A detailed 3D rendering of a COVID-19 virus particle, showing its characteristic spike proteins and internal structure, set against a dark blue background.

Computer Vision

## **COVID-19-DIAGNOSE-USING-LUNGS-X-RAYS**

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# OUTLINE

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Phase 1: Easy Round - **CNN MODEL**

Phase 2: Medium Round - **CNN MODEL-PRE\_TRAINED**

Phase 3: Hard Round-**Integration on web**

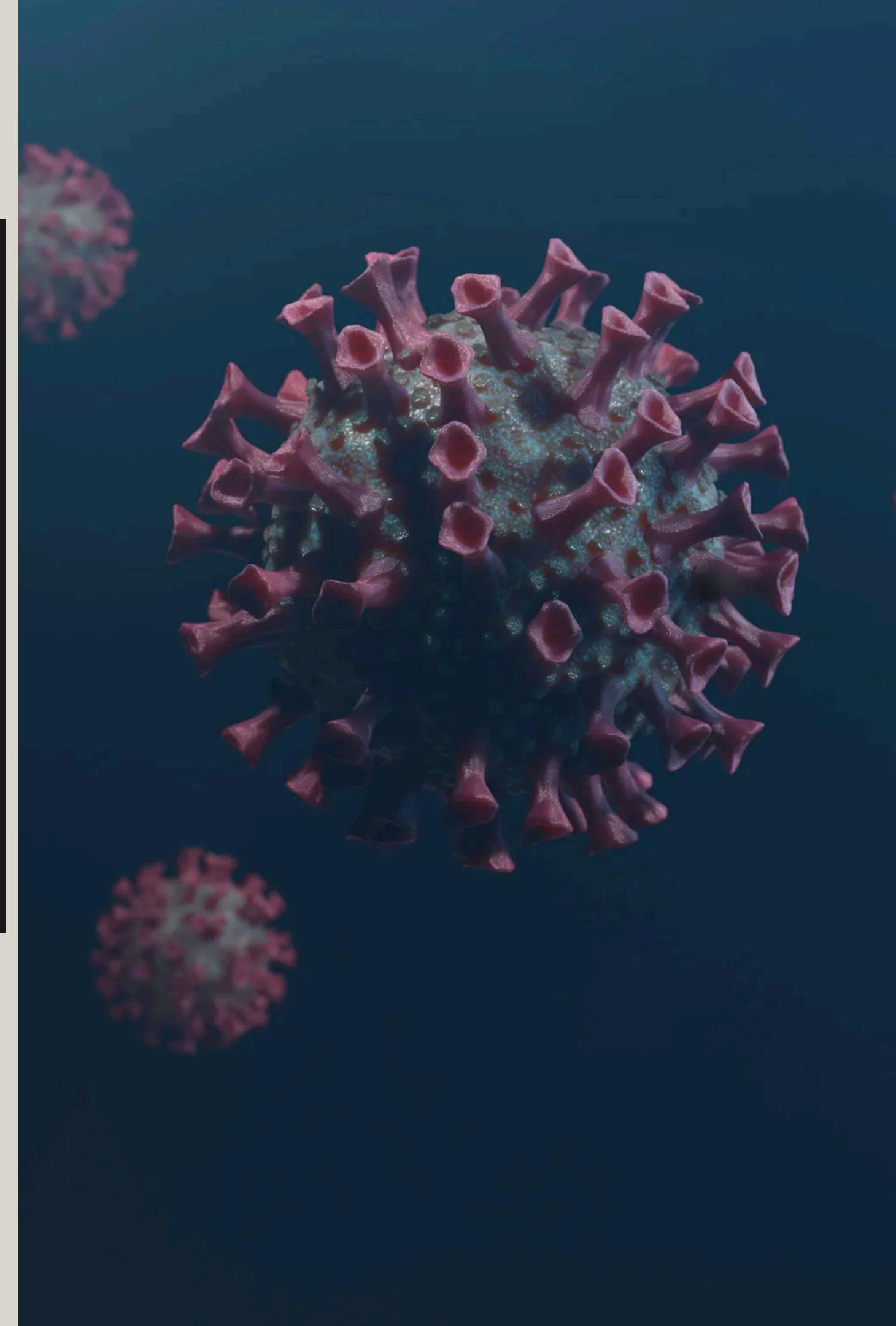


# PHASE 1: CNN-MODEL

# Libraries

---

```
from keras.models import Sequential  
from keras.layers import Convolution2D  
from keras.layers import MaxPool2D  
from keras.layers import Flatten  
from keras.layers import Dense  
import matplotlib.pyplot as plt  
import numpy as np  
from tensorflow.keras.preprocessing import image
```



# preprocessing

---

testing

```
testing_data_generator = ImageDataGenerator(  
    rescale=1.0/255  
)
```

tranining

```
✓ training_data_generator = ImageDataGenerator(  
    Click to collapse the range.  
    shear_range=0.2,  
    zoom_range=0.2,  
    horizontal_flip=True  
)
```

```
training_dataset = training_data_generator.flow_from_directory(  
    'Covid-19/train',  
    target_size=(64,64),  
    batch_size=32,  
    class_mode='categorical'  
)  
]
```

Python

Found 251 images belonging to 3 classes.

```
testing_dataset = testing_data_generator.flow_from_directory(  
    'Covid-19/test',  
    target_size=(64,64),  
    batch_size=32,  
    class_mode='categorical'  
)  
]
```

Python

Found 66 images belonging to 3 classes.

