

# Project 4

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## Business understanding

The medical dataset comes from Kermay et al. contains a set of x-ray images of pediatric patients. The images will show whether the patients have pneumonia or not. Our task is to build a model that can classify whether a given patient has pneumonia given a chest x-ray image. Since this is an Image Classification problem, we will solve it with Deep Learning.

## The Dataset

The dataset that we will use for image classification is the chese\_xray which contains two categories: Pneumonia and Normal. The data was downloaded from <https://data.mendeley.com/datasets/rscbjbr9sj/3> (<https://data.mendeley.com/datasets/rscbjbr9sj/3>) to the local drive and unzipped. The data set is assigned into two folders (train and test) and contains a subfolder for each Pneumonia and Normal category. In each of the folders, there are a lot of x-ray images. To check how many samples were in each category, we used the OS.listdir methods.

```
In [2]: import numpy as np
import os, shutil
import pandas as pd
from keras.preprocessing.image import ImageDataGenerator
from tensorflow.keras.applications import VGG19
from tensorflow.keras import layers
from tensorflow.keras import models
from tensorflow.keras import optimizers
import matplotlib.pyplot as plt
from sklearn.metrics import classification_report, confusion_matrix
import seaborn as sns
```

```
In [3]: list_train_normal = os.listdir('chest_xray/train/normal')
list_train_PNEUMONIA = os.listdir('chest_xray/train/PNEUMONIA')
list_test_normal = os.listdir('chest_xray/test/normal')
list_test_PNEUMONIA = os.listdir('chest_xray/test/PNEUMONIA')

len(list_train_normal), len(list_train_PNEUMONIA), len(list_test_normal), len(list_test_PNI
```

```
Out[3]: (1349, 3884, 235, 390)
```

In the train folder, there is a regular folder that contains 1349 images and the PNEUMONIA folder, which includes 3884 images. The NORMAL folder in the test folder contains 235 pictures, and the PNEUMONIA folder contains 390 images. The images in each folder are too large for the modeling since our local computer is not very powerful for multiple testing. Therefore, we need to downsample the dataset to find the optimal

model and parameter first. We are then using the entire dataset to train and test our model. Based on our earlier experience, we will use 20% of the total dataset to model our model. We also need to make 10% of the training data to the validation dataset.

## Plan

1. Downsample the data set by randomly choosing 20% of the initial training and testing images to the new data\_org\_subset folder. Make a new validation folder and randomly select 5% of the pictures from the training folder.
2. Define the trained generator, validation generator, and test generator.
3. Build a baseline model.
4. Build the deep learning model base on the Pretrained CNN (VGG19) by adding a few fully connected layers. Then, train the model with selected images.
5. Retrain the model with complete training data.
6. Evaluate the model with the test images.

### **1. Rebuild the data subset folder with 20% of the original images**

In [5]: *# define the old and new direction of dataset*

*# define a new method to transfer the images between two folder*

```
def transfer(no_of_files, source, dest):  
    for i in range(no_of_files):  
        #Variable random_file stores the name of the random file chosen  
        random_file=np.random.choice(os.listdir(source))  
        # print("%d} %s"%(i+1,random_file))  
        source_file="%s/%s"%(source,random_file)  
        dest_file=dest  
        #"shutil.move" function moves file from one directory to another  
        shutil.copy(source_file,dest_file)
```

*# set the propotion of images transfered to the new folders p\_val, p\_train, p\_test and*

*# define a new method to creat and transfer images*

```
def make_subset (old_dir, new_root_dir, p_val, p_train, p_test) :
```

*# make the root dir folder*

```
os.mkdir(new_root_dir)
```

*# define the name of subset to save all the images in different categories*

```
dir_names = ['train', 'val', 'test']
```

```
cat_names = ['normal', 'PNEUMONIA']
```

```
for d in dir_names:
```

```
    new_dir = os.path.join(new_root_dir, d)
```

```
    os.mkdir(new_dir)
```

*# make the source dir to train and test folder, since we donot have validation in train*

*# we make it to train folder*

```
    if d == 'val':
```

```
        source_dir = os.path.join(old_dir, 'train')
```

```
    else:
```

```
        source_dir = os.path.join(old_dir, d)
```

```
    for cat in cat_names:
```

```
        new_cat = os.path.join(new_dir, cat)
```

```
        source = os.path.join(source_dir, cat )
```

```
        os.mkdir(new_cat)
```

```
        no_of_files = len(os.listdir(source))
```

