

Predicting Lung Disease Using Deep Learning

Please download the dataset from the below url

<https://www.kaggle.com/paultimothymooney/chest-xray-pneumonia>

In [53]: `!pwd`

`/Users/mybeast/Documents`

In [1]: `# import the libraries as shown below`

```
from keras.layers import Input, Lambda, Dense, Flatten
from keras.models import Model

#we will create generic code which can be used for other base models as well
#from keras.applications.resnet50 import ResNet50
from keras.applications.vgg16 import VGG16
from keras.applications.vgg16 import preprocess_input
from keras.preprocessing import image
from keras.preprocessing.image import ImageDataGenerator
from keras.models import Sequential
import numpy as np
from glob import glob
import matplotlib.pyplot as plt
```

In [2]: `# re-size all the images to this`

```
IMAGE_SIZE = [224, 224]
```

```
train_path = 'Datasets/train'
```

```
valid_path = 'Datasets/test'
```

In [3]: `# Import the Vgg 16 library as shown below and add preprocessing layer to the model`
`# Here we will be using imagenet weights`

```
vgg = VGG16(input_shape=IMAGE_SIZE + [3], weights='imagenet', include_top=False)
```

Downloading data from https://storage.googleapis.com/tensorflow/keras-applications/vgg16/vgg16_weights_tf_dim_ordering_tf_kernels_notop.h5

58892288/58889256 [=====] - 26s 0us/step

58900480/58889256 [=====] - 26s 0us/step

In [4]: `# don't train existing weights`

```
for layer in vgg.layers:
    layer.trainable = False
```

In [5]: `# useful for getting number of output classes in order to know how many output classes there are`
`folders = glob('Datasets/train/*')`

```
In [6]: # our layers - you can add more if you want  
x = Flatten()(vgg.output)
```

```
In [7]: prediction = Dense(len(folders), activation='softmax')(x)  
  
# create a model object  
model = Model(inputs=vgg.input, outputs=prediction) #create a model with vgg
```

```
In [8]: # view the structure of the model  
model.summary()
```

Model: "model"

Layer (type)	Output Shape	Param #
=====		
input_1 (InputLayer)	[(None, 224, 224, 3)]	0
block1_conv1 (Conv2D)	(None, 224, 224, 64)	1792
block1_conv2 (Conv2D)	(None, 224, 224, 64)	36928
block1_pool (MaxPooling2D)	(None, 112, 112, 64)	0
block2_conv1 (Conv2D)	(None, 112, 112, 128)	73856
block2_conv2 (Conv2D)	(None, 112, 112, 128)	147584
block2_pool (MaxPooling2D)	(None, 56, 56, 128)	0
block3_conv1 (Conv2D)	(None, 56, 56, 256)	295168
block3_conv2 (Conv2D)	(None, 56, 56, 256)	590080
block3_conv3 (Conv2D)	(None, 56, 56, 256)	590080
block3_pool (MaxPooling2D)	(None, 28, 28, 256)	0
block4_conv1 (Conv2D)	(None, 28, 28, 512)	1180160
block4_conv2 (Conv2D)	(None, 28, 28, 512)	2359808
block4_conv3 (Conv2D)	(None, 28, 28, 512)	2359808
block4_pool (MaxPooling2D)	(None, 14, 14, 512)	0
block5_conv1 (Conv2D)	(None, 14, 14, 512)	2359808
block5_conv2 (Conv2D)	(None, 14, 14, 512)	2359808
block5_conv3 (Conv2D)	(None, 14, 14, 512)	2359808
block5_pool (MaxPooling2D)	(None, 7, 7, 512)	0
flatten (Flatten)	(None, 25088)	0
dense (Dense)	(None, 2)	50178
=====		
Total params: 14,764,866		
Trainable params: 50,178		
Non-trainable params: 14,714,688		

```
In [9]: # compile model
model.compile(
    loss='categorical_crossentropy',
    optimizer='adam',
```

```
metrics=['accuracy']  
)
```

In [10]: *# Use the Image Data Generator to import the images from the dataset*
from keras.preprocessing.image **import** ImageDataGenerator

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

In [11]: *# Make sure you provide the same target size as initialied for the image size*
training_set = train_datagen.flow_from_directory('Datasets/train',
 target_size = (224, 224),
 batch_size = 32,
 class_mode = 'categorical')

Found 5216 images belonging to 2 classes.

