

# PNEUMONIA DETECTION

**USING MACHINE LEARNING** 

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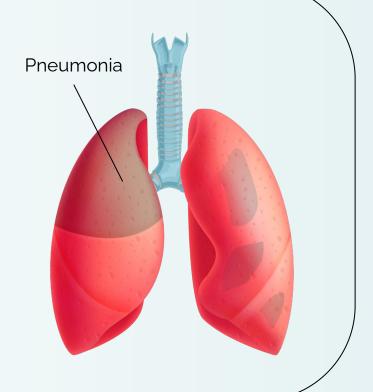
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Introduction to Pneumonia Detection



#### **Introduction to Pneumonia Detection**



#### What is Pneumonia?

A respiratory disease that affects the lungs, which could ultimately result in death.



#### How Do We Identify It?

One major sign of pneumonia is white spots in the lungs, which can be identified through image detection of chest X-rays.



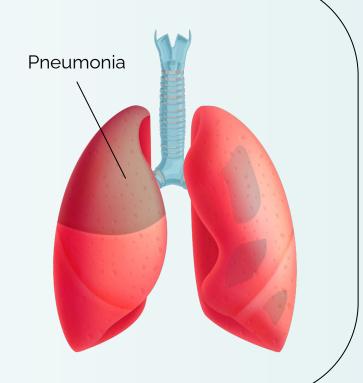
#### **Further Details**

X-rays work by passing radiation through the body, which can create an accurate image because the radiation passes through some materials more easily than others. The waves will pass right through air and water; in comparison, they will not penetrate through fat, tissue, or bone well.

Diagnosing pneumonia <u>early</u> is crucial, as it could significantly worsen if not detected.

02

**Modeling** 



## Train vs. Test Data, Healthy vs. Pneumonia Images

```
### YOUR CODE HERE
metadata.groupby(['split']).count()
sns.countplot(x = 'split', data = metadata)
### END CODE
<matplotlib.axes. subplots.AxesSubplot at 0x7ff3f46430d0>
   2000
  1750
   1500
  1250
1000
   750
    500
   250
                 train
                                         test
                             split
```

```
[16] # grab our seaborn visualization toolbox!
     import seaborn as sns
     ### YOUR CODE HERE
     metadata.groupby(['class']).count()
     sns.countplot(x = 'class', data = metadata)
     ### END CODE
     <matplotlib.axes._subplots.AxesSubplot at 0x7ff3f46e3650>
        1200
        1000
         800
        600
         400
         200
                       0.0
                                             1.0
                                 dass
```

## MLP & Dense Neural Networks

```
#YOUR CODE HERE
from sklearn.neural_network import MLPClassifier

(train_data, train_labels) = get_train_data(flatten = True)
  (test_data, test_labels) = get_test_data(flatten = True)

model = MLPClassifier(hidden_layer_sizes=(5))

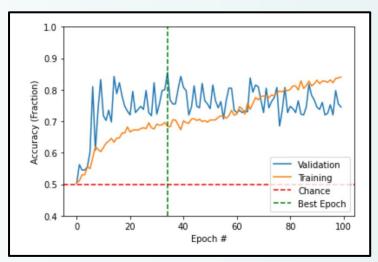
model.fit(train_data, train_labels)

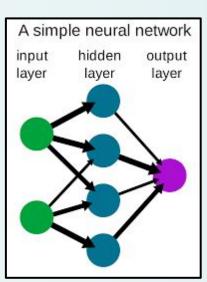
pred = model.predict(test_data)

accuracy = accuracy_score(test_labels, pred)
print(accuracy)

#YOUR CODE HERE

0.5
```

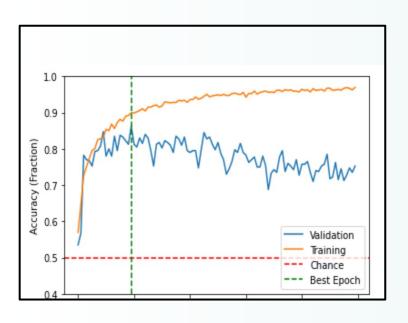


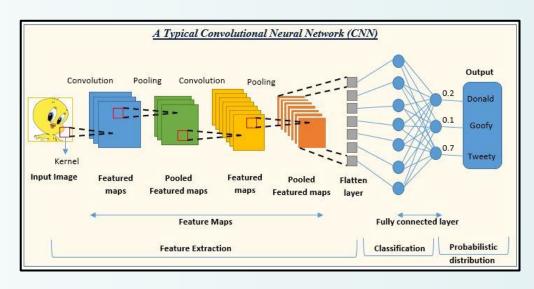


MLP Classifier Neural Network Code

Dense Neural Network Accuracy Graph

## **CNNs (Convolutional Neural Networks)**





Input Layer: Images of lungs

Convolutional Layer: 64 x 64 layers (3 layers)