*# set the number of copy to 20% from source folder. For the validation folder,*

```
        if d == 'val':
```

```
            no_of_copy = int(p_val * no_of_files)
```

```
        if d == 'train':
```

```
            no_of_copy = int(p_train * no_of_files)
```

```
        if d == 'test':
```

```
            no_of_copy = int(p_test * no_of_files )
```

```
        #print('d = ', d)
```

```
        print('copy {} of files in {} total files from {} to {}'.format(no_of_copy,no_of_files,source,new_cat))  
        transfer(no_of_copy, source, new_cat)
```

```
old_dir = 'chest_xray/'
```

```
new_root_dir = 'data_org_subset/'
```

```
make_subset(old_dir, new_root_dir, p_val = 0.05, p_train = 0.2, p_test = 0.2)
```

```

copy 269 of files in 1349 total files from chest_xray/train\normal to data_org_subset/train\normal
copy 776 of files in 3884 total files from chest_xray/train\PNEUMONIA to data_org_subset/train\PNEUMONIA
copy 67 of files in 1349 total files from chest_xray/train\normal to data_org_subset/val\normal
copy 194 of files in 3884 total files from chest_xray/train\PNEUMONIA to data_org_subset/val\PNEUMONIA
copy 47 of files in 235 total files from chest_xray/test\normal to data_org_subset/test\normal
copy 78 of files in 390 total files from chest_xray/test\PNEUMONIA to data_org_subset/test\PNEUMONIA

```

We copied 20% of the training and testing images from the original folder. We also made a new folder for validation and randomly selected 5% of the images from the training folder.

```

In [3]: old_dir = 'chest_xray/'
new_root_dir = 'data_org_subset/'

```

## 2. Define the train generator, validation generator and test generator.

```

In [5]: # define the direction for train , validation and test folder

train_dir = '{}train'.format(new_root_dir)
validation_dir = '{}val/'.format(new_root_dir)
test_dir = '{}test/'.format(new_root_dir)

train_datagen = ImageDataGenerator(rescale=1./255,
                                   rotation_range=40,
                                   width_shift_range=0.2,
                                   height_shift_range=0.2,
                                   shear_range=0.2,
                                   zoom_range=0.2,
                                   horizontal_flip=True,
                                   fill_mode='nearest')

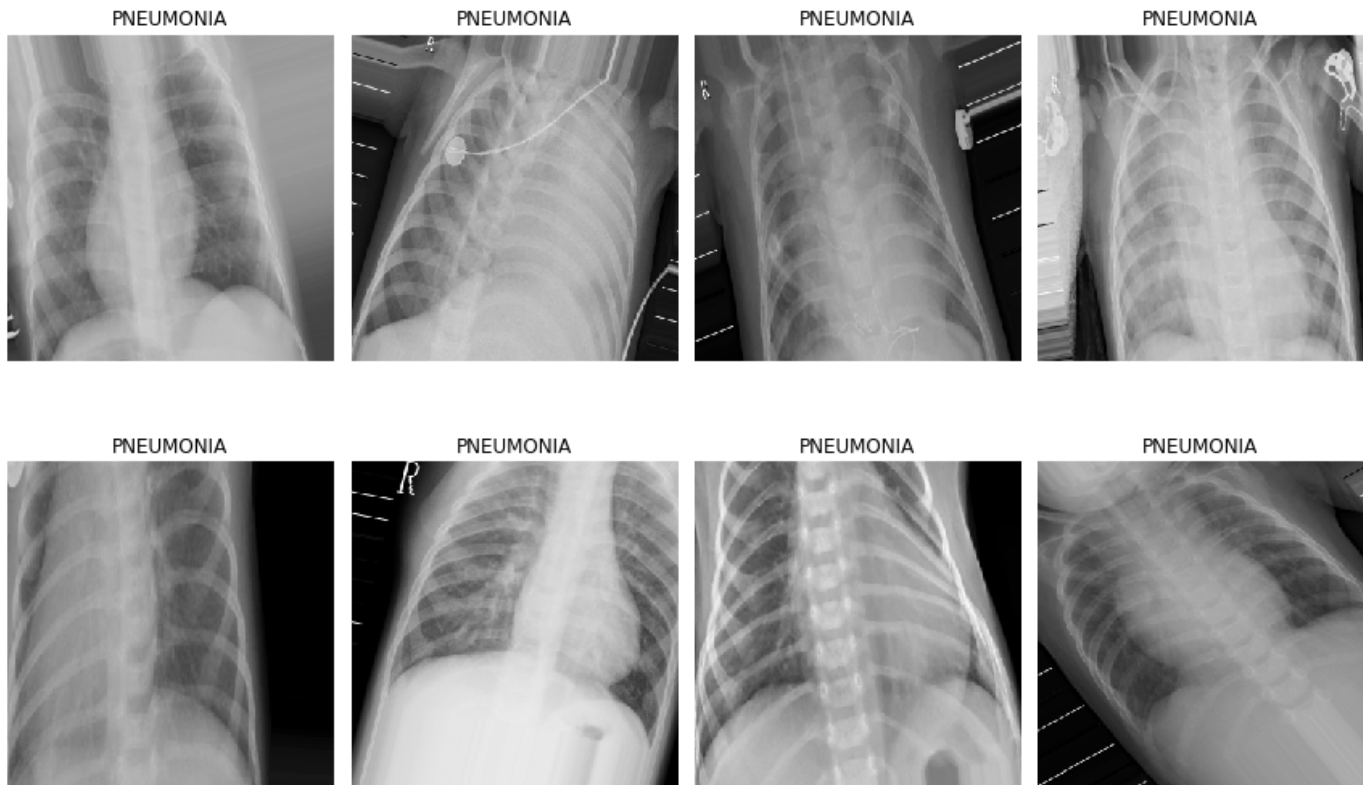
train_generator = train_datagen.flow_from_directory(train_dir,
                                                    target_size=(300, 300),
                                                    batch_size= 20,
                                                    class_mode='categorical')

# Get all the data in the directory split/validation (200 images), and reshape them
val_generator = ImageDataGenerator(rescale=1./255).flow_from_directory(validation_dir,
                                                                    target_size=(300, 300),
                                                                    batch_size=20,
                                                                    class_mode='categorical')

