

COVID-19 DETECTION FROM X-RAY IMAGES USING CONVOLUTIONAL NEURAL NETWORK (CNN)



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Project Source



Source: <https://www.kaggle.com/technick/covid-detection-from-xray>

Introduction



- The need for auxiliary diagnostic tools has increased as there is lack of automated toolkits to detect COVID-19.
- Researchers state that combining clinical image features with laboratory results may help in early detection of COVID-19 [1,2,3].
- Recent findings obtained using radiology imaging techniques suggest that such images contain salient information about the COVID-19 virus [4].
- In this project, a Convolutional Neural Network (CNN) is designed to automatically diagnose COVID-19 using x-ray images

Libraries



Kera's

- High-level neural networks library

TensorFlow

- High performance numerical computation based library for ML applications

Numpy

- Fundamental scientific computations

Pandas

- Data Analysis:** pandas is a fast, powerful, flexible and easy to use open source data analysis and manipulation tool

Matplotlib

- Data visualization

```
from keras import backend as K
from keras.preprocessing.image import ImageDataGenerator, load_img, img_to_array
from keras.models import Sequential, Model
from keras.layers import Conv2D, MaxPooling2D, GlobalAveragePooling2D
from keras.layers import Activation, Dropout, BatchNormalization, Flatten, Dense, AvgPool2D, MaxPool2D
from keras.models import Sequential, Model
from keras.applications.vgg16 import VGG16, preprocess_input
from keras.optimizers import Adam, SGD, RMSprop

import tensorflow as tf

import os
import numpy as np
import pandas as pd

import matplotlib.pyplot as plt
%matplotlib inline
```

Data Set



Data Set Overview

No. of X-ray Images = 1098

There are two classes:

- 1) Covid (751)
- 2) Normal (347)

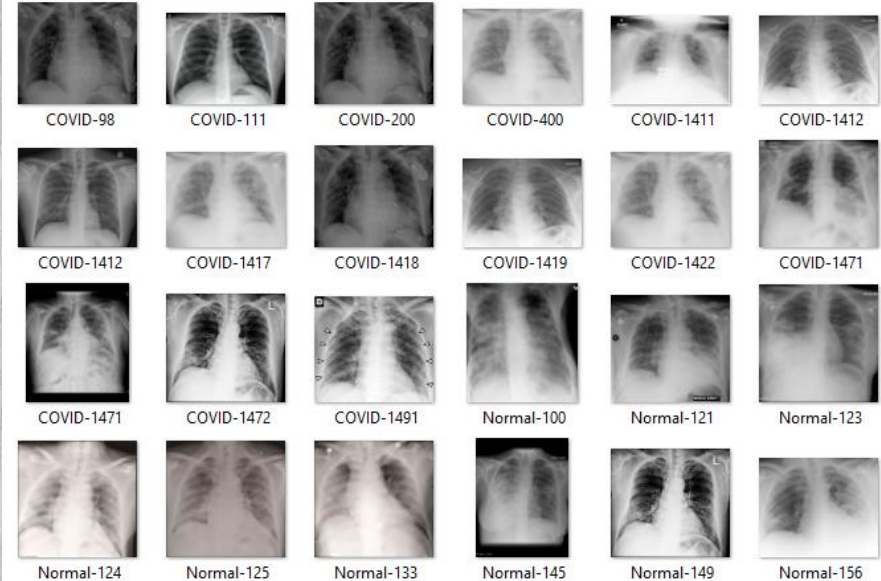
Dataset containing X-ray Images from two classes ('covid', 'normal') is loaded into "DATASET_DIR"

Images from both the classes are placed in their respective folders

```
DATASET_DIR = "../input/covid-19-x-ray-10000-images/dataset"
```

```
os.listdir(DATASET_DIR)
```

```
['covid', 'normal']
```



