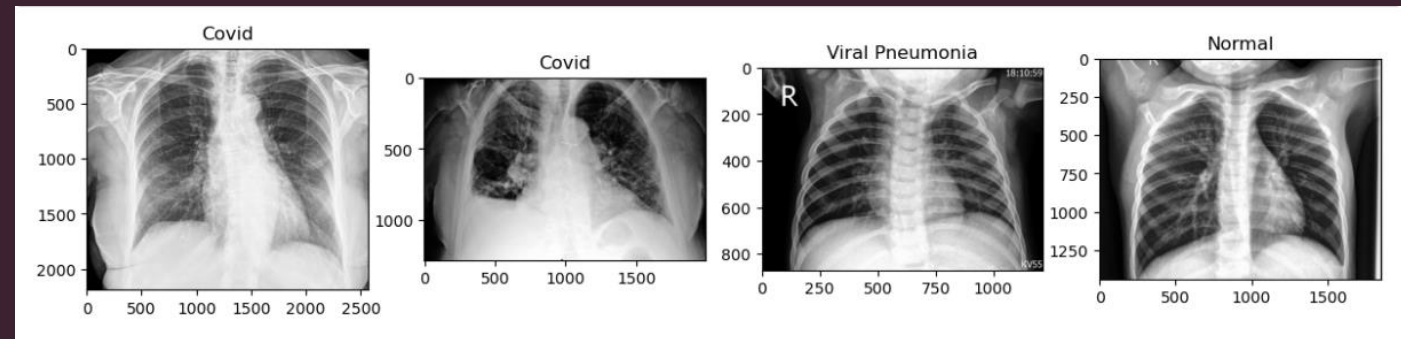


# Covid-19 chest Xray classification

Using machine learning and  
Deep Learning Methods



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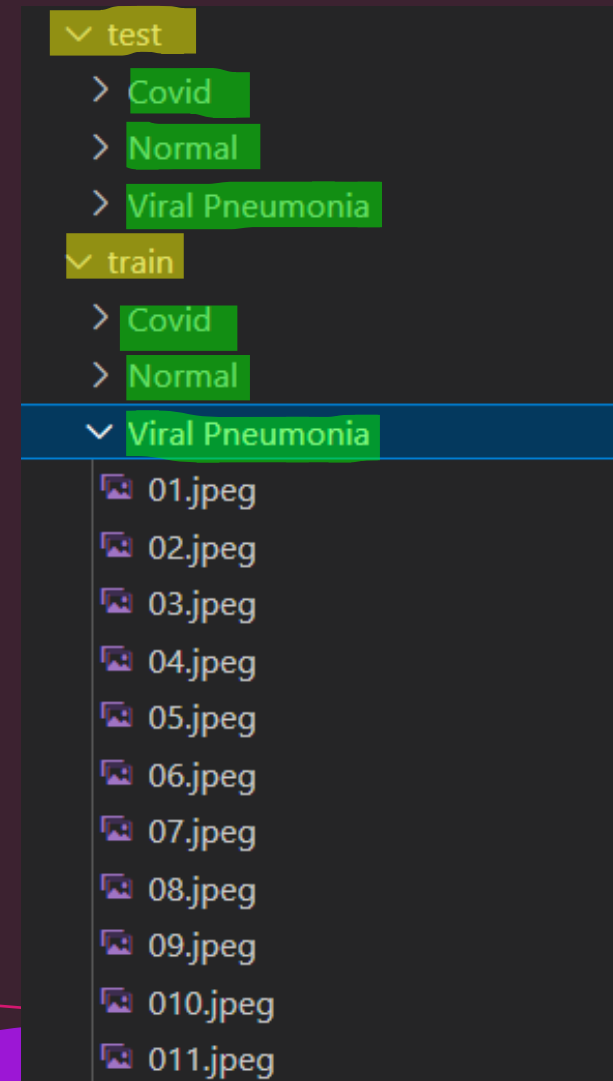
# INTRODUCTION

- This project focuses on classifying chest X-ray images into three categories:
  1. COVID-19
  2. viral pneumonia
  3. healthy people
- Chest X-ray classification plays a vital role in diagnosing respiratory diseases, especially during pandemics like COVID-19.
- This project aims to develop a reliable CNN model that will overcome ML model scores and will be capable of accurately categorizing chest X-ray images into relevant classes.