```

Found 957 images belonging to 2 classes.  
Found 255 images belonging to 2 classes.

```
In [7]: # plotsome of the train set images we resampled in the train dataset
plt.figure(figsize=(12, 8))
for i in range(0, 8):
    plt.subplot(2, 4, i+1)
    for X_batch, Y_batch in train_generator:
        image = X_batch[0]
        dic = {0: 'NORMAL', 1: 'PNEUMONIA'}
        plt.title(dic[Y_batch[0][0]])
        plt.axis('off')
        plt.imshow(np.squeeze(image), cmap='gray', interpolation='nearest')
        break
plt.tight_layout()
plt.show()
```



We plot some of the images in the training dataset. However, I can not tell which one is a case of pneumonia and which one is a normal case just by looking at the pictures. So now we will train the computer with a Pretrained CNN model to predict whether the picture belongs to pneumonia or normal case.

### 3. Build a baseline model

```
In [11]: model = models.Sequential()
model.add(layers.Conv2D(32, (3, 3), activation='relu',
                        input_shape=(300, 300, 3)))
model.add(layers.MaxPooling2D((2, 2)))
model.add(layers.Conv2D(64, (3, 3), activation='relu'))
model.add(layers.MaxPooling2D((2, 2)))
model.add(layers.Conv2D(128, (3, 3), activation='relu'))
model.add(layers.MaxPooling2D((2, 2)))
model.add(layers.Conv2D(128, (3, 3), activation='relu'))
model.add(layers.MaxPooling2D((2, 2)))
model.add(layers.Flatten())
model.add(layers.Dense(64, activation='relu'))
model.add(layers.Dense(128, activation='relu'))
model.add(layers.Dense(256, activation='relu'))
model.add(layers.Dense(512, activation='relu'))
model.add(layers.Dense(2, activation='softmax'))

model.compile(loss='categorical_crossentropy',
              optimizer=optimizers.RMSprop(lr=1e-4),
              metrics=['acc'])
model.summary()
```

Model: "sequential\_1"

Layer (type)	Output Shape	Param #
=====		
conv2d_4 (Conv2D)	(None, 298, 298, 32)	896
max_pooling2d_4 (MaxPooling 2D)	(None, 149, 149, 32)	0
conv2d_5 (Conv2D)	(None, 147, 147, 64)	18496
max_pooling2d_5 (MaxPooling 2D)	(None, 73, 73, 64)	0
conv2d_6 (Conv2D)	(None, 71, 71, 128)	73856
max_pooling2d_6 (MaxPooling 2D)	(None, 35, 35, 128)	0
conv2d_7 (Conv2D)	(None, 33, 33, 128)	147584
max_pooling2d_7 (MaxPooling 2D)	(None, 16, 16, 128)	0
flatten_1 (Flatten)	(None, 32768)	0
dense_5 (Dense)	(None, 64)	2097216
dense_6 (Dense)	(None, 128)	8320
dense_7 (Dense)	(None, 256)	33024
dense_8 (Dense)	(None, 512)	131584
dense_9 (Dense)	(None, 2)	1026
=====		
Total params: 2,512,002		