Data Analysis



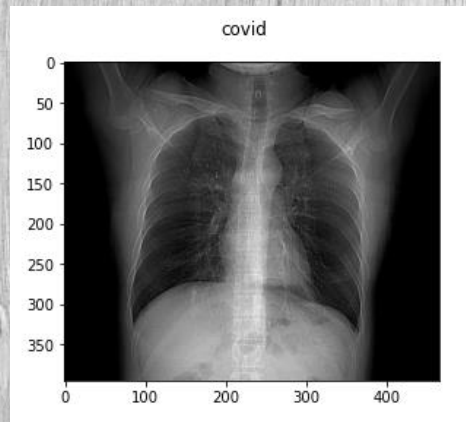
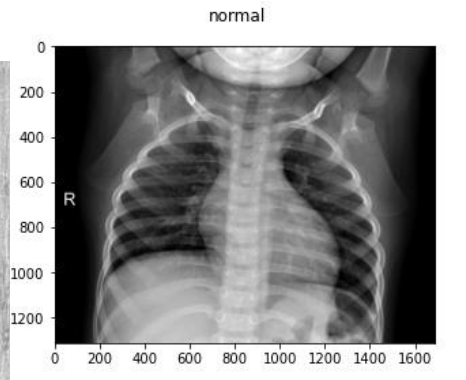
```
import glob
import matplotlib.pyplot as plt
import matplotlib.image as mpimg
%matplotlib inline

normal_images = []
for img_path in glob.glob(DATASET_DIR + '/normal/*'):
    normal_images.append(mpimg.imread(img_path))

fig = plt.figure()
fig.suptitle('normal')
plt.imshow(normal_images[0], cmap='gray')

covid_images = []
for img_path in glob.glob(DATASET_DIR + '/covid/*'):
    covid_images.append(mpimg.imread(img_path))

fig = plt.figure()
fig.suptitle('covid')
plt.imshow(covid_images[0], cmap='gray')
```



Convolutional Neural Network (CNN)



Hyper parameters Initialization:

- All the images are transformed into fixed size **WIDTH** and **HEIGHT** (i.e., 150 x 150).

CHANNELS:

- 3 (for R, G and B)

INPUT_SHAPE:

- Give array as an input to CNN

NB_CLASSES:

- binary

EPOCHS:

- Epoch is when an ENTIRE dataset is passed forward and backward through the neural network only ONCE.

BATCH_SIZE:

- Since one epoch is too big to feed to the computer at once we divide it in several smaller batches.

```
IMG_W = 150
```

```
IMG_H = 150
```

```
CHANNELS = 3
```

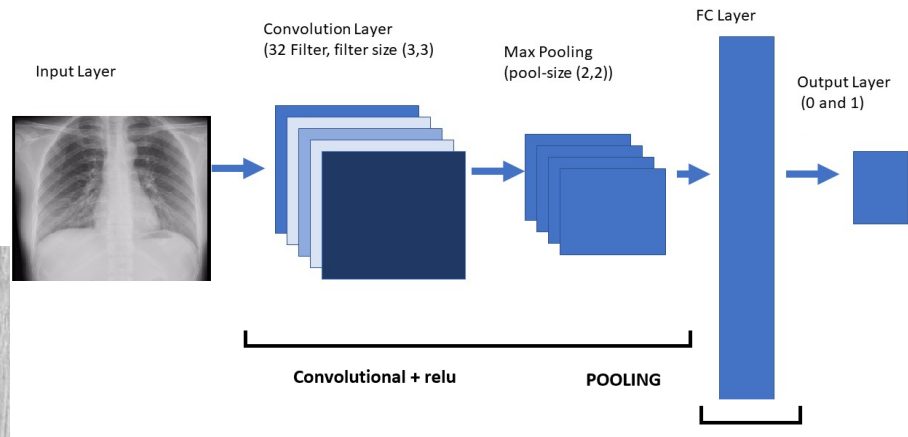
```
INPUT_SHAPE = (IMG_W, IMG_H, CHANNELS)
```

```
NB_CLASSES = 2
```

```
EPOCHS = 48
```

```
BATCH_SIZE = 6
```


Convolutional Neural Network (CNN)



Conv2D

- Conv2D is used to create a convolution filter

Filter Size:

- 32, 32, 64, 128

Kernel Size:

- 3 x 3

Activation Function

- RELU** is used as an activation function to add non-linearity

MaxPooling2D

- Reduce the amount of parameters and computation in the network

```
model = Sequential()
model.add(Conv2D(32, (3, 3), input_shape=INPUT_SHAPE))
model.add(Activation('relu'))
model.add(MaxPooling2D(pool_size=(2, 2)))

model.add(Conv2D(32, (3, 3)))
model.add(Activation('relu'))
model.add(MaxPooling2D(pool_size=(2, 2)))

model.add(Conv2D(64, (3, 3)))
model.add(Activation("relu"))
model.add(Conv2D(250, (3, 3)))
model.add(Activation("relu"))

model.add(Conv2D(128, (3, 3)))
model.add(Activation("relu"))
model.add(AvgPool2D(2, 2))
model.add(Conv2D(64, (3, 3)))
```

Convolutional Neural Network (CNN)

Flatten Layer

- Used for converting a two-dimensional matrix of features into a vector that can be fed into a fully connected neural network classifier

Dense Layer

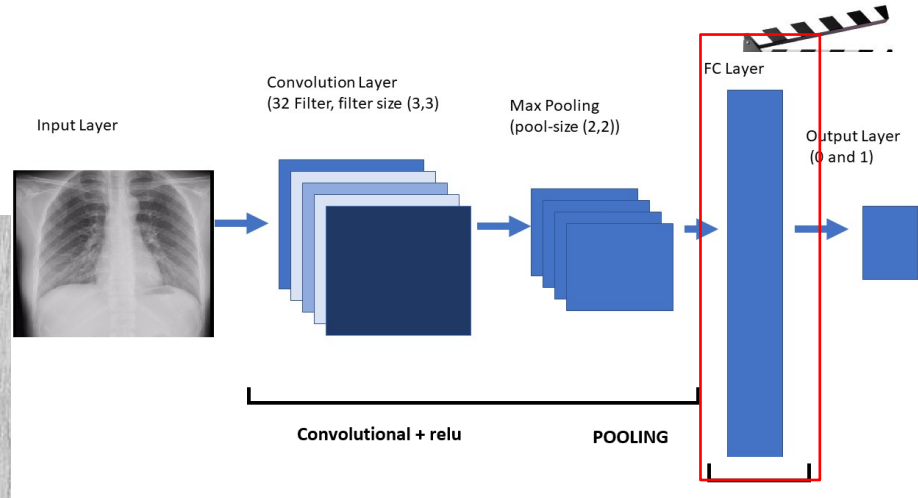
- Each neuron in dense layer receives input from all the neurons in the previous layer, thus densely connected.

Dropout

- Used in regular interval for generalization purpose

Sigmoid

- method is used as an activation method for output layer.



```
model.add(Conv2D(256, (2, 2)))
model.add(Activation("relu"))
model.add(MaxPool2D(2, 2))

model.add(Flatten())
model.add(Dense(32))
model.add(Dropout(0.25))
model.add(Dense(1))
model.add(Activation("sigmoid"))
```

Convolutional Neural Network (CNN)



Compile Deep CNN model:

- For a **binary classification** like our example, the **typical loss function** is the **binary cross-entropy / log loss**

Optimizer:

- RMSprop

Metrics:

- Accuracy

$$Accuracy = \frac{TP+TN}{TP+FP+TN+FN}$$

```
model.compile(loss='binary_crossentropy',  
              optimizer='rmsprop',  
              metrics=['accuracy'])
```

```
model.summary()
```

Model: "sequential_1"