# GETTING KNOW THE DATA

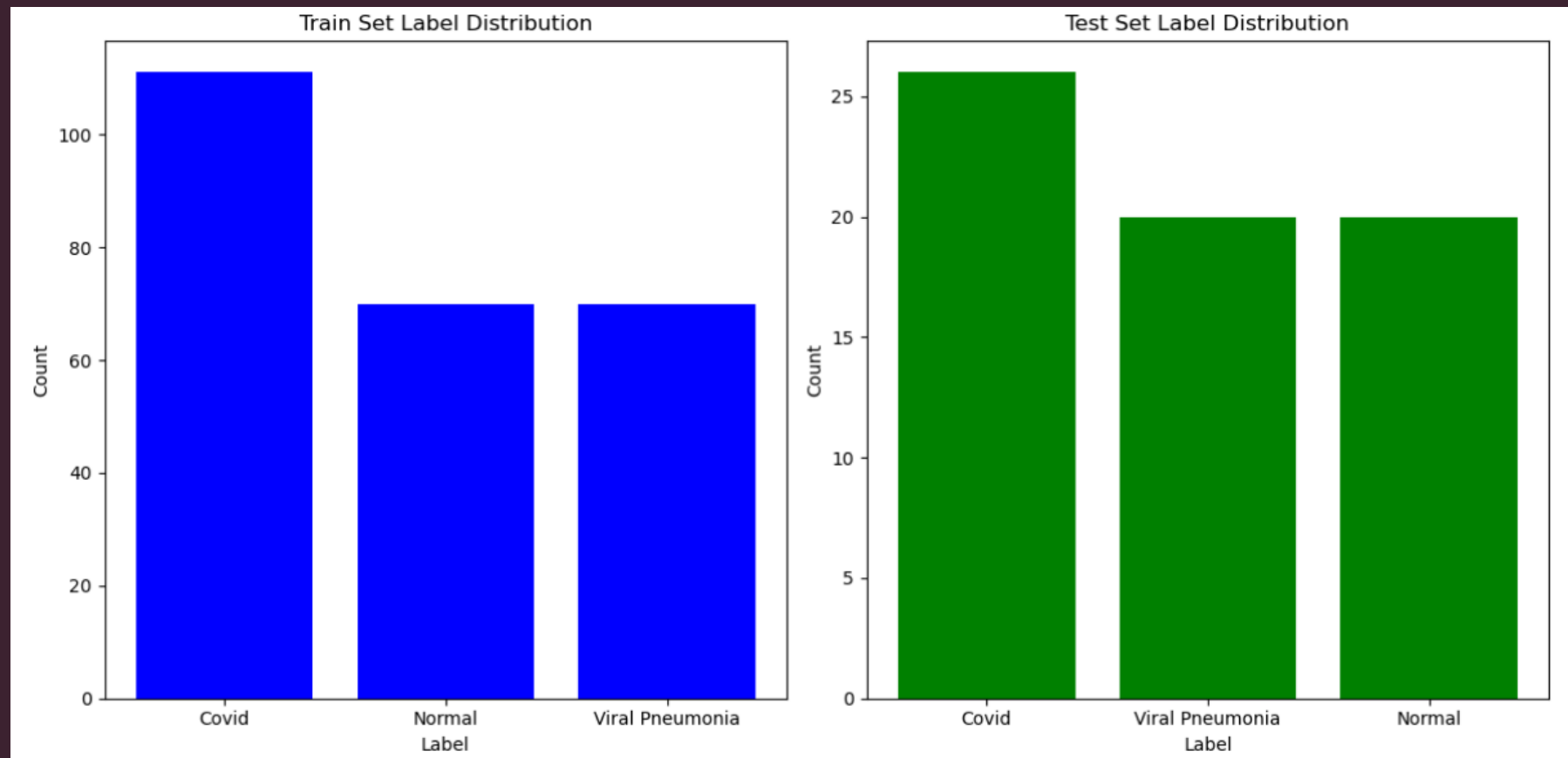
- This Data is organized in 2 folders which are separated to its label into different folders for each cluster.
- We extracted the images by iterating the folders and append it to lists.

```
def preprocess_data():  
    # Load and preprocess train and test data  
    train_dir = 'train'  
    test_dir = 'test'  
  
    train_filepaths = []  
    train_labels = []  
    for label in os.listdir(train_dir):  
        label_dir = os.path.join(train_dir, label)  
        for filename in os.listdir(label_dir):  
            filepath = os.path.join(label_dir, filename)  
            train_filepaths.append(filepath)  
            train_labels.append(label)  
  
    test_filepaths = []  
    test_labels = []  
    for label in os.listdir(test_dir):  
        label_dir = os.path.join(test_dir, label)  
        for filename in os.listdir(label_dir):  
            filepath = os.path.join(label_dir, filename)  
            test_filepaths.append(filepath)  
            test_labels.append(label)
```