Trainable params: 2,512,002  
Non-trainable params: 0

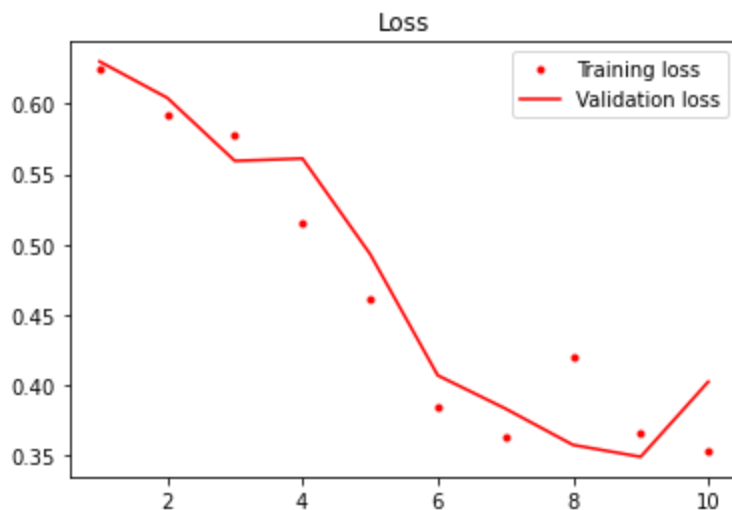
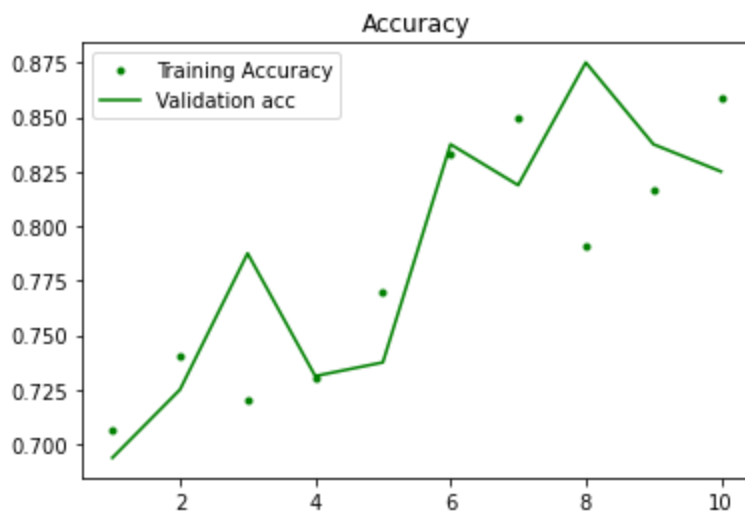
---

```
In [12]: history = model.fit(train_generator,  
                             steps_per_epoch=15,  
                             epochs=10,  
                             validation_data=val_generator,  
                             validation_steps=8)
```

```
Epoch 1/10  
15/15 [=====] - 29s 2s/step - loss: 0.6246 - acc: 0.7067 - val_loss: 0.6301 - val_acc: 0.6938  
Epoch 2/10  
15/15 [=====] - 28s 2s/step - loss: 0.5921 - acc: 0.7407 - val_loss: 0.6043 - val_acc: 0.7250  
Epoch 3/10  
15/15 [=====] - 25s 2s/step - loss: 0.5780 - acc: 0.7200 - val_loss: 0.5595 - val_acc: 0.7875  
Epoch 4/10  
15/15 [=====] - 24s 2s/step - loss: 0.5158 - acc: 0.7300 - val_loss: 0.5612 - val_acc: 0.7312  
Epoch 5/10  
15/15 [=====] - 22s 1s/step - loss: 0.4616 - acc: 0.7700 - val_loss: 0.4933 - val_acc: 0.7375  
Epoch 6/10  
15/15 [=====] - 23s 2s/step - loss: 0.3852 - acc: 0.8333 - val_loss: 0.4072 - val_acc: 0.8375  
Epoch 7/10  
15/15 [=====] - 22s 1s/step - loss: 0.3629 - acc: 0.8500 - val_loss: 0.3836 - val_acc: 0.8188  
Epoch 8/10  
15/15 [=====] - 22s 1s/step - loss: 0.4203 - acc: 0.7912 - val_loss: 0.3578 - val_acc: 0.8750  
Epoch 9/10  
15/15 [=====] - 22s 1s/step - loss: 0.3656 - acc: 0.8167 - val_loss: 0.3493 - val_acc: 0.8375  
Epoch 10/10  
15/15 [=====] - 21s 1s/step - loss: 0.3532 - acc: 0.8586 - val_loss: 0.4027 - val_acc: 0.8250
```

In [13]: *# Plot the accuracy and loss for train and validation.*

```
def plot_acc(history):  
    train_acc = history.history['acc']  
    val_acc = history.history['val_acc']  
    train_loss = history.history['loss']  
    val_loss = history.history['val_loss']  
    epoch = range(1, len(train_acc) + 1)  
    plt.plot(epoch, train_acc, 'g.', label='Training Accuracy')  
    plt.plot(epoch, val_acc, 'g', label='Validation acc')  
    plt.title('Accuracy')  
    plt.legend()  
    plt.figure()  
    plt.plot(epoch, train_loss, 'r.', label='Training loss')  
    plt.plot(epoch, val_loss, 'r', label='Validation loss')  
    plt.title('Loss')  
    plt.legend()  
    plt.show()  
plot_acc(history)
```



The acc and loss curve of training are pretty close to the validation scores in each step. However, the number is only 85% which is not very high. Thus we will continue to try the model with pretrain network VGG19.