Layer (type)	Output Shape	Param #

conv2d_7 (Conv2D)	(None, 148, 148, 32)	896

activation_7 (Activation)	(None, 148, 148, 32)	0

max_pooling2d_3 (MaxPooling2)	(None, 74, 74, 32)	0

conv2d_8 (Conv2D)	(None, 72, 72, 32)	9248

activation_8 (Activation)	(None, 72, 72, 32)	0

max_pooling2d_4 (MaxPooling2)	(None, 36, 36, 32)	0

conv2d_9 (Conv2D)	(None, 34, 34, 64)	18496

Convolutional Neural Network (CNN)



activation_9 (Activation)	(None, 34, 34, 64)	0

conv2d_10 (Conv2D)	(None, 32, 32, 256)	144256

activation_10 (Activation)	(None, 32, 32, 256)	0

conv2d_11 (Conv2D)	(None, 30, 30, 128)	288128

activation_11 (Activation)	(None, 30, 30, 128)	0

average_pooling2d_2 (Average)	(None, 15, 15, 128)	0

conv2d_12 (Conv2D)	(None, 13, 13, 64)	73792

activation_12 (Activation)	(None, 13, 13, 64)	0

average_pooling2d_3 (Average)	(None, 6, 6, 64)	0

conv2d_13 (Conv2D)	(None, 5, 5, 256)	65792

activation_13 (Activation)	(None, 5, 5, 256)	0

max_pooling2d_5 (MaxPooling2)	(None, 2, 2, 256)	0

flatten (Flatten)	(None, 1024)	0

dense (Dense)	(None, 32)	32800

dropout (Dropout)	(None, 32)	0

dense_1 (Dense)	(None, 1)	33

activation_14 (Activation)	(None, 1)	0
=====		
Total params: 633,435		
Trainable params: 633,435		
Non-trainable params: 0		

Training and Testing

ImageDataGenerator:

- increase number of images by augmenting a few images

Shear_range:

- shear_range is for randomly applying shearing transformations

Zoom_range:

- randomly zooming inside pictures

Horizontal flip:

- randomly flipping half of the images horizontally

Validation_split:

- Set 0.3 (**70% data for testing** and **30% for training**)

target_size

- size of your input images, every image will be resized to this size.

batch_size:

- no. of images to be yielded from the generator per batch.

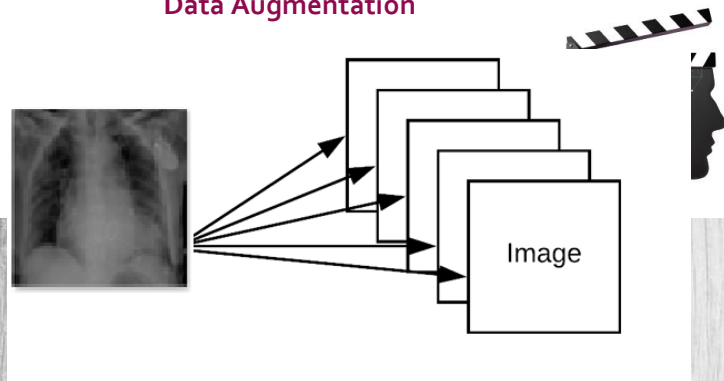
class_mode:

- set **"binary"** if you have only two classes to predict,

shuffle:

- set **True** if you want to shuffle the order of the image that is being yielded, else set **False**.

Data Augmentation



```
train_datagen = ImageDataGenerator(rescale=1./255,  
    shear_range=0.2,  
    zoom_range=0.2,  
    horizontal_flip=True,  
    validation_split=0.3)
```

```
train_generator = train_datagen.flow_from_directory(  
    DATASET_DIR,  
    target_size=(IMG_H, IMG_W),  
    batch_size=BATCH_SIZE,  
    class_mode='binary',  
    subset='training')
```

```
validation_generator = train_datagen.flow_from_directory(  
    DATASET_DIR,  
    target_size=(IMG_H, IMG_W),  
    batch_size=BATCH_SIZE,  
    class_mode='binary',  
    shuffle= False,  
    subset='validation')
```