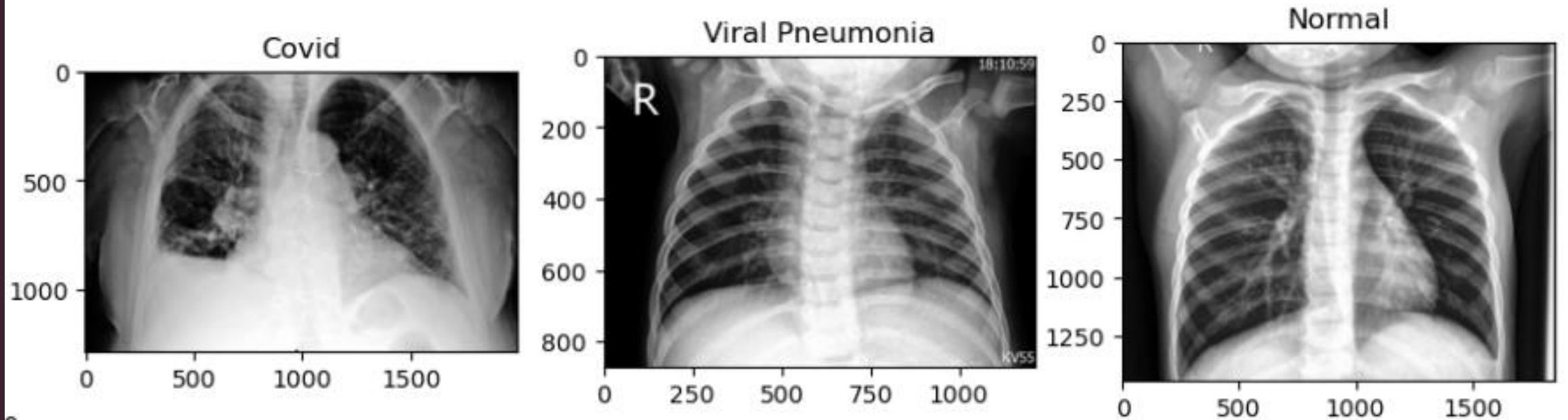
# GETTING KNOW THE DATA

- Although The data is a bit biased to the “Covid” cluster, we have enough data such that the model will be able to learn all the labels well



# GETTING KNOW THE DATA

- In addition we noticed the sizes of the images are different



# PRE-PROCESSING

- During the loading of the images, we implemented a preprocessing function to convert all the images to the same size and the same scale

```
def preprocess_image(image_path, target_size=(256, 256)):
    img = cv2.imread(image_path)
    img_resized = cv2.resize(img, target_size)
    img_normalized = img_resized.astype("float32") / 255.0
    return img_normalized
```