## 4. Build the model base on pretrain network VGG19.



In [27]:

```
# defined the pretrained model VGG19 and add more layer to the network.
cnn_base = VGG19(weights='imagenet',
                  include_top=False,
                  input_shape=(300, 300, 3))

# Define Model Architecture
model = models.Sequential()
model.add(cnn_base)
model.add(layers.Flatten())
model.add(layers.Dense(64, activation='relu'))
model.add(layers.Dense(128, activation='relu'))
model.add(layers.Dense(256, activation='relu'))
model.add(layers.Dense(128, activation='relu'))
model.add(layers.Dense(2, activation='softmax'))

cnn_base.trainable = False

model.summary()
```

Model: "sequential"

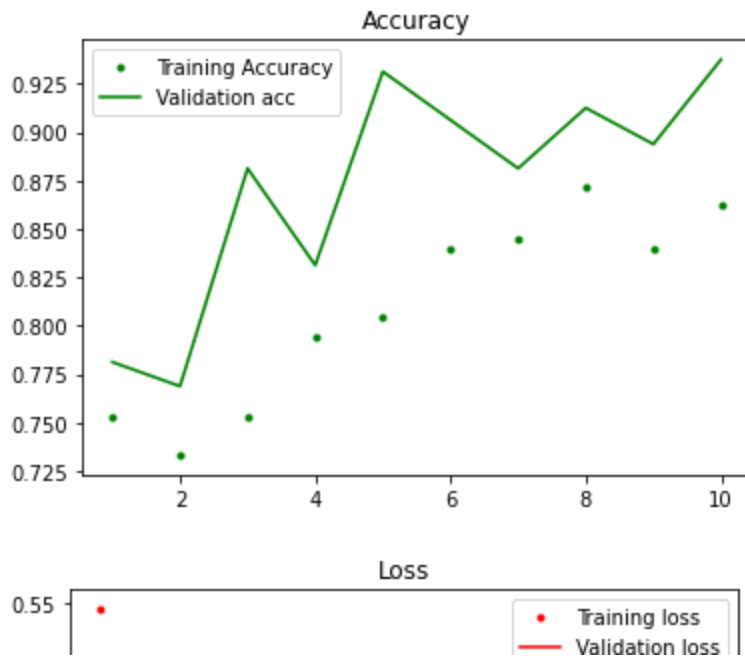
Layer (type)	Output Shape	Param #
=====		
vgg19 (Functional)	(None, 9, 9, 512)	20024384
flatten (Flatten)	(None, 41472)	0
dense (Dense)	(None, 64)	2654272
dense_1 (Dense)	(None, 128)	8320
dense_2 (Dense)	(None, 256)	33024
dense_3 (Dense)	(None, 128)	32896
dense_4 (Dense)	(None, 2)	258
=====		
Total params: 22,753,154		
Trainable params: 2,728,770		
Non-trainable params: 20,024,384		

```
In [9]: # Compilation
model.compile(loss='categorical_crossentropy',
              optimizer=optimizers.RMSprop(learning_rate=2e-5),
              metrics=['acc'])

# Fitting the Model
history = model.fit(train_generator,
                    steps_per_epoch=15,
                    epochs=10,
                    validation_data=val_generator,
                    validation_steps=8)
```

```
Epoch 1/10
15/15 [=====] - 103s 7s/step - loss: 0.5447 - acc: 0.7533 - val_
loss: 0.4575 - val_acc: 0.7812
Epoch 2/10
15/15 [=====] - 100s 7s/step - loss: 0.4953 - acc: 0.7333 - val_
loss: 0.4226 - val_acc: 0.7688
Epoch 3/10
15/15 [=====] - 107s 7s/step - loss: 0.4545 - acc: 0.7533 - val_
loss: 0.3840 - val_acc: 0.8813
Epoch 4/10
15/15 [=====] - 108s 7s/step - loss: 0.4147 - acc: 0.7946 - val_
loss: 0.3535 - val_acc: 0.8313
Epoch 5/10
15/15 [=====] - 110s 7s/step - loss: 0.4017 - acc: 0.8047 - val_
loss: 0.3721 - val_acc: 0.9312
Epoch 6/10
15/15 [=====] - 112s 8s/step - loss: 0.3587 - acc: 0.8400 - val_
loss: 0.3239 - val_acc: 0.9062
Epoch 7/10
15/15 [=====] - 108s 7s/step - loss: 0.3368 - acc: 0.8451 - val_
loss: 0.3544 - val_acc: 0.8813
Epoch 8/10
15/15 [=====] - 109s 7s/step - loss: 0.2969 - acc: 0.8721 - val_
loss: 0.2719 - val_acc: 0.9125
Epoch 9/10
15/15 [=====] - 111s 8s/step - loss: 0.3515 - acc: 0.8400 - val_
loss: 0.3143 - val_acc: 0.8938
Epoch 10/10
15/15 [=====] - 104s 7s/step - loss: 0.3330 - acc: 0.8620 - val_
loss: 0.2501 - val_acc: 0.9375
```

```
In [12]: # Plot the accuracy and loss for train and validation.
plot_acc(history)
```



The acc and loss curve of training gave us a pretty good score, and the validation scores are going to a similar range in each step. Thus, we can use the same model for better training on the entire training dataset.

```
In [13]: #save model
model.save('results_on_partial_dataset.h5')
```