# DATASET

Training Images

Class: 0



Class: 0

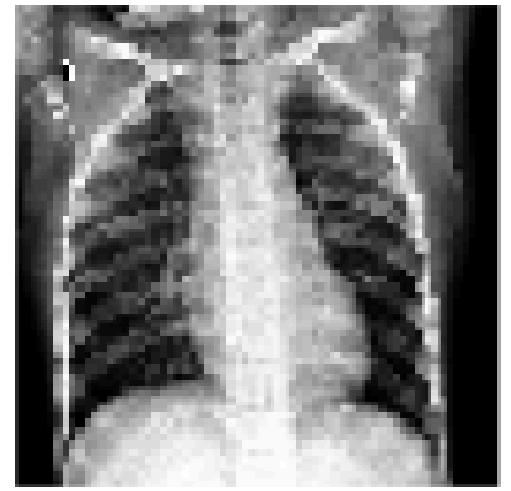


Class: 1

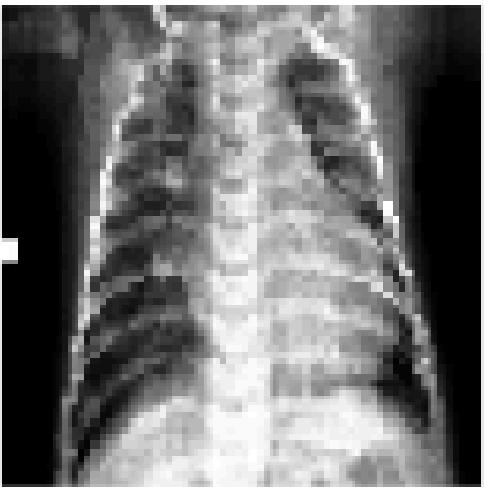


Testing Images

Class: 1



Class: 2



Class: 2



# MODEL

## CNN

```
classifier = Sequential()
classifier.add(Convolution2D(32,(3,3),input_shape=(64,64,3), activation='relu'))
classifier.add(MaxPool2D(pool_size=(2,2)))

classifier.add(Convolution2D(16,(3,3), activation='relu'))
classifier.add(MaxPool2D(pool_size=(2,2)))

classifier.add(Flatten())

classifier.add(Dense(units= 128, activation='relu'))
classifier.add(Dense(units= 3, activation='softmax'))

classifier.compile(optimizer='adam', loss='categorical_crossentropy', metrics=[ 'accuracy'])
```

# SUMMARY

Layer (type)	Output Shape	Param #
conv2d_6 (Conv2D)	(None, 62, 62, 32)	896
max_pooling2d_6 (MaxPooling2D)	(None, 31, 31, 32)	0
conv2d_7 (Conv2D)	(None, 29, 29, 16)	4,624
max_pooling2d_7 (MaxPooling2D)	(None, 14, 14, 16)	0
flatten_3 (Flatten)	(None, 3136)	0
dense_6 (Dense)	(None, 128)	401,536
dense_7 (Dense)	(None, 3)	387

Total params: 407,443 (1.55 MB)

Trainable params: 407,443 (1.55 MB)

Non-trainable params: 0 (0.00 B)

# TRAINING

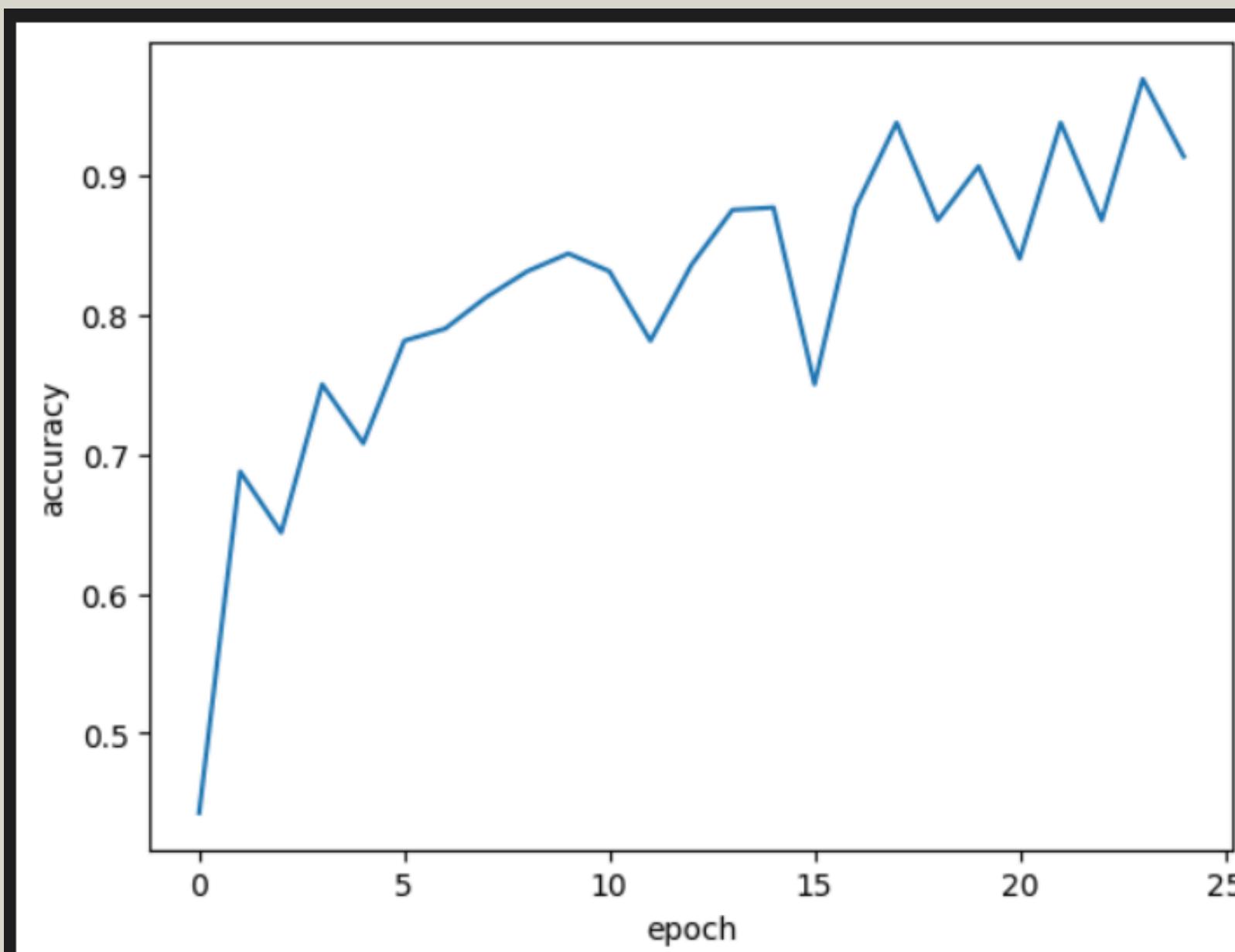
```
history = classifier.fit(  
    training_dataset,  
    steps_per_epoch=int(251/32),  
    epochs=25,  
    validation_data=testing_dataset,  
    validation_steps= int(66/32)  
)
```

