```
# command to download the dataset (covid 19) the specified URL
!wget http://cb.lk/covid_19
# unzipping the dataset : covid_19
!unzip covid 19
# import libraries
import numpy as np
import matplotlib.pyplot as plt
import keras
from keras.layers import *
from keras.models import *
from keras.preprocessing import image
# CNN based model using keras (with tensorflow backend)
classifier = Sequential() # we have created an object(classifier:modle name) of class sequ
## we are using conv2D layer
## we are not using VGG layer because it works on around 180 million features so our model
classifier.add(Conv2D(32,kernel_size=(3,3),activation = 'relu',input_shape = (224,224,3)))
classifier.add(Conv2D(64,(3,3),activation = 'relu'))
# using 2 convolution layers of kernel size =(3,3) is similar to using one single convolut
# but we don't perfer that becauser to increase non linearity in the model using more relu
classifier.add(MaxPooling2D(pool_size = (2,2)))
classifier.add(Dropout(0.25)) # to find overfitting in the model
classifier.add(Conv2D(64,(3,3),activation = 'relu'))
classifier.add(MaxPooling2D(pool size = (2,2)))
classifier.add(Dropout(0.25))
classifier.add(Conv2D(128,(3,3),activation = 'relu'))
# as we go deep into model we are increasing the number of convolutional layer because we
# features (i.e complex features)
classifier.add(MaxPooling2D(pool_size = (2,2)))
classifier.add(Dropout(0.25))
classifier.add(Flatten());
classifier.add(Dense(64,activation = 'relu'))
classifier.add(Dropout(0.5))
classifier.add(Dense(1,activation = 'sigmoid'))
classifier.compile(loss = keras.losses.binary_crossentropy,optimizer = 'adam',metrics =['&
# adam use gradient descent algorithm for optimizing
classifier.summary()
     Model: "sequential"
```

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 222, 222, 32)	896
conv2d_1 (Conv2D)	(None, 220, 220, 64)	18496
max_pooling2d (MaxPooling2D)	(None, 110, 110, 64)	0
dropout (Dropout)	(None, 110, 110, 64)	0
conv2d_2 (Conv2D)	(None, 108, 108, 64)	36928
max_pooling2d_1 (MaxPooling2	(None, 54, 54, 64)	0
dropout_1 (Dropout)	(None, 54, 54, 64)	0
conv2d_3 (Conv2D)	(None, 52, 52, 128)	73856
max_pooling2d_2 (MaxPooling2	(None, 26, 26, 128)	0
dropout_2 (Dropout)	(None, 26, 26, 128)	0
flatten (Flatten)	(None, 86528)	0
dense (Dense)	(None, 64)	5537856
dropout_3 (Dropout)	(None, 64)	0
dense_1 (Dense)	(None, 1)	65
Total params: 5,668,097		

## training the model
# firstly setting up the data ready for training i.e data preprocessing with the help of I
train\_datagen = image.ImageDataGenerator(
 rescale = 1./255,
 shear\_range = 0.2,

snear\_range = 0.2,
zoom\_range = 0.2,
horizontal\_flip = True

Trainable params: 5,668,097 Non-trainable params: 0

)
validation\_datagen = image.ImageDataGenerator(rescale = 1./255)
# we are dividing by 255 for normalisation

# here images get loaded one by one and get reshaped to specified dimensions

Found 224 images belonging to 2 classes.

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train_data.class_indices
```

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{'Covid': 0, 'Normal': 1}
validation_data = validation_datagen.flow_from_directory('CovidDataset/Val',
                                      target_size = (224,224),
                                      batch_size = 32,
                                    class mode = 'binary')
# here images get loaded one by one and get reshaped to specified dimensions
    Found 60 images belonging to 2 classes.
validation data.class indices
    {'Covid': 0, 'Normal': 1}
final_model = classifier.fit_generator(train_data,
                     steps_per_epoch= 7,
                    epochs = 10,
                    validation_data = validation_data,
                    validation_steps = 2)
    /usr/local/lib/python3.7/dist-packages/keras/engine/training.py:1915: UserWarning: `
     warnings.warn('`Model.fit_generator` is deprecated and '
    Epoch 1/10
    7/7 [============== ] - 60s 2s/step - loss: 2.4544 - accuracy: 0.5109
    Epoch 2/10
    Epoch 3/10
    7/7 [============ ] - 10s 1s/step - loss: 0.6194 - accuracy: 0.6369
    Epoch 4/10
    7/7 [============== ] - 10s 1s/step - loss: 0.4704 - accuracy: 0.7643
    Epoch 5/10
    Epoch 6/10
    7/7 [============ ] - 10s 1s/step - loss: 0.2847 - accuracy: 0.8927
    Epoch 7/10
    7/7 [============= ] - 10s 2s/step - loss: 0.3579 - accuracy: 0.8288
    Epoch 8/10
    Epoch 9/10
    7/7 [============ ] - 10s 1s/step - loss: 0.1903 - accuracy: 0.9034
    Epoch 10/10
    7/7 [============ ] - 10s 1s/step - loss: 0.1394 - accuracy: 0.9562
## with help of "grad CAM" technique we can see and visualize how our model is differentia
## part of network our model is focusing on cilensing map
## can read about class activation
## saving the model
classifier.save("COVID 19 model.h5")
##Through Keras, models can be saved in three formats:
#YAML format
#JSON format
#HDF5 format
```

```
#YAML and JSON files store only model structure, whereas, HDF5 file stores complete neural
# we can load the model when ever required
classifier = load model("COVID 19 model.h5")
import os # it is a standard library in python
# setup for confusion matrix
y_validation = []
y_predict = []
for i in os.listdir("./CovidDataset/Val/Normal/"):
  img = image.load_img("./CovidDataset/Val/Normal/"+i,target_size = (224,224))
  img = image.img_to_array(img)
  img = np.expand_dims(img,axis = 0)
 pred = classifier.predict_classes(img)
 y_predict.append(pred[0,0])
 y validation.append(1)
for i in os.listdir("./CovidDataset/Val/Covid/"):
  img = image.load_img("./CovidDataset/Val/Covid/"+i,target_size = (224,224))
  img = image.img_to_array(img)
  img = np.expand_dims(img,axis = 0)
  pred = classifier.predict_classes(img)
 y_predict.append(pred[0,0])
 y_validation.append(0)
y_validation = np.array(y_validation)
y_predict = np.array(y_predict)
     /usr/local/lib/python3.7/dist-packages/keras/engine/sequential.py:450: UserWarning:
       warnings.warn('`model.predict_classes()` is deprecated and '
# now importing sklearn for confusion matrix
from sklearn.metrics import confusion_matrix
# sklearn provides a selection of efficient tools for machine learning and statistical moc
#including classification, regression, clustering and dimensionality reduction via a consi
con matrix = confusion matrix(y validation,y predict)
import seaborn as sns
#Seaborn is an open-source Python library built on top of matplotlib.
#It is used for data visualization and exploratory data analysis.
#Seaborn works easily with dataframes and the Pandas library.
sns.heatmap(con_matrix,cmap = "ocean_r",annot = True)
# Heatmap is defined as a graphical representation of data using colors to visualize the 	imes
# cmap is for color map (ex : ocean, ocean_r, pink, pink_r, plasma )
# annot : When we pass bool 'True' value to annot then the value will show on each cell of
```

<matplotlib.axes.\_subplots.AxesSubplot at 0x7fa815153890>