## 5. Retrain the model with full training data.

```
In [14]: # remake the folder of train, val, test folder for full dataset.
#Transfer 90% of train images to new train and 10% of train images to new validation folder
#Transfer 100% of test to new test folder
old_dir = 'chest_xray/'
new_root_dir = 'data_org_full/'
make_subset(old_dir, new_root_dir, p_val = 0.1, p_train = 0.9, p_test = 1)
```

```
copy 1214 of files in 1349 total files from chest_xray/train\normal to data_org_full/train\normal
copy 3495 of files in 3884 total files from chest_xray/train\PNEUMONIA to data_org_full/train\PNEUMONIA
copy 134 of files in 1349 total files from chest_xray/train\normal to data_org_full/val\normal
copy 388 of files in 3884 total files from chest_xray/train\PNEUMONIA to data_org_full/val\PNEUMONIA
copy 235 of files in 235 total files from chest_xray/test\normal to data_org_full/test\normal
copy 390 of files in 390 total files from chest_xray/test\PNEUMONIA to data_org_full/test\PNEUMONIA
```

```

In [42]: train_dir = '{}train'.format(new_root_dir)
validation_dir = '{}val/'.format(new_root_dir)
test_dir = '{}test/'.format(new_root_dir)

full_train_datagen = ImageDataGenerator(rescale=1./255,
                                        rotation_range=40,
                                        width_shift_range=0.2,
                                        height_shift_range=0.2,
                                        shear_range=0.2,
                                        zoom_range=0.2,
                                        horizontal_flip=True,
                                        fill_mode='nearest')

full_train_generator = full_train_datagen.flow_from_directory(train_dir,
                                                             target_size=(300, 300),
                                                             batch_size= 20,
                                                             class_mode='categorical')

# Get all the data in the directory split/validation (, and reshape them
full_val_generator = ImageDataGenerator(rescale=1./255).flow_from_directory(validation_dir,
                                                                              target_size=(300, 300),
                                                                              batch_size=20,
                                                                              class_mode='categorical')

```

Found 3132 images belonging to 2 classes.

Found 492 images belonging to 2 classes.

```
In [43]: # recompile the model and fit to the full training dataset.
model.compile(loss='categorical_crossentropy',
              optimizer=optimizers.RMSprop(learning_rate=2e-5),
              metrics=['acc'])

history = model.fit(full_train_generator,
                    steps_per_epoch=15,
                    epochs=20,
                    validation_data=full_val_generator,
                    validation_steps=10)
```

```
Epoch 1/20
15/15 [=====] - 103s 7s/step - loss: 0.2719 - acc: 0.8870 - va
l_loss: 0.2330 - val_acc: 0.9100
Epoch 2/20
15/15 [=====] - 106s 7s/step - loss: 0.2238 - acc: 0.9167 - va
l_loss: 0.2073 - val_acc: 0.9250
Epoch 3/20
15/15 [=====] - 106s 7s/step - loss: 0.2244 - acc: 0.8900 - va
l_loss: 0.2235 - val_acc: 0.9050
Epoch 4/20
15/15 [=====] - 106s 7s/step - loss: 0.2212 - acc: 0.9100 - va
l_loss: 0.2736 - val_acc: 0.8900
Epoch 5/20
15/15 [=====] - 103s 7s/step - loss: 0.2360 - acc: 0.9007 - va
l_loss: 0.3543 - val_acc: 0.8200
Epoch 6/20
15/15 [=====] - 105s 7s/step - loss: 0.2231 - acc: 0.9033 - va
l_loss: 0.2692 - val_acc: 0.8850
Epoch 7/20
15/15 [=====] - 105s 7s/step - loss: 0.1796 - acc: 0.9300 - va
l_loss: 0.2150 - val_acc: 0.9250
Epoch 8/20
15/15 [=====] - 104s 7s/step - loss: 0.1946 - acc: 0.9100 - va
l_loss: 0.1999 - val_acc: 0.9100
Epoch 9/20
15/15 [=====] - 104s 7s/step - loss: 0.2327 - acc: 0.8967 - va
l_loss: 0.2443 - val_acc: 0.9200
Epoch 10/20
15/15 [=====] - 104s 7s/step - loss: 0.2043 - acc: 0.9000 - va
l_loss: 0.2135 - val_acc: 0.9100
Epoch 11/20
15/15 [=====] - 104s 7s/step - loss: 0.2299 - acc: 0.9067 - va
l_loss: 0.1807 - val_acc: 0.9250
Epoch 12/20
15/15 [=====] - 106s 7s/step - loss: 0.2253 - acc: 0.9067 - va
l_loss: 0.1874 - val_acc: 0.9200
Epoch 13/20
15/15 [=====] - 105s 7s/step - loss: 0.2054 - acc: 0.9100 - va
l_loss: 0.2096 - val_acc: 0.9250
Epoch 14/20
15/15 [=====] - 103s 7s/step - loss: 0.2368 - acc: 0.9178 - va
l_loss: 0.3129 - val_acc: 0.8550
Epoch 15/20
15/15 [=====] - 107s 7s/step - loss: 0.1851 - acc: 0.9433 - va
l_loss: 0.2631 - val_acc: 0.8850
Epoch 16/20
15/15 [=====] - 105s 7s/step - loss: 0.2751 - acc: 0.8867 - va
l_loss: 0.1733 - val_acc: 0.9450
Epoch 17/20
15/15 [=====] - 105s 7s/step - loss: 0.2585 - acc: 0.8967 - va
```