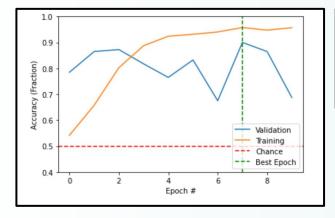
## **Transfer Learning Model (VGG16)**

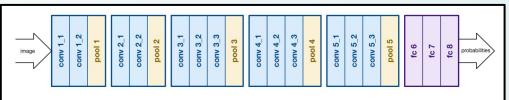
```
[14] ### YOUR CODE HERE
    train_data, train_labels = get_train_data()
    test_data, test_labels = get_test_data()

    transfer = TransferClassifier(name = 'VGG16')

    transfer.fit(train_data, train_labels, epochs = 10, validation_data = (test_data, test_labels), shuffle = True, callbacks
    plot_acc(transfer.history)

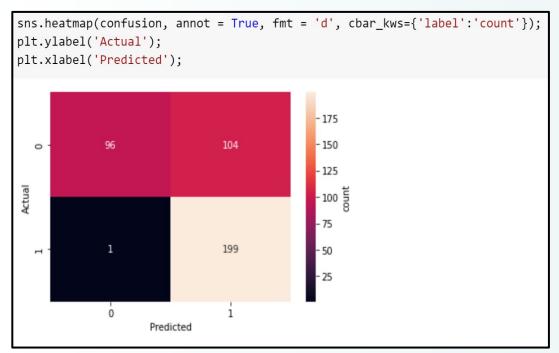
### END CODE
```

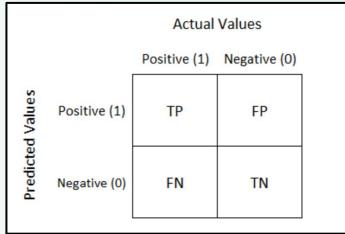




I decided to train the data using a new, expert CNN model (VGG16). This particular model has experience with over 14 million images, so it should be useful when distinguishing between healthy and pneumonia x-rays.

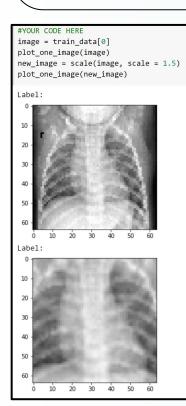
### **Confusion Matrix**

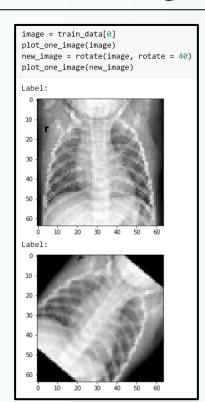


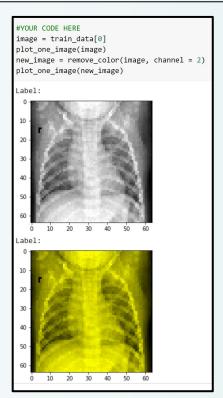


The confusion matrix is a way to evaluate the performance of the model by analyzing the number of true and false positives and negatives. In this matrix, I analyzed the Transfer Learning VGG16 Model.

## Augmentation







In order to allow the CNN model to handle various forms of scans. I trained it using augmented data. This augmentation improves the model and allows it to create more accurate predictions by exposing the model to "imperfect" scans. Imperfect scans mean tilted, rotated, or discolored.

# **Application (Field Data in CNNs)**

```
train_data, train_labels = get_train_data()
test_data, test_labels = get_test_data()
field_data, field_labels = get_field_data()

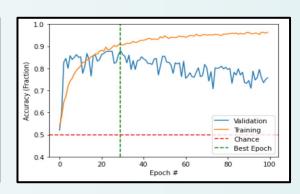
### YOUR CODE HERE

cnnnew = CNNClassifier(num_hidden_layers = 2)

cnnnew_history = cnnnew.fit(train_data, train_labels, epochs = 100, validation_data = (test_data, test_labels), shuffle = True, callbacks = [monitor])
plot_acc(cnnnew_history)

y_pred = (cnnnew.predict(field_data) > 0.5).astype("int32")
print(accuracy_score(field_labels, y_pred))

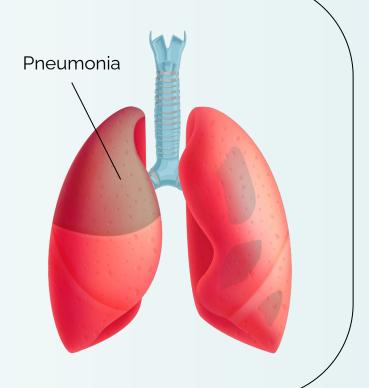
### END CODE
```



Here, I incorporated field data into the new CNN model. Though it didn't impact the model's accuracy greatly, the field data introduced it model to a larger dataset. This made the model more applicable to the real world.

03

**ML App Deployment** 



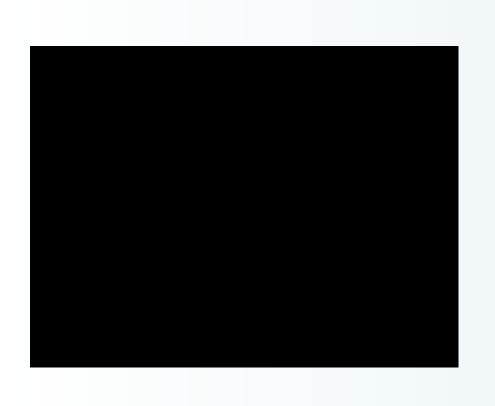
# Model Deployment



- Website using local Streamlit server (in development)
- Coded with Python
- Takes picture input and outputs "Healthy" or "Pneumonia"
- Possible to update to include TB

```
#@title Run this to host our website! { display-mode: "form"
3 app = Flask( name )
4 run with ngrok(app)
5 CORS(app)
7 simple webpage = "simple-webpage.html"
8 simple html = codecs.open(simple_webpage, 'r').read()
10 @app.route("/")
11 def home():
      return simple html
14 app.run()
```

# Streamlit - Model Code



# Old HTML Server Demo