Accuracy:

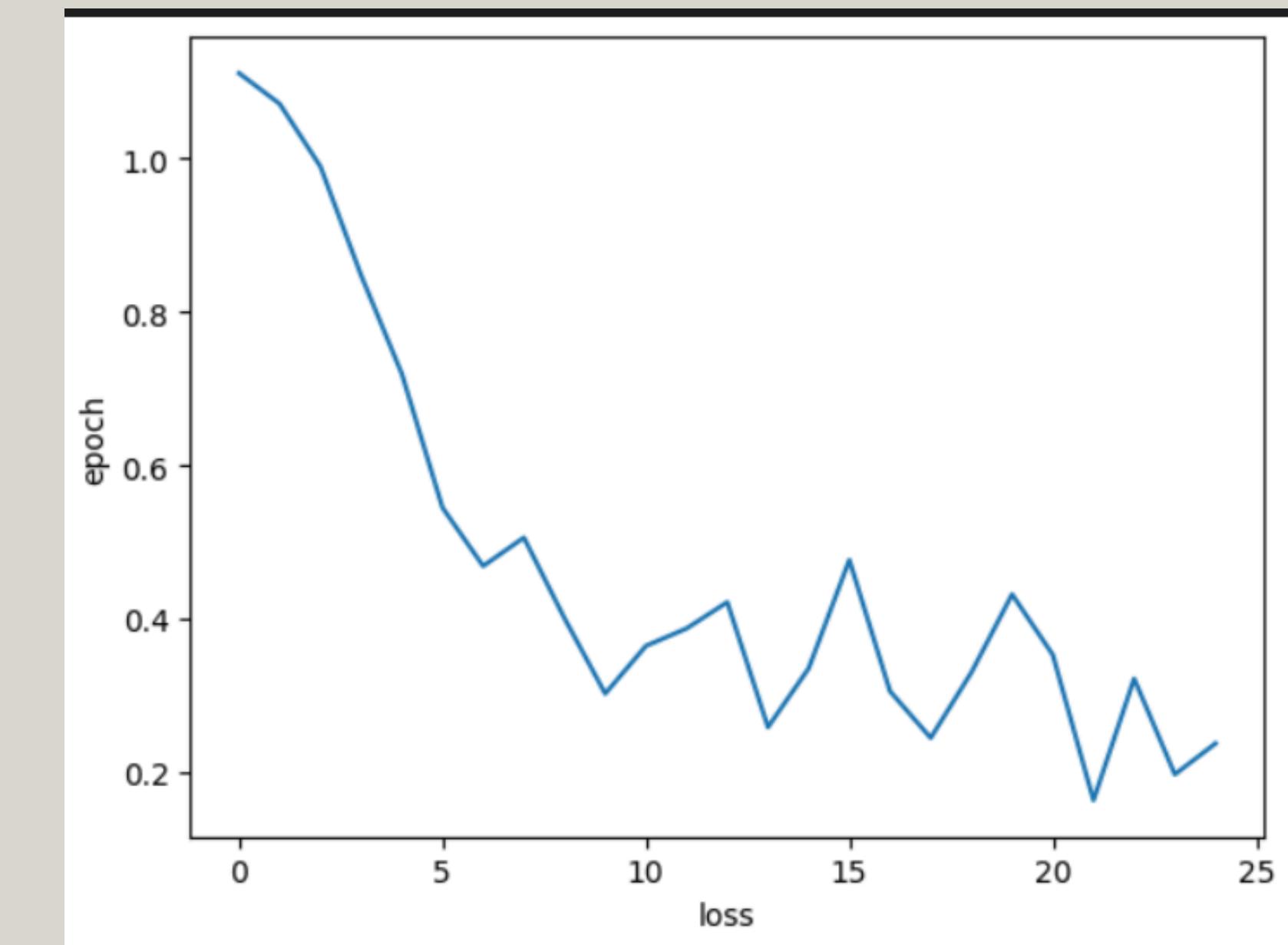
```
Training Accuracy: 0.9083665609359741  
Test Accuracy: 0.9242424368858337
```