l\_loss: 0.2074 - val\_acc: 0.9350

Epoch 18/20

15/15 [=====] - 104s 7s/step - loss: 0.2191 - acc: 0.8967 - val\_loss: 0.3136 - val\_acc: 0.8450

Epoch 19/20

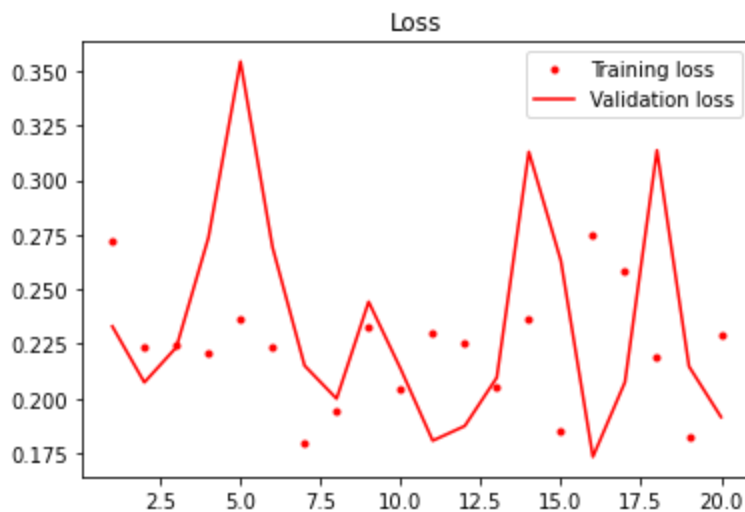
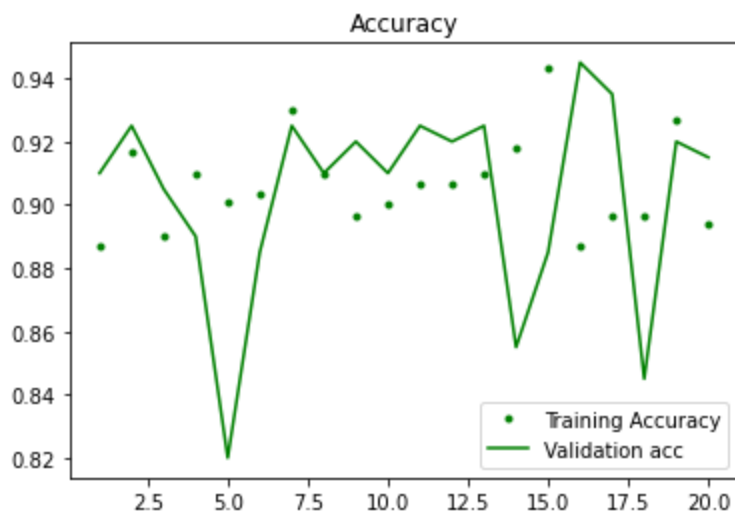
15/15 [=====] - 104s 7s/step - loss: 0.1822 - acc: 0.9267 - val\_loss: 0.2146 - val\_acc: 0.9200

Epoch 20/20

15/15 [=====] - 102s 7s/step - loss: 0.2287 - acc: 0.8938 - val\_loss: 0.1915 - val\_acc: 0.9150

```
In [50]: #save model
model.save('results_on_full_dataset.h5')
```

```
In [52]: plot_acc(history)
```



In this fitting, both training accuracy and validation accuracy are very high. Even though the fluctuation of validation accuracy is bigger than training accuracy, both accuracies generally had the same trend.

## 6. Evaluate the model with the test images.

In [ ]: *# Get all the data in the directory split/test (180 images), and reshape them*

```
test_generator = ImageDataGenerator(rescale=1./255).flow_from_directory(test_dir,
                                                                    target_size=(300, 300),
                                                                    batch_size=399,
                                                                    class_mode='categorical',
                                                                    shuffle=False)

# generate the test_labels which is the y_true data
test_images, test_labels = next(test_generator)
```

Generate the test\_labels for making the confusion box later.

```
In [48]: test_datagen = ImageDataGenerator(rescale=1./255,
                                           rotation_range=40,
                                           width_shift_range=0.2,
                                           height_shift_range=0.2,
                                           shear_range=0.2,
                                           zoom_range=0.2,
                                           horizontal_flip=True,
                                           fill_mode='nearest')
test_generator2 = test_datagen.flow_from_directory(test_dir,
                                                  target_size=(300, 300),
                                                  batch_size=20,
                                                  class_mode='categorical',
                                                  shuffle=False)

test_loss, test_acc = model.evaluate(test_generator2, steps=10)
```

Found 399 images belonging to 2 classes.