- Afterwards we separated them into Numpy correspondence arrays

```
x_train = np.array([preprocess_image(filepath) for filepath in train_filepaths])
y_train = np.array(train_labels)
x_test = np.array([preprocess_image(filepath) for filepath in test_filepaths])
y_test = np.array(test_labels)

# Convert labels to numerical values
label_to_index = {label: i for i, label in enumerate(np.unique(y_train))}
y_train = np.array([label_to_index[label] for label in y_train])
y_test = np.array([label_to_index[label] for label in y_test])

return x_train, y_train, x_test, y_test, train_filepaths, train_labels, test_filepaths, test_labels
```

# PRE-PROCESSING

- And made sure the training data would shuffle

```
x_train, y_train, x_test, y_test, train_filepaths, train_labels, test_filepaths, test_labels = preprocess_data()

# Shuffle indices
shuffled_indices = np.random.permutation(len(x_train))

# Shuffle x_train and y_train accordingly
x_train = x_train[shuffled_indices]
y_train = y_train[shuffled_indices]
```

- Creating a Pandas Data Frames – (251 samples in the training set, and 66 in test set)

	filepath	label
0	test\Covid\0100.jpeg	Covid
1	test\Covid\0102.jpeg	Covid
2	test\Covid\0105.png	Covid
3	test\Covid\0106.jpeg	Covid
4	test\Covid\0108.jpeg	Covid

	filepath	label
0	train\Covid\01.jpeg	Covid
1	train\Covid\010.png	Covid
2	train\Covid\012.jpeg	Covid
3	train\Covid\015.jpg	Covid
4	train\Covid\019.png	Covid

```
Num of samples in train set: 200
Num of samples in validation set: 51
Num of samples in test set: 66
```

- We also split the training set into a validation set

# SOFTMAX MODEL

- We prefer to use Softmax which is commonly preferred when dealing with multi-class classification problems.
- Softmax assumes your classes are mutually exclusive. This means a data point can only belong to one class at a time.
- In addition, Softmax outputs a probability distribution for each class. The sum of these probabilities for all classes will always be 1. This makes it ideal for scenarios where you want the model to predict the most likely class along with its confidence level



# SOFTMAX MODEL

- We developed our SoftmaxModel class using TensorFlow and Keras. After experimenting with various configurations, we achieved optimal performance with a learning rate of 0.001 and the Adam optimizer

```
Epoch 48/50
7/7 [=====] - 0s 38ms/step - loss: 2.1556e-06 - accuracy: 1.0000 - val_loss: 0.7499 - val_accuracy: 0.9608
Epoch 49/50
7/7 [=====] - 0s 39ms/step - loss: 2.0829e-06 - accuracy: 1.0000 - val_loss: 0.7493 - val_accuracy: 0.9608
Epoch 50/50
7/7 [=====] - 0s 34ms/step - loss: 2.0239e-06 - accuracy: 1.0000 - val_loss: 0.7487 - val_accuracy: 0.9608
3/3 [=====] - 0s 17ms/step - loss: 2.2056 - accuracy: 0.8939
Test Accuracy: 0.8939393758773804
```

# ‘ADAM’ OPTIMIZER

- Adam (Adaptive Moment Estimation) helps minimize the loss function by iteratively adjusting the learning rate for each parameter in the model based on past gradients by fast convergence (if the loss is high – the adjustment in the learning rate will be much more significant than if the loss is low)

# SIMPLE NN MODEL

- Using TensorFlow and Keras, we built a simple Neural network of Sequential model, with ReLU activation function.

```
model = Sequential([
    Flatten(input_shape=(256, 256, 3)), # Flatten input images
    Dense(256, activation='relu'), # First dense layer with ReLU activation
    Dense(128, activation='relu'), # Second dense layer with ReLU activation
    Dense(64, activation='relu'), # third dense layer with ReLU activation
    Dense(num_classes, activation='softmax') # Output layer with softmax activation
])
```