# Visualization



validation accuracy



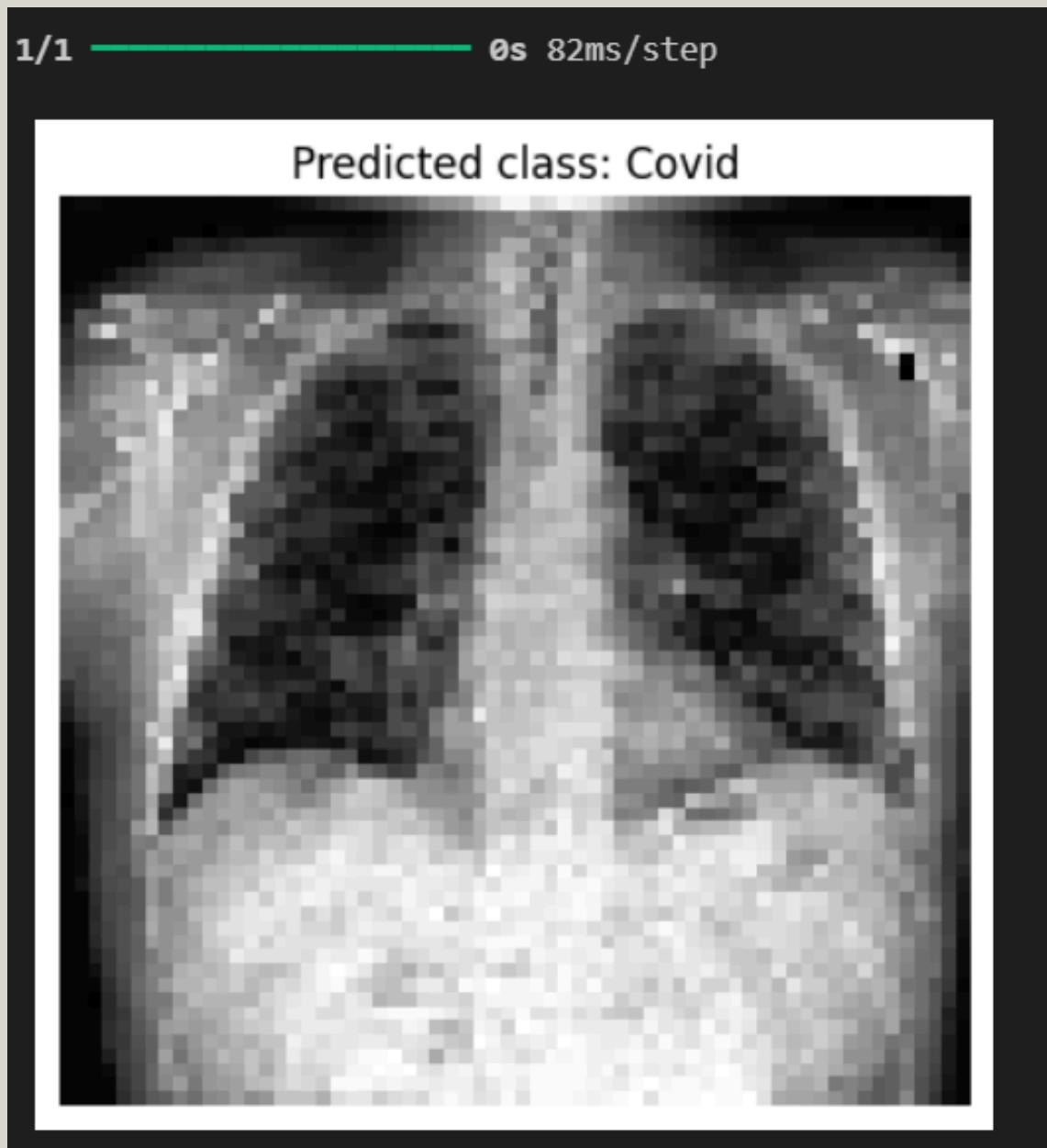
validation loss

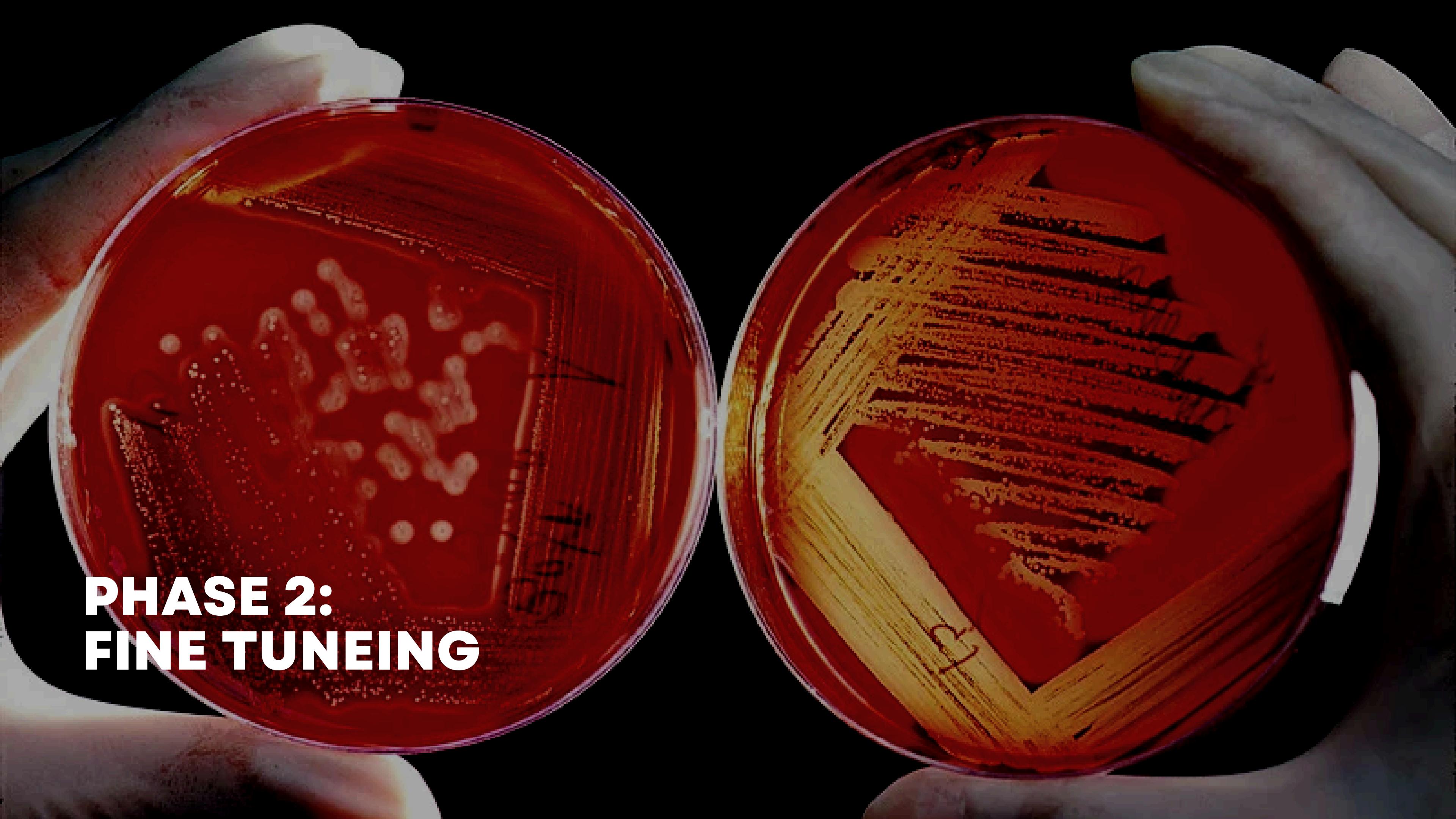
# prediction

```
img_path = r'Covid-19\test\Covid\0100.jpeg'
img = image.load_img(img_path, target_size=(64, 64))
img_array = image.img_to_array(img)
img_array = np.expand_dims(img_array, axis=0)
img_array = img_array / 255.0 # Normalize the image as per the

# Predict the class label using the ResNet50 model
prediction_cnn = classifier.predict(img_array)
predicted_class_cnn = np.argmax(prediction_cnn)
class_labels = ["Covid", "Normal", "Viral Pneumonia"]
predicted_label_resnet50 = class_labels[predicted_class_cnn]

# Display the image
plt.imshow(img)
plt.axis('off') # Hide axes
plt.title("Predicted class: " + predicted_label_resnet50) #
plt.show()
```