10/10 [=====] - 42s 4s/step - loss: 0.0685 - acc: 0.9750

The test accuracy of the model on test dataset are 95% which is very high also.

```
In [12]: # calculate the predicitons
preds = model.predict(test_generator2, verbose = 1 )

20/20 [=====] - 92s 5s/step
```

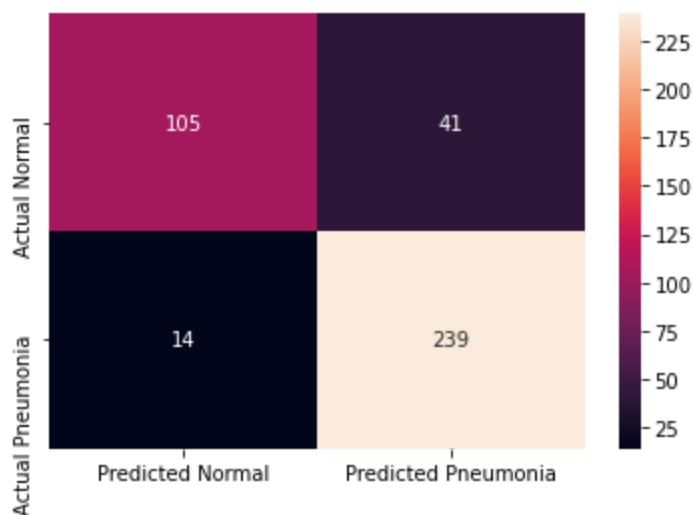
```
In [54]: predictions = preds.copy()
predictions[predictions <= 0.6] = 0
predictions[predictions > 0.6] = 1
```

We calculate the predictions of the classificaiton with our model.

In [57]: *# calculate the confusion box for final model*

```
cm = pd.DataFrame(data=confusion_matrix(test_labels[:,0], predictions[:,0], labels=[0,1]),
                  columns=["Predicted Normal", "Predicted Pneumonia"])
sns.heatmap(cm,annot=True,fmt="d")
```

Out[57]: <AxesSubplot:>



In [58]: *# print the scores for normal and pneumonia categories*

```
print(classification_report(y_true=test_labels[:,0],y_pred=predictions[:,0],target_names =
```

	precision	recall	f1-score	support
NORMAL	0.88	0.72	0.79	146
PNEUMONIA	0.85	0.94	0.90	253
accuracy			0.86	399
macro avg	0.87	0.83	0.84	399
weighted avg	0.86	0.86	0.86	399

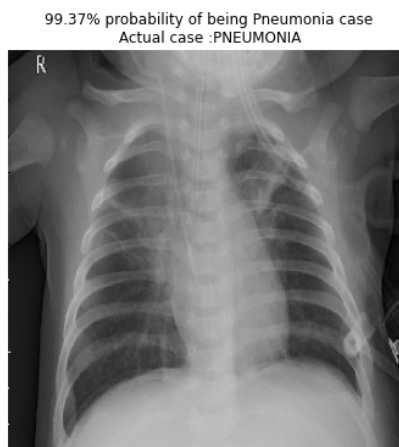
The confusion box shows that the TP and TN predictions are much higher than the FN and FP results. The f1-score for both normal and pneumonia data are 0.79 and 0.9, which are very reasonable too.



```
In [25]: # print some of the predicted images with percentage of predictions
test_generator.reset()
x=np.concatenate([test_generator.next()[0] for i in range(test_generator.__len__())])
y=np.concatenate([test_generator.next()[1] for i in range(test_generator.__len__())])
print(x.shape)
print(y.shape)
dic = {0:'NORMAL', 1:'PNEUMONIA'}
plt.figure(figsize=(20,14))

#for i in range(0+200, 9+200):
for idx, i in enumerate(np.random.randint(1, 388, 6)):
    plt.subplot(2, 3, idx+1)
    if preds[i, 0] >= 0.5:
        out = ('{:.2%} probability of being Pneumonia case'.format(preds[i][0]))
    else:
        out = ('{:.2%} probability of being Normal case'.format(1-preds[i][0]))
    plt.title(out + "\n Actual case : " + dic[y[i][0]])
    plt.imshow(np.squeeze(x[i]))
    plt.axis('off')
plt.show()
```

(399, 300, 300, 3)  
(399, 2)



We randomly plot some of the pictures from the test folder and give the prediction and actual case of the picture. The prediction and actual results are identical to each other in our samples.

## **Conclusion**

Based on 20% of the whole dataset, we created a CNN model based on a Pretrained model (VGG19), which can classify X-ray images as Pneumonia cases or Normal cases. The model was then retrained with the whole dataset and tested with the separated test images. The accuracy of the prediction is around 95%.