In [12]: test_set = test_datagen.flow_from_directory('Datasets/test',
 target_size = (224, 224),
 batch_size = 32,
 class_mode = 'categorical')

Found 624 images belonging to 2 classes.

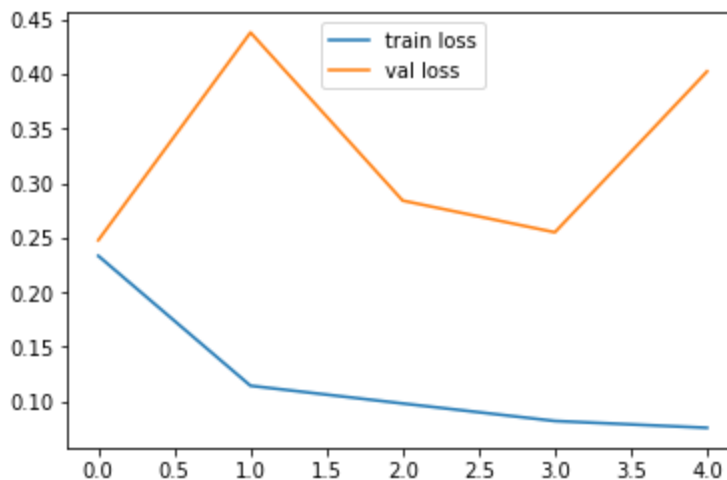
In [13]: *# fit the model*
Run the cell. It will take some time to execute
r = model.fit_generator(
 training_set,
 validation_data=test_set,
 epochs=5,
 steps_per_epoch=len(training_set),
 validation_steps=len(test_set)
)

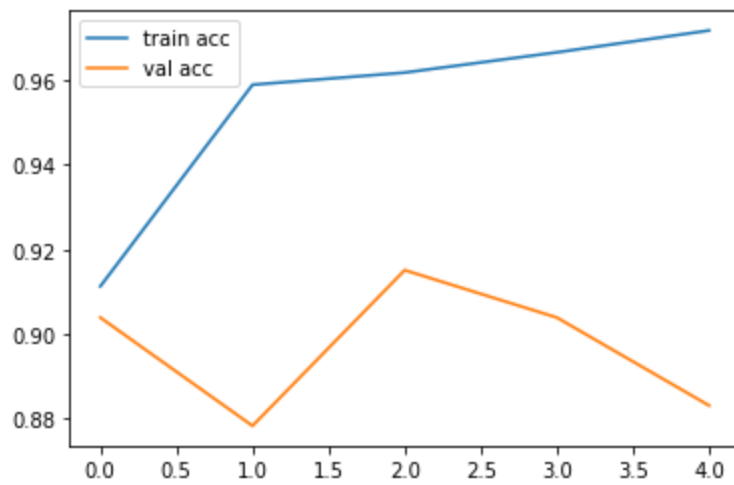
/Users/mybeast/opt/anaconda3/lib/python3.8/site-packages/keras/engine/training.py:1972: UserWarning: `Model.fit_generator` is deprecated and will be removed in a future version. Please use `Model.fit`, which supports generators.
warnings.warn("`Model.fit_generator` is deprecated and "

Epoch 1/5
163/163 [=====] - 3461s 21s/step - loss: 0.2333 - accuracy: 0.9112 - val_loss: 0.2477 - val_accuracy: 0.9038
Epoch 2/5
163/163 [=====] - 2407s 15s/step - loss: 0.1143 - accuracy: 0.9590 - val_loss: 0.4380 - val_accuracy: 0.8782
Epoch 3/5
163/163 [=====] - 2128s 13s/step - loss: 0.0981 - accuracy: 0.9618 - val_loss: 0.2839 - val_accuracy: 0.9151
Epoch 4/5
163/163 [=====] - 2122s 13s/step - loss: 0.0820 - accuracy: 0.9666 - val_loss: 0.2549 - val_accuracy: 0.9038
Epoch 5/5
163/163 [=====] - 2100s 13s/step - loss: 0.0758 - accuracy: 0.9718 - val_loss: 0.4023 - val_accuracy: 0.8830

```
In [17]: # plot the loss
plt.plot(r.history['loss'], label='train loss')
plt.plot(r.history['val_loss'], label='val loss')
plt.legend()
plt.show()
plt.savefig('LossVal_loss')

# plot the accuracy
plt.plot(r.history['accuracy'], label='train acc')
plt.plot(r.history['val_accuracy'], label='val acc')
plt.legend()
plt.show()
plt.savefig('AccVal_acc')
```