A close-up photograph of two petri dishes containing bacterial cultures on agar media. The dish on the left shows a dense, irregular cluster of white bacterial colonies. The dish on the right shows a more organized, linear arrangement of colonies, suggesting a different stage or condition of growth. Both dishes are set against a dark background.

## PHASE 2: FINE TUNEING

# Libraries

---

```
from tensorflow.keras.models import Sequential  
from tensorflow.keras.models import Model  
from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense  
from tensorflow.keras.applications import ResNet50  
from tensorflow.keras.applications import VGG16  
from tensorflow.keras.preprocessing.image import ImageDataGenerator  
from tensorflow.keras.preprocessing import image  
from sklearn.metrics import precision_score, recall_score, f1_score  
import matplotlib.pyplot as plt  
import numpy as np
```

# FINE TUNING

```
# Define your CNN architecture
model = Sequential()
model.add(Conv2D(32, (3, 3), activation='relu', input_shape=(64, 64, 3)))
model.add(MaxPooling2D((2, 2)))
model.add(Conv2D(64, (3, 3), activation='relu'))
model.add(MaxPooling2D((2, 2)))
model.add(Conv2D(128, (3, 3), activation='relu'))
model.add(MaxPooling2D((2, 2)))
model.add(Flatten())
model.add(Dense(128, activation='relu'))
model.add(Dense(3, activation='softmax'))
```

## RESNET\_50

```
# Load the ResNet50 model with pre-trained weights, excluding the top layer
base_model_resnet50 = ResNet50(weights='imagenet', include_top=False, input_shape

x = base_model_resnet50.output
x = Flatten()(x)
x = Dense(128, activation='relu')(x)
predictions = Dense(3, activation='softmax')(x)

model_resnet50 = Model(inputs=base_model_resnet50.input, outputs=predictions)

for layer in base_model_resnet50.layers:
    layer.trainable = False
```

Python

+ Code + Markdown

## VVG16

```
base_model_vgg16 = VGG16(weights='imagenet', include_top=False, input_shape=(64,

x = base_model_vgg16.output
x = Flatten()(x)
x = Dense(128, activation='relu')(x)
predictions_vgg16 = Dense(3, activation='softmax')(x)

model_vgg16 = Model(inputs=base_model_vgg16.input, outputs=predictions_vgg16)

for layer in base_model_vgg16.layers:
    layer.trainable = False
```