- We compiled the model with the previous configurations, and fit the model with 50 epochs, with batch size of 32

```
Epoch 48/50
7/7 [=====] - 2s 286ms/step - loss: 0.0859 - accuracy: 0.9900 - val_loss: 0.5557 - val_accuracy:
0.9608
Epoch 49/50
7/7 [=====] - 2s 295ms/step - loss: 0.0427 - accuracy: 0.9950 - val_loss: 0.5248 - val_accuracy:
0.9608
Epoch 50/50
7/7 [=====] - 2s 288ms/step - loss: 0.0011 - accuracy: 1.0000 - val_loss: 1.0230 - val_accuracy:
0.9412
3/3 [=====] - 0s 52ms/step - loss: 2.2867 - accuracy: 0.8485
Test Accuracy: 0.8484848737716675
```

# SIMPLE NN MODEL

- The current model's performance might be suffering because it's a fully connected neural network with several dense layers. This architecture might be too complex for this specific task, especially if the images contain a lot of irrelevant pixels.

# CNN MODEL

- Using a convolutional neural network, we achieved a much better score
- We convolved the data and reduced its dimension by MaxPooling2D and the dropout method
- This process required a few attempts and adjustments with the parameters, such that in the final stages we could use fully connected NN.
- We compiled the model with the same previous configuration (Adam optimizer, loss - categorical\_crossentropy, and accuracy metric)