Fit the model



Model.fit_generator:

steps_per_epoch

- value as the total number of training data points divided by the batch size. Once Keras hits this step count it knows that it's a new epoch.

```
history = model.fit_generator(  
    train_generator,  
    steps_per_epoch = train_generator.samples // BATCH_SIZE,  
    validation_data = validation_generator,  
    validation_steps = validation_generator.samples // BATCH_SIZE,  
    epochs = EPOCHS)
```

```
Epoch 1/48  
11/11 [=====] - 5s 454ms/step - loss: 0.9752 -  
accuracy: 0.7302 - val_loss: 0.5168 - val_accuracy: 0.8750  
Epoch 2/48  
11/11 [=====] - 5s 427ms/step - loss: 0.6296 -  
accuracy: 0.6984 - val_loss: 0.5840 - val_accuracy: 0.8750  
Epoch 3/48  
11/11 [=====] - 5s 450ms/step - loss: 0.7205 -  
accuracy: 0.7273 - val_loss: 0.6117 - val_accuracy: 0.8750  
Epoch 4/48  
11/11 [=====] - 5s 423ms/step - loss: 0.6340 -  
accuracy: 0.7143 - val_loss: 0.4250 - val_accuracy: 0.8750
```

```
Epoch 48/48  
11/11 [=====] - 5s 445ms/step - loss: 0.0291 -  
accuracy: 1.0000 - val_loss: 0.0364 - val_accuracy: 0.9583
```

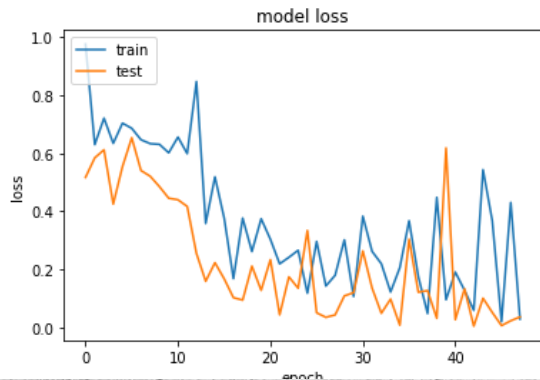
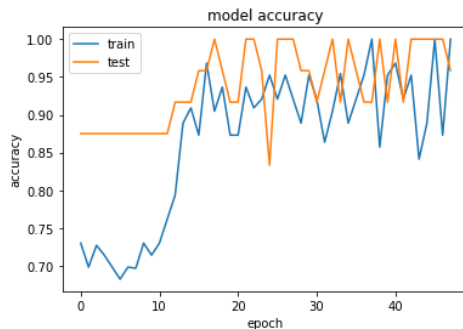

Performance Analysis



- The epochs history shows that accuracy gradually increases and achieved **+95% accuracy** on both training and validation set.
- The loss value of the model **gradually decreases** as the number of training **epoch increases**.

```
plt.plot(history.history['accuracy'])
plt.plot(history.history['val_accuracy'])
plt.title('model accuracy')
plt.ylabel('accuracy')
plt.xlabel('epoch')
plt.legend(['train', 'test'], loc='upper left')
plt.show()
```

```
plt.plot(history.history['loss'])
plt.plot(history.history['val_loss'])
plt.title('model loss')
plt.ylabel('loss')
plt.xlabel('epoch')
plt.legend(['train', 'test'], loc='upper left')
plt.show()
```



Conclusion



- CNN model has been proven useful in detecting “covid” via X-ray images data.
- The model has tendency to predict true positives with significant accuracy
- However, the size of data is too small to draw a generic conclusion

References



- [1] Wu F., Zhao S., Yu B. A new coronavirus associated with human respiratory disease in China. Nature. 2020;579(7798):265–269. [[PMC free article](#)] [[PubMed](#)] [[Google Scholar](#)]
- [2] Huang C., Wang Y. Clinical features of patients infected with 2019 novel coronavirus in Wuhan, China. Lancet. 2020;395(10223):497–506.
- [3] World Health Organization . World Health Organization (WHO); 2020. Pneumonia of Unknown Cause–China. Emergencies Preparedness, Response, Disease Outbreak News.
- [4] Wu Z., McGoogan J.M. Characteristics of and important lessons from the coronavirus disease 2019 (COVID-19) outbreak in China: summary of a report of 72 314 cases from the Chinese Center for Disease Control and Prevention. Jama. 2020;323(13):1239–1242.

THANKS!