Python

# RESULTS

## RESNET\_50

FINE TUNED\_cnn

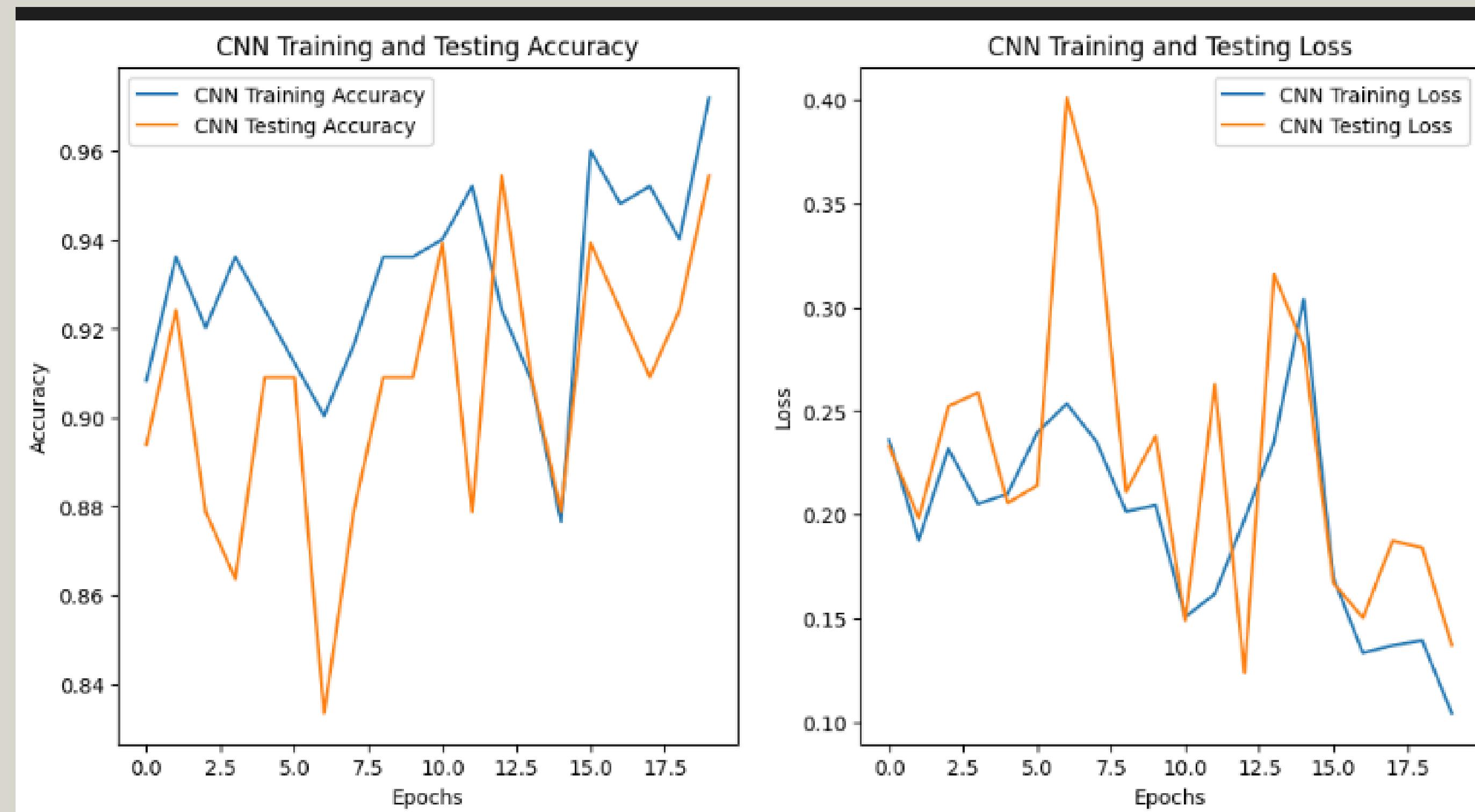
Training Accuracy: 0.9442231059074402  
Test Accuracy: 0.9545454382896423

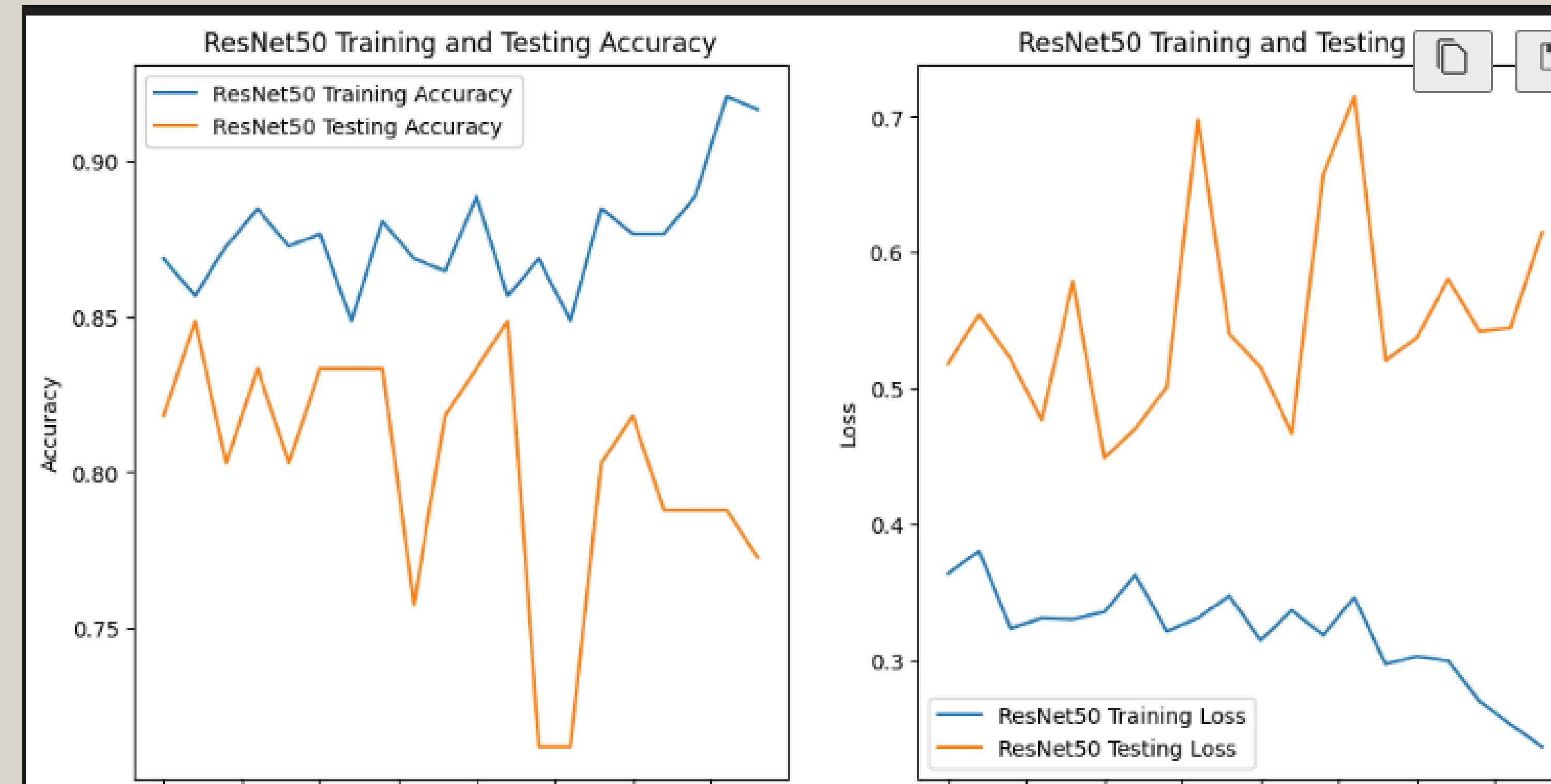
VGG\_16

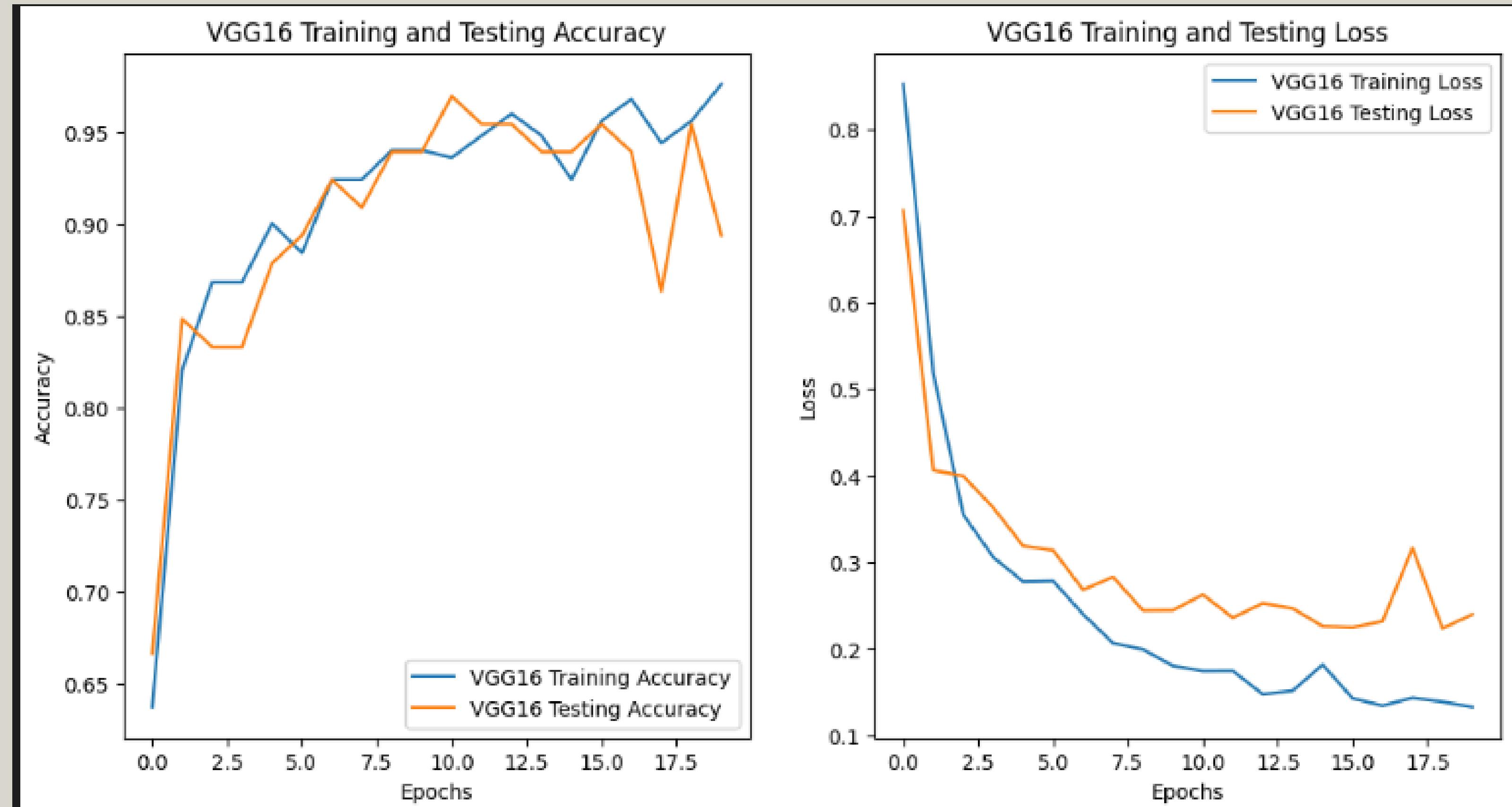
VGG16 Training Accuracy: 0.9721115827560425  
VGG16 Test Accuracy: 0.8939393758773804

ResNet50 Training Accuracy: 0.8764940500259399  
ResNet50 Test Accuracy: 0.7727272510528564

# Visualization







A photograph of two petri dishes containing bacterial cultures. The dish on the left shows several distinct, white, irregular colonies on a red agar medium. The dish on the right contains a clear, linear DNA ladder, with visible bands of increasing size moving from top to bottom. Both dishes are held by hands wearing white gloves against a dark background.

**YOLO:  
IDENTIFICATION**

# yolo interaction with vgg16

```
# Function to detect defects using YOLO and classify using VGG16
def detect_defects_yolo(image_path, net, output_layers, classes, vgg_model):
    img = cv2.imread(image_path)
    height, width, channels = img.shape

    blob = cv2.dnn.blobFromImage(img, 0.00392, (416, 416), (0, 0, 0), True, crop=False)
    net.setInput(blob)
    outs = net.forward(output_layers)

    class_ids = []
    confidences = []
    boxes = []

    for out in outs:
        for detection in out:
            scores = detection[5:]
            class_id = np.argmax(scores)
            confidence = scores[class_id]
            if confidence > 0.5: # Confidence threshold
                center_x = int(detection[0] * width)
                center_y = int(detection[1] * height)
                w = int(detection[2] * width)
                h = int(detection[3] * height)
                x = int(center_x - w / 2)
                y = int(center_y - h / 2)
                boxes.append([x, y, w, h])
                confidences.append(float(confidence))
                class_ids.append(class_id)
```

```
indexes = cv2.dnn.NMSBoxes(boxes, confidences, 0.5, 0.4)
detected_regions = []
detected_boxes = []
detected_labels = []

for i in range(len(boxes)):
    if i in indexes:
        x, y, w, h = boxes[i]
        detected_regions.append(img[y:y+h, x:x+w]) # Extract region of interest
        detected_boxes.append((x, y, x+w, y+h)) # Store box coordinates

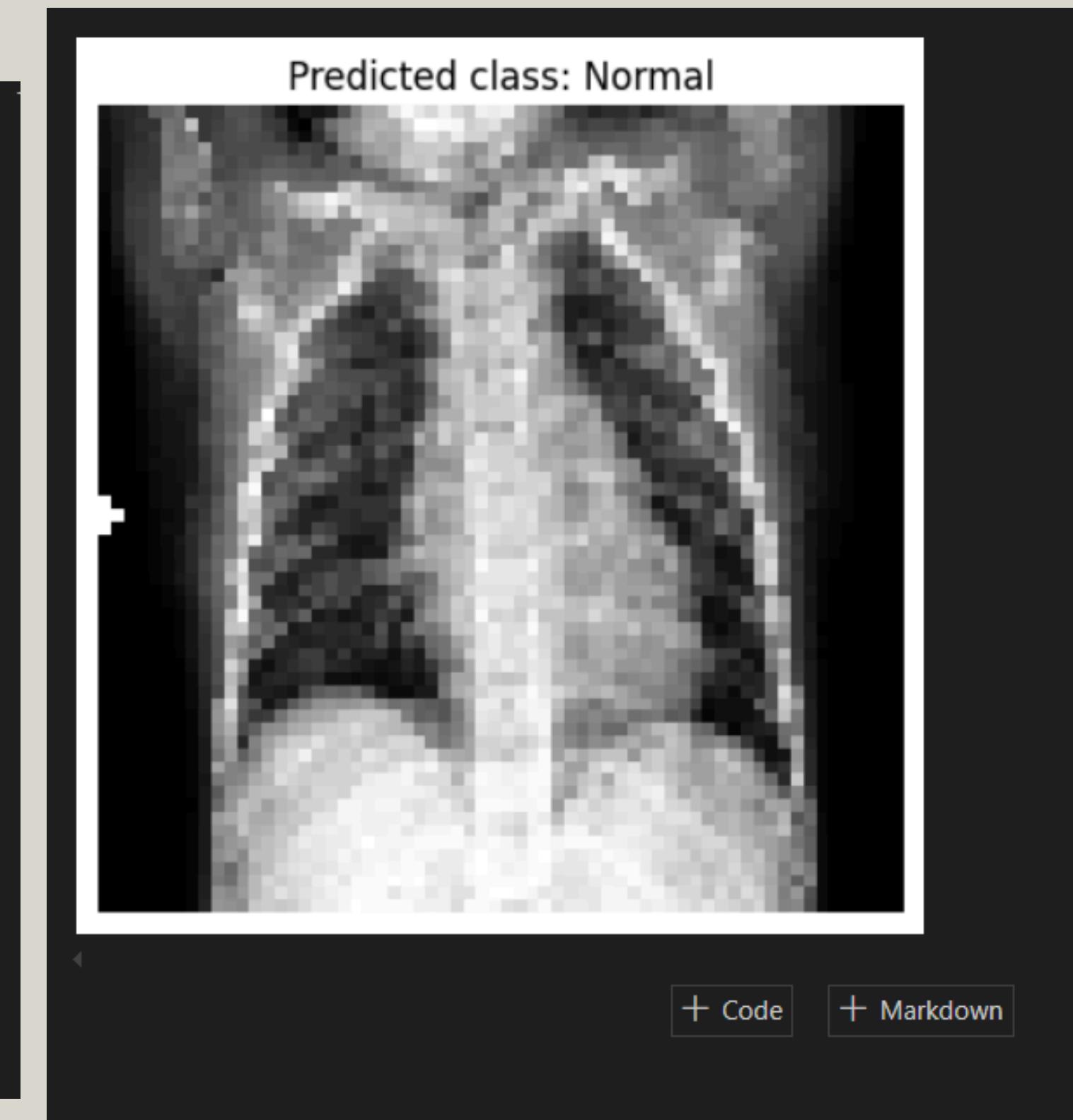
    # Perform classification using VGG16 model
    region_img = cv2.resize(img[y:y+h, x:x+w], (64, 64)) # Resize image to VGG16 input size
    region_img = cv2.cvtColor(region_img, cv2.COLOR_BGR2RGB) # Convert to RGB (VGG16 expects RGB)
    region_img = region_img.astype(np.float32) / 255.0 # Normalize

    # Predict using VGG16 model
    prediction = vgg_model.predict(np.expand_dims(region_img, axis=0))
    predicted_class = np.argmax(prediction)
    label = classes[predicted_class]
    detected_labels.append(label)

    # Draw rectangle and label on the original image
    cv2.rectangle(img, (x, y), (x + w, y + h), (0, 255, 0), 2)
    cv2.putText(img, label, (x, y - 10), cv2.FONT_HERSHEY_SIMPLEX, 0.5, (0, 255, 0))

return img, detected_regions, detected_boxes, detected_labels
```

```
WARNING:absl:Compiled the loaded model, but the compiled metrics have yet  
1/1 ━━━━━━ 0s 237ms/step
```



# Flask



# Detection with flask

COVID-19 Detection App

Choose file No file chosen



*Thank you for your  
attention!*