Epoch 10/10

13/13 [=====] - 17s 1s/step - loss: 0.0037 - accuracy: 1.0000

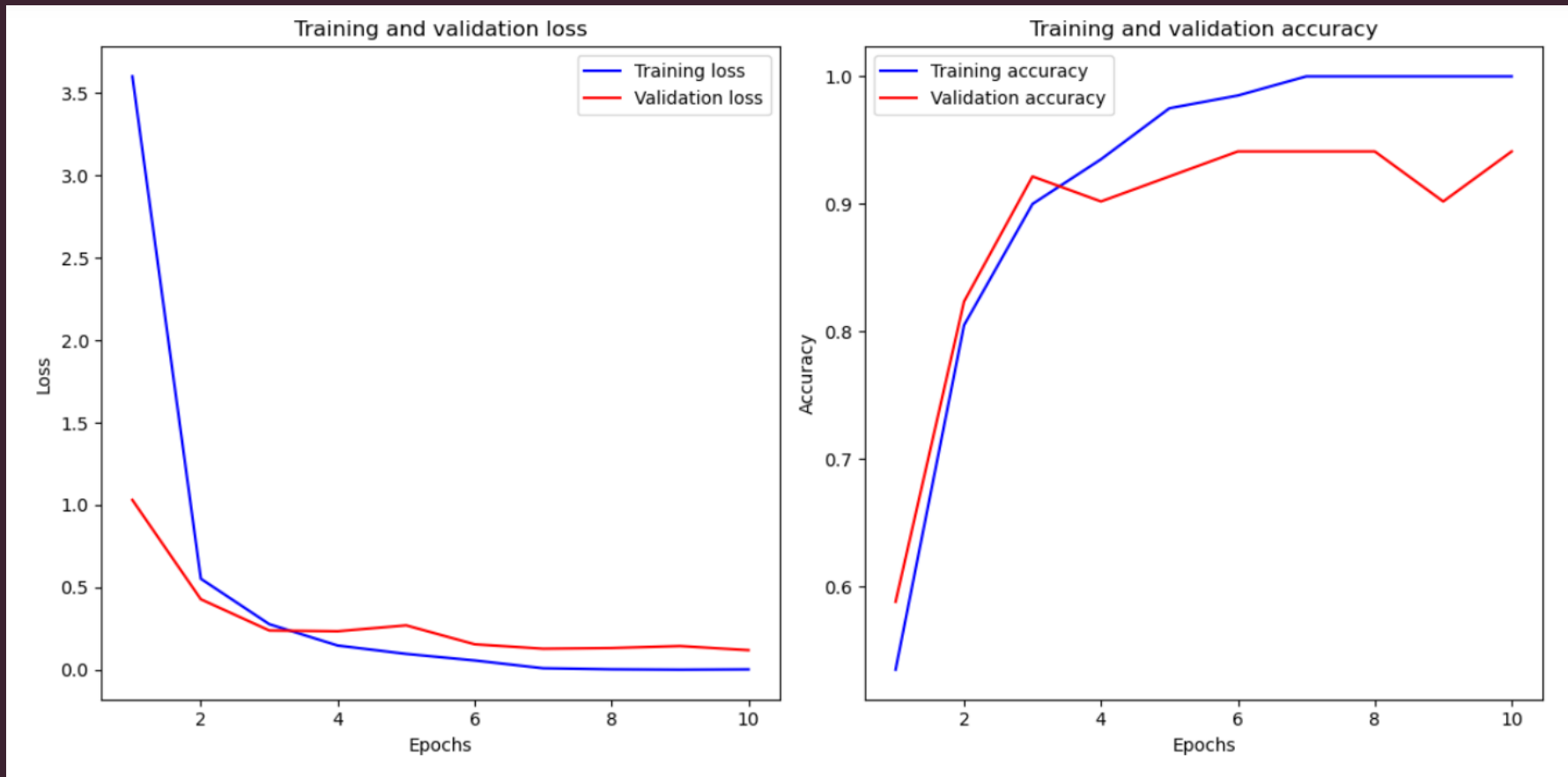
9412

3/3 [=====] - 1s 299ms/step - loss: 0.1114 - accuracy: 0.9545

Test Accuracy: 0.9545454382896423

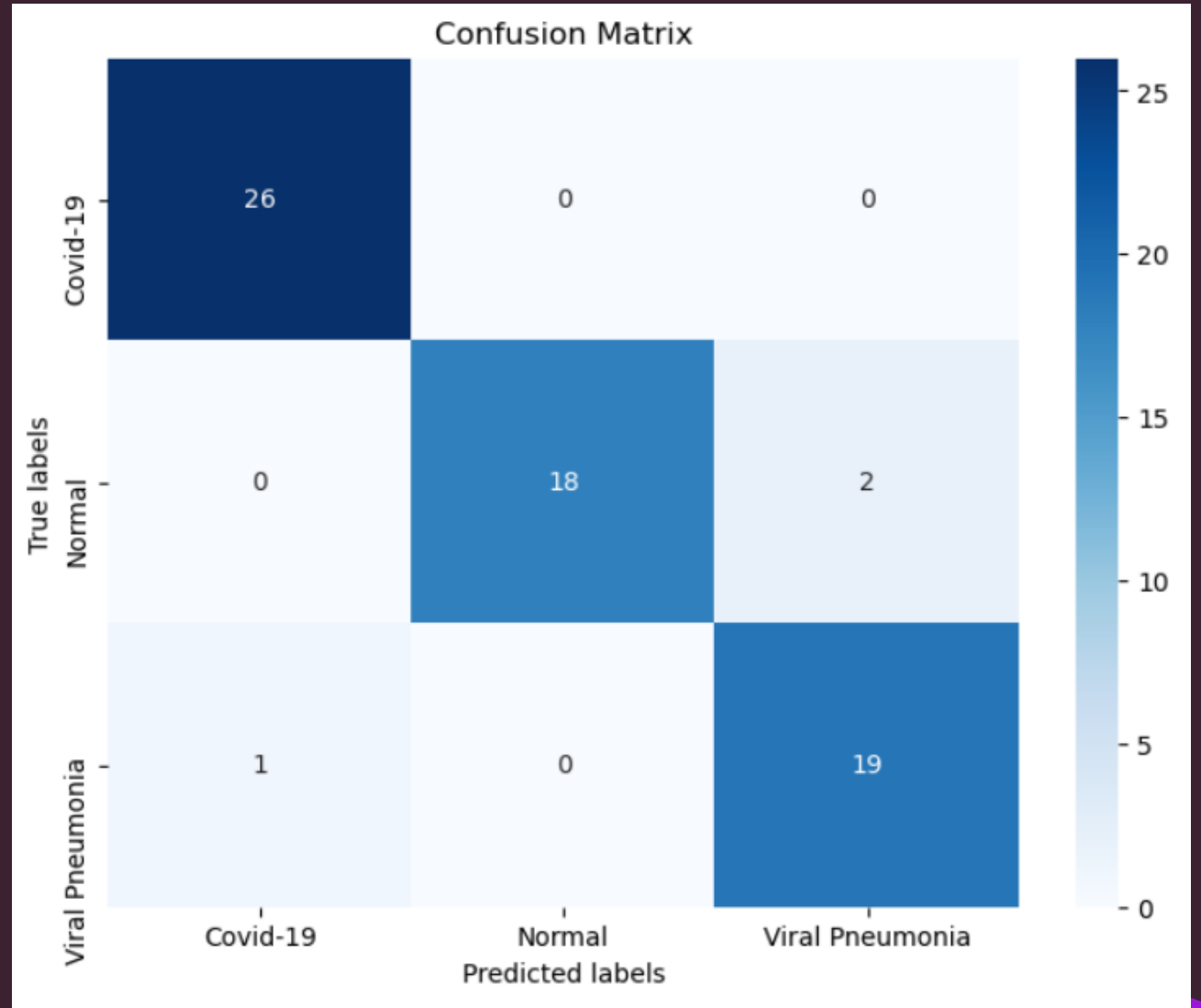
# CNN MODEL

- Plotting the accuracy score relative to the loss score
- We would expect the loss and accuracy of the training and the validation sets to go together, and as we can see - it is.



# CONFUSION MATRIX

- Following this confusion matrix we can see that the model predicted correctly 63 from 66 samples, with the following mistakes



# METRICS COMPARING

- There are several considerations in the direction of the error (e.g. in cases of false negative/false negative)
- We wanted to see the results of the model also with the F1-score or Precision indices, which would be the recommended indices since it is important to identify positive cases accurately To be able to provide the necessary medical treatment.



# METRICS COMPARING

- As you can see when it comes to a COVID-19 patient there is even higher accuracy, which indicates the reliability and suitability of the model for the desired case

Covid-19:

Precision: 0.9629629629629629

Recall: 1.0

F1 Score: 0.9811320754716981

Normal:

Precision: 1.0

Recall: 0.9

F1 Score: 0.9473684210526316

Viral Pneumonia:

Precision: 0.9047619047619048

Recall: 0.95

F1 Score: 0.9268292682926829

# DATA AUGMENTATION

- In an attempt to improve the model's score, we implemented data augmentation techniques such as rescaling, rotation, shear, and zoom. This was motivated by the fact that the current dataset primarily consists of frontal chest X-ray images with similar appearances.
- While training with these augmented samples didn't lead to a performance increase, it's reassuring to know that, according to our understanding, X-ray institutes capture chest X-rays in a standardized position and orientation.
- This suggests that the original dataset, despite its lack of variety, possesses sufficient reliability.

# CONCLUSIONS

- As you can see, the use of a convolutional network in our model provides an almost complete prediction of Covid-19 among patients who came for a chest x-ray.
- In this problem, and in many other problems, just as it is necessary to deepen the level of learning to several layers, it is necessary not to go too deep, thus not only to burden the complexity of the model, but inclining it to learn of data that does not help to solve the problem, and to use simple methods to reduce the dimensions required for the problem.