Load test images

```
In []:

x_test = []
y_test = []

for feature, label in Dataset_test:
    x_test.append(feature)
    y_test.append(label)

x_test = np.array(x_test).reshape(-1, img_size, img_size, 1)
x_test = x_test / 255
y_test = np.array(y_test)
```

Evaluate model

```
In []:

predictions = model_origin.predict_classes(x_test)
predictions = predictions.reshape(1,-1)[0]
print(classification_report(y_test, predictions, target_names = ['Negative','Positive'])
)
print(confusion_matrix(y_test, predictions))
```

VGG-16

```
In [ ]:
```

```
from keras.preprocessing.image import ImageDataGenerator
from tensorflow.keras.optimizers import Adam
from tensorflow.keras import Model, layers
import tensorflow as tf
import keras

import matplotlib.pyplot as plt
from matplotlib.image import imread
import cv2
```

Exclude test set from the original dataset

```
import os
import shutil
cnt=0
for f in os.listdir("/content/data/train/Positive"):
   if cnt<=1000:
      shutil.move("/content/data/train/Positive/"+f,"/content/data/test/Positive/"+f)</pre>
```

```
cnt+=1
cnt=0

for f in os.listdir("/content/data/train/Negative"):
   if cnt<=1000:
      shutil.move("/content/data/train/Negative/"+f,"/content/data/test/Negative/"+f)
      cnt+=1</pre>
```

Preparing the Dataset

```
In []:

def load_images_from_folder(folder):
    images = []
    for filename in os.listdir(folder):
        img = cv2.imread(os.path.join(folder,filename))
        if img is not None:
            images.append(img)
        if len(images) == 4:
            break

fig=plt.figure(figsize=(10,12))

for img,x in zip(images,range(1, 5)):
        ax=fig.add_subplot(1, 4, x)
        ax.imshow(img)
        ax.set_title(img.shape)
```

Load pretrained VGG-16 model

```
In [ ]:
```

Add fully connected layers to the network:

- Flatten
- Dense layer with 256 node and relu activation
- Drop out with the rate of 0.4
- · softmax layer with 2 node

```
In [ ]:
```

```
last_layer = pretrained_model.get_layer('block5_pool')
last_output = last_layer.output

x = layers.Flatten()(last_output)
x = layers.Dense(256, activation='relu')(x)
x = layers.Dropout(0.4)(x)
x = layers.Dense(2, activation='softmax')(x)

model_vgg = Model(pretrained_model.input, x)

model_vgg.summary()
model_vgg.compile(optimizer = Adam(lr=0.0005),
```

```
loss = 'sparse_categorical_crossentropy',
metrics = ['accuracy'])
```

Split dataset to trainning and validation subset

```
In [ ]:
```

Train model with 10 epochs

```
In [ ]:
```

Plot the trainning experiments

```
In [ ]:
```

```
acc = history.history['accuracy']
val_acc = history.history['val_accuracy']
loss = history.history['loss']
val_loss = history.history['val_loss']
epochs = range(len(acc))

plt.plot(epochs, acc, 'r', label='Training accuracy')
plt.plot(epochs, val_acc, 'b', label='Validation accuracy')
plt.title('Training and validation accuracy')

plt.figure()

plt.plot(epochs, loss, 'r', label='Training Loss')
plt.plot(epochs, val_loss, 'b', label='Validation Loss')
plt.title('Training and validation loss')

plt.legend()
plt.show()
```

Testing and evaluate the model

```
In [ ]:
```

```
shuffle = False,
    class_mode='binary',
    batch_size=1)

filenames = test_generator.filenames
nb_samples = len(filenames)

predict = model_vgg.predict_generator(test_generator, steps = nb_samples)
predictions = np.argmax(predict,axis=1)
print(classification_report(test_generator.classes, predictions, target_names = ['Negative', 'Positive']))
print(confusion_matrix(test_generator.classes, predictions))
```