<Figure size 432x288 with 0 Axes>

```
In [15]: # save it as a h5 file

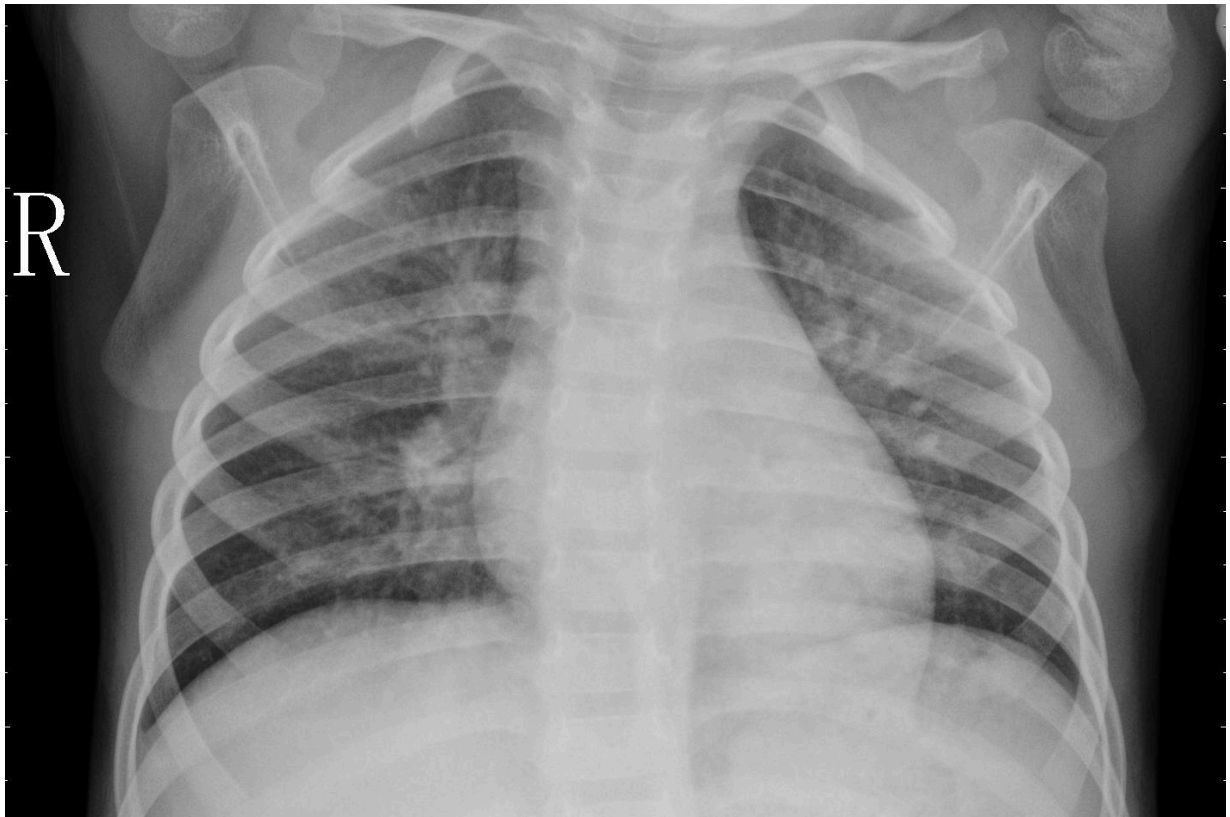
import tensorflow as tf

from keras.models import load_model

model.save('model_vgg16.h5')
```

```
In [80]: from IPython.display import display
from PIL import Image

NeumonialPath="Datasets/val/PNEUMONIA/person1949_bacteria_4880.jpeg"
# NormalPath = "Datasets/val/NORMAL/NORMAL2-IM-1431-0001.jpeg"
display(Image.open(NeumonialPath))
```



```
In [81]: from tensorflow.keras.models import load_model
import numpy as np

model = load_model('model_vgg16.h5')

img = image.load_img(NeumonialPath,target_size=(224, 224))

x = image.img_to_array(img)
x = np.expand_dims(x,axis=0)
img_data = preprocess_input(x)
classes = model.predict(img_data)
print(classes)
if classes[0][0] > classes[0][1]:
    print('X-Ray image is NORMAL')
else:
    print('X-Ray image is Having NEUMONIAL')

[[0. 1.]]
X-Ray image is Having NEUMONIAL
```

In []: