

PNEUMONIA CLASSIFICATION

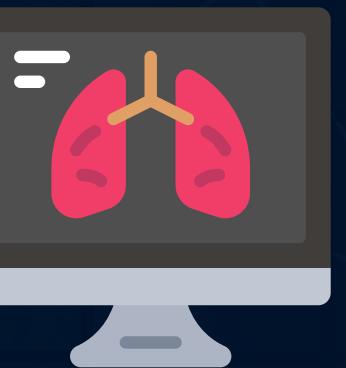


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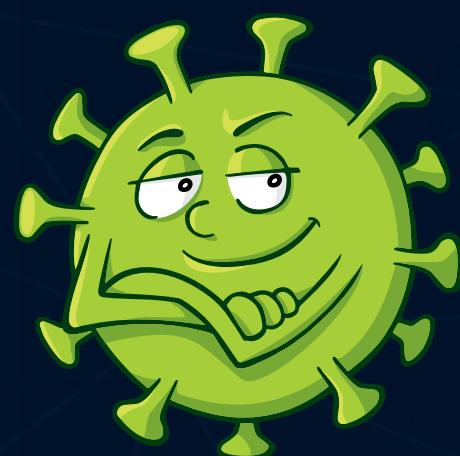
08 Deployment



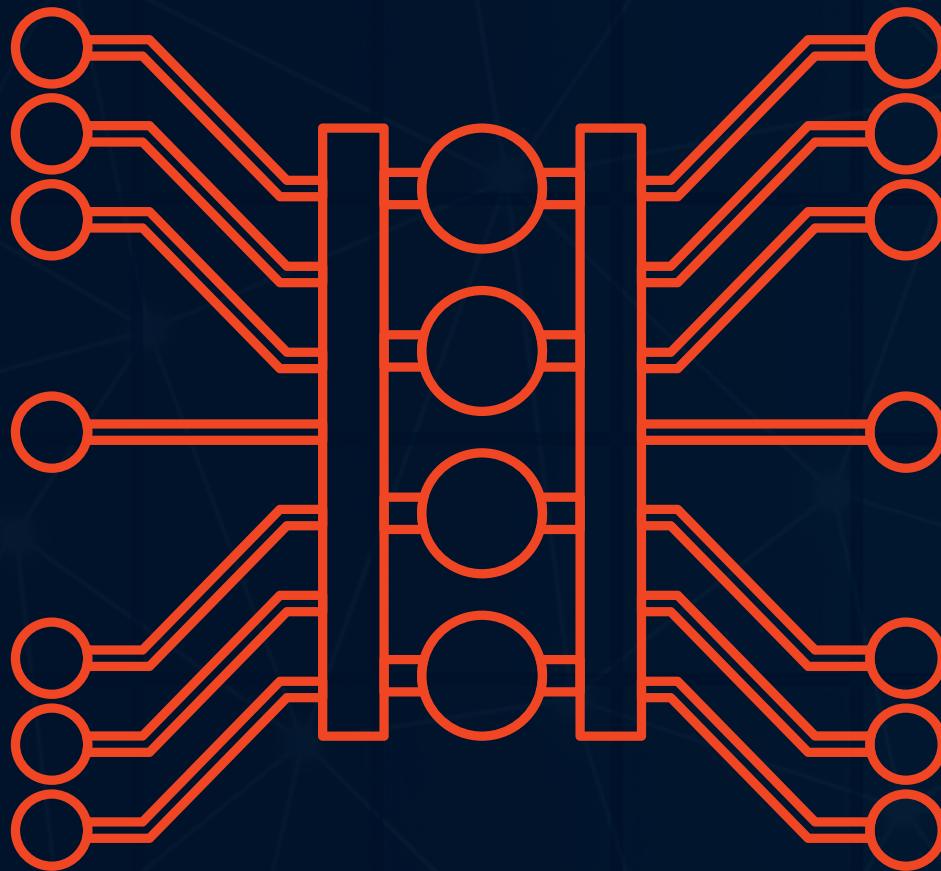
PNEUMONIA PROBLEM



Pneumonia is a life-threatening lung infection that requires prompt diagnosis and treatment. It's a prevalent respiratory disease and a respiratory infection that primarily affects the lungs, causing inflammation in the air sacs. It can be caused by bacteria, viruses, or fungi and is characterized by symptoms such as cough, fever, difficulty breathing, and chest pain. Pneumonia can range from mild to severe, with complications more likely in the elderly and individuals with weakened immune systems. Diagnosis often involves chest X-rays and laboratory tests.



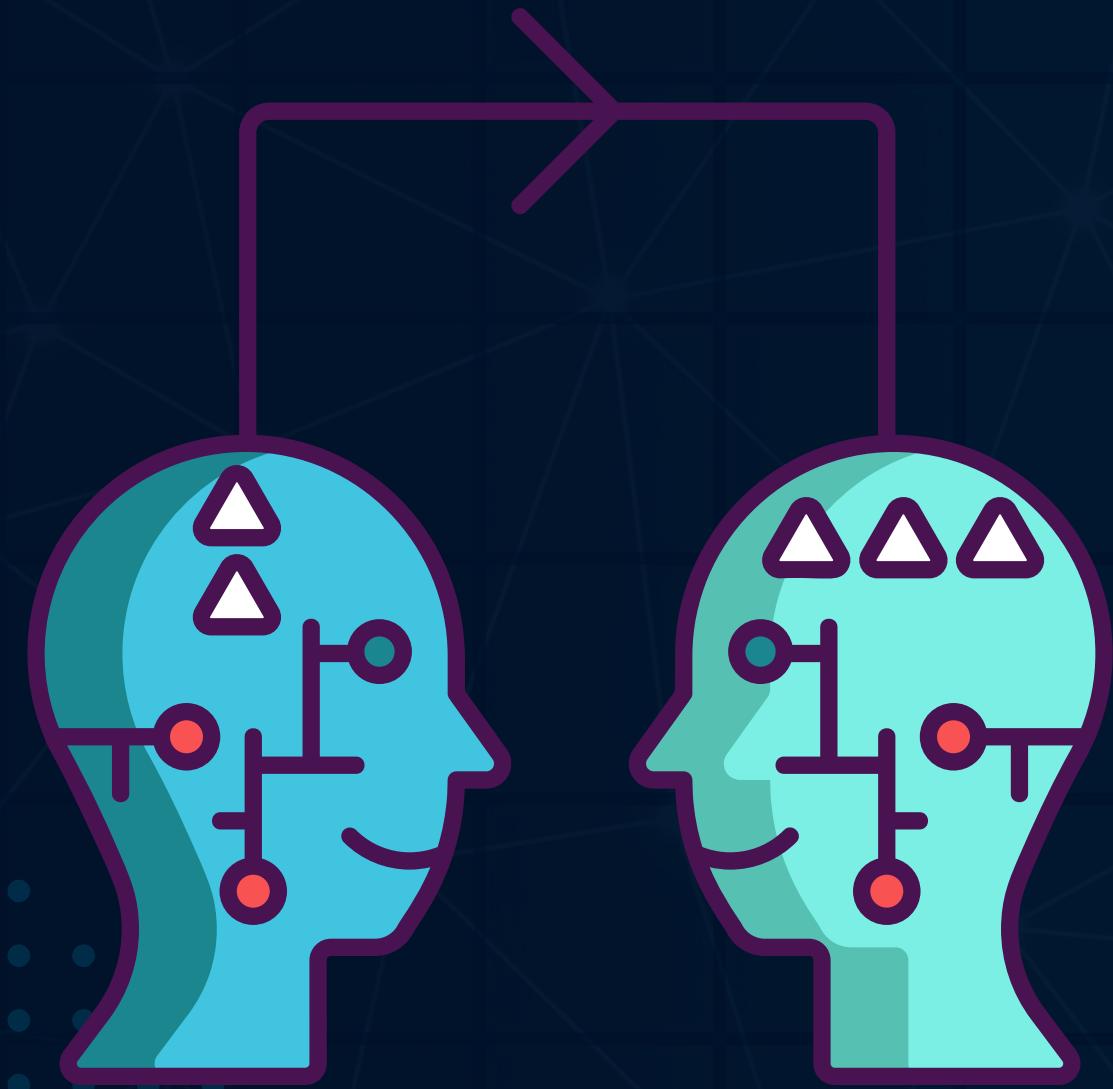
ABSTRACT



The use of deep learning techniques, particularly convolutional neural networks (CNNs), has emerged as a promising approach for pneumonia classification using medical imaging data.

These models demonstrate the potential to enhance diagnostic accuracy, assist healthcare professionals in making informed decisions, and improve the overall management of pneumonia cases. By analyzing radiological images, these models can accurately differentiate between pneumonia and healthy lungs, aiding in the early detection and treatment of the disease.

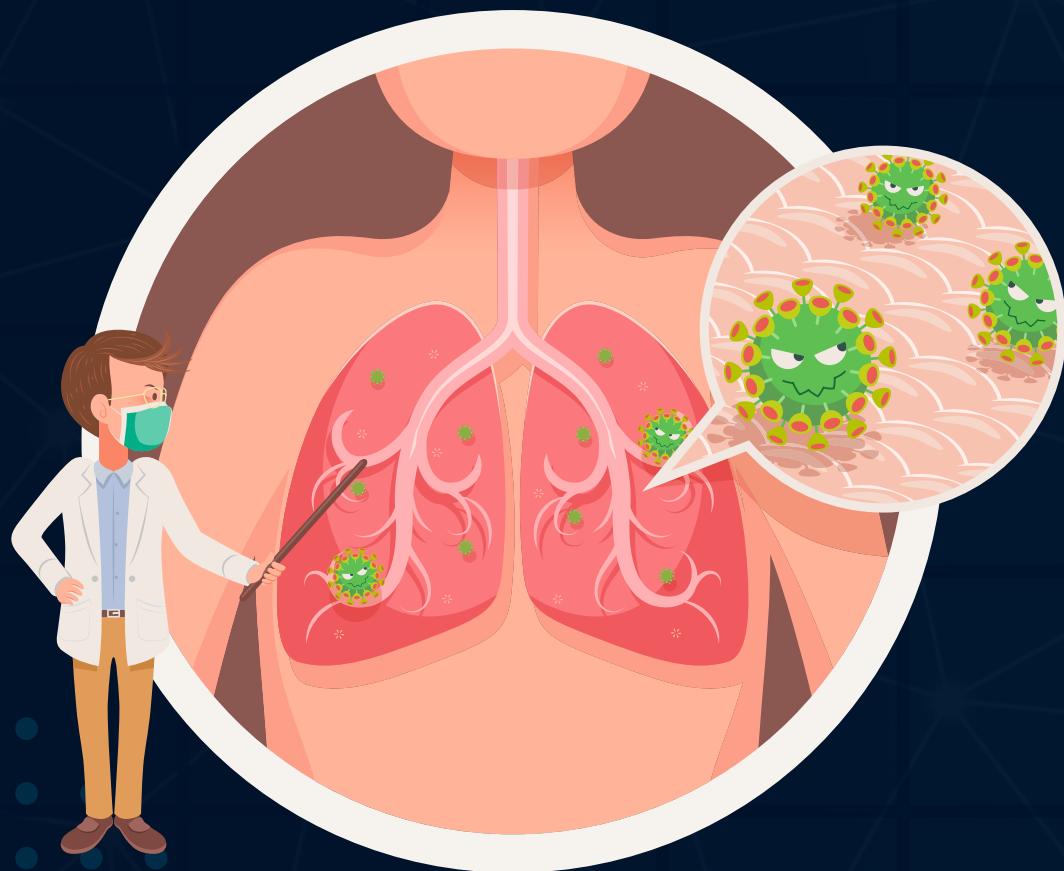




Nonetheless, the process of training deep learning models from scratch demands abundant labeled data and significant computational resources, which may not be easily accessible. To overcome this challenge, transfer learning comes into play, enabling us to harness the power of pre-trained CNN models that have already undergone extensive training on vast image datasets. By adapting these pre-trained models to new tasks using smaller datasets, transfer learning empowers us to make efficient use of existing knowledge and achieve desirable outcomes.

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INTRODUCTION



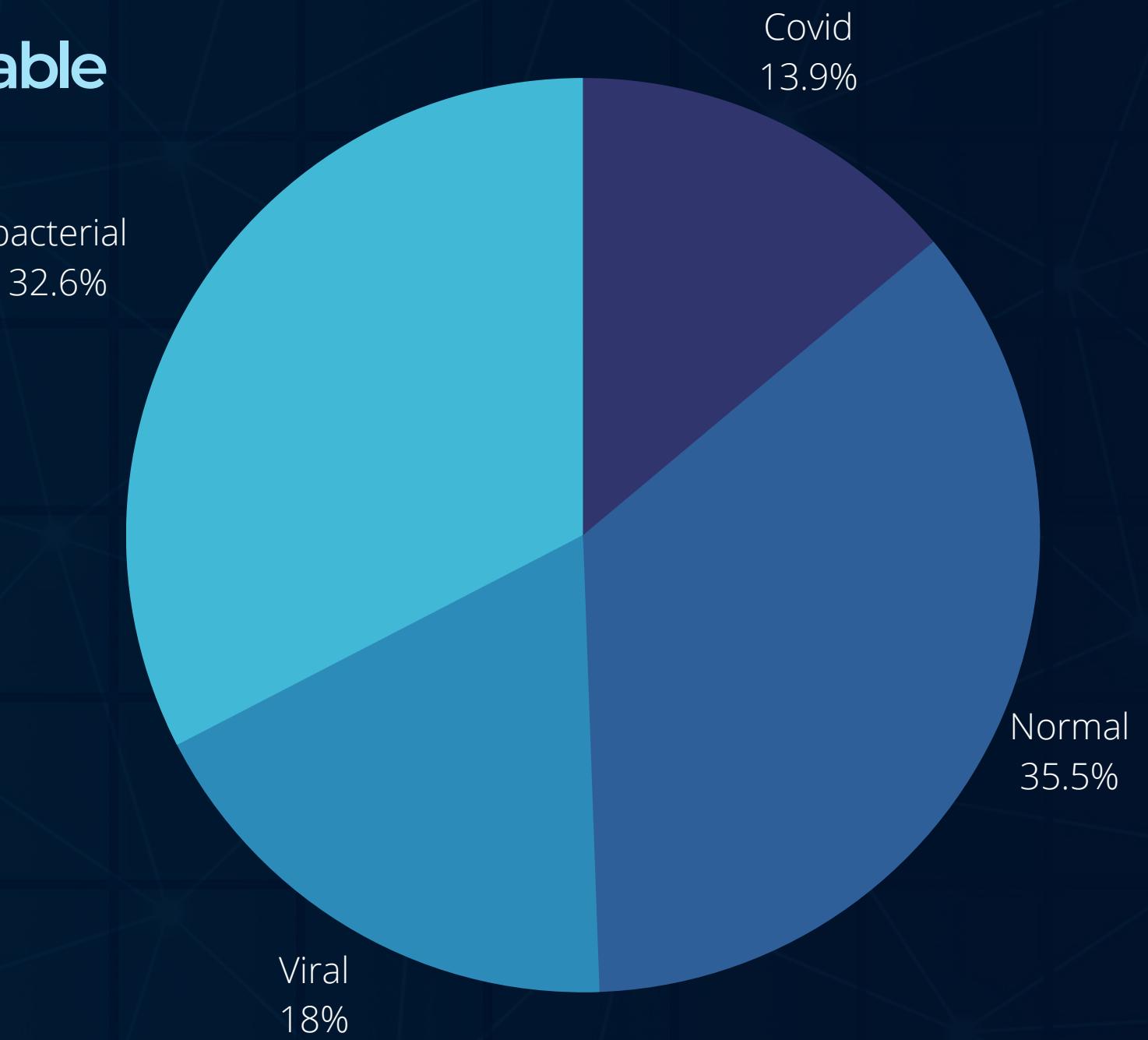
In the field of Pneumonia classification, our objective is to develop a robust deep learning model capable of accurately categorizing chest X-ray images as either pneumonia-infected or non-infected using state-of-the-art techniques like convolutional neural networks (CNNs) and transfer learning models. By employing architectures such as densenet121, mobilenetv2, xception, and vgg16, we aim to achieve high accuracy in the classification of pneumonia cases based on the analysis of medical images. This Project aims to contribute to the early and accurate diagnosis of pneumonia, enabling timely intervention and improved patient outcomes.

04

OUR DATASET

This is a combined curated dataset of COVID-19 Chest X-ray images obtained by collating 15 public available datasets.

- Curated X-Ray Dataset
 - Normal
 - Pneumonia-Bacterial
 - Pneumonia-Viral
 - COVID-19



04

OUR DATASET

- The present dataset contains:
 - 1281 COVID-19 X-Rays
 - 3270 Normal X-Rays
 - 1656 viral-pneumonia X-Rays
 - 3001 bacterial-pneumonia X-Rays.

REVENUE

4,000

3,000

2,000

1,000

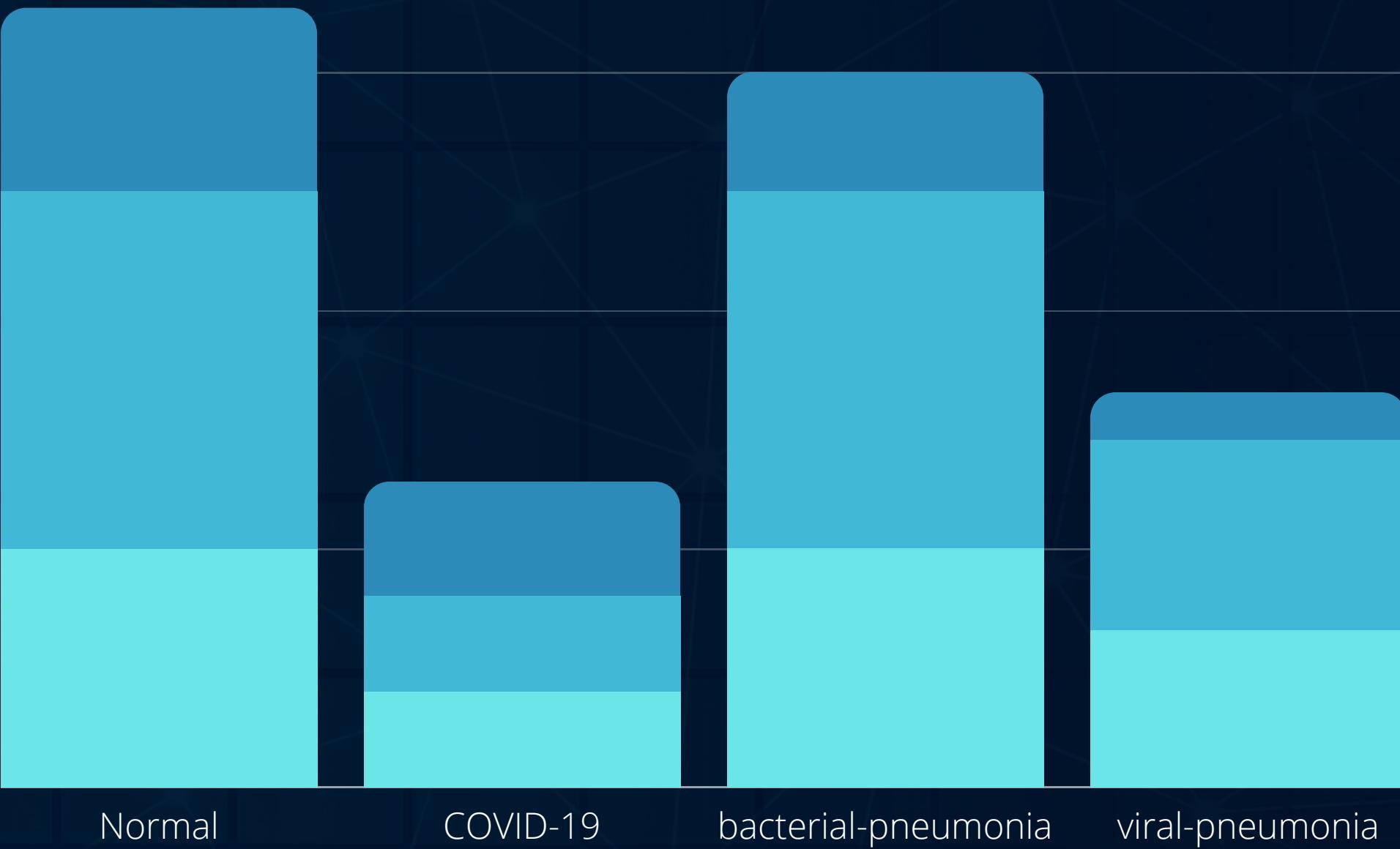
0

Normal

COVID-19

bacterial-pneumonia

viral-pneumonia



04

OUR DATASET

TEST
PERSENTAGE



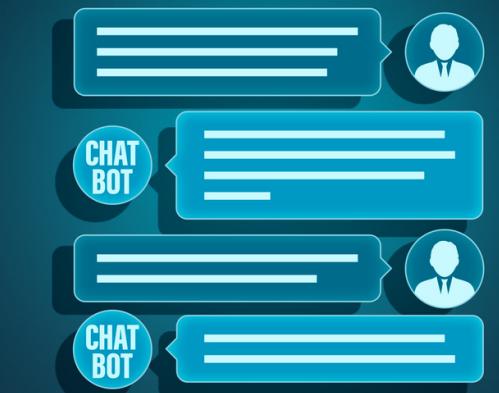
15%

TRAIN
PERSENTAGE



85%

DIVEDING DATA TO TRAIN AND TEST





PREPROCESSING



Checking the size of images in different categories :

- COVID,
- Normal,
- Bacterial
- Viral.



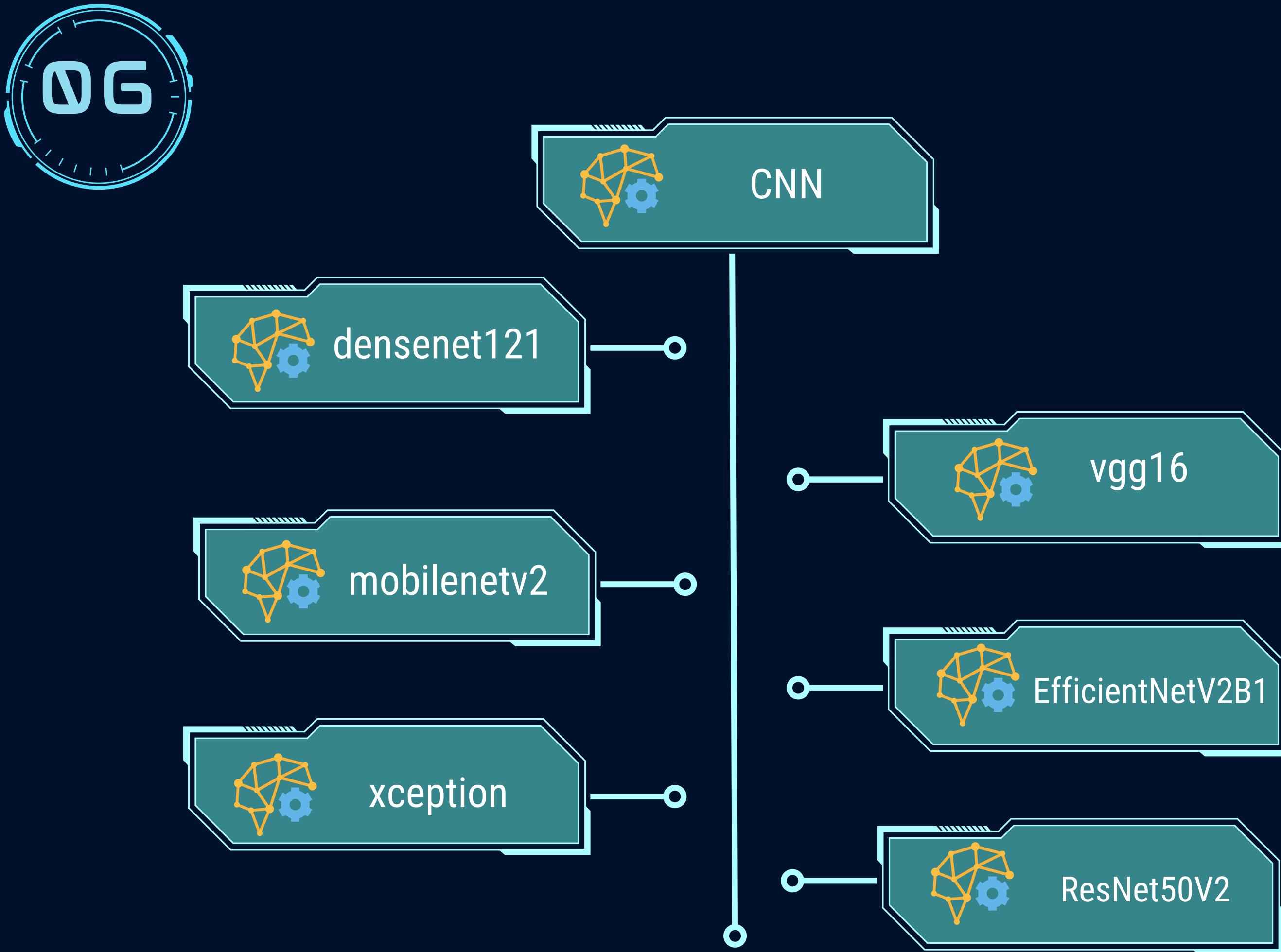
reading JPG images from each category's directory, retrieves their shapes, and generates a frequency count of the unique image shapes using Pandas. The resulting count provides information on the distribution of image sizes within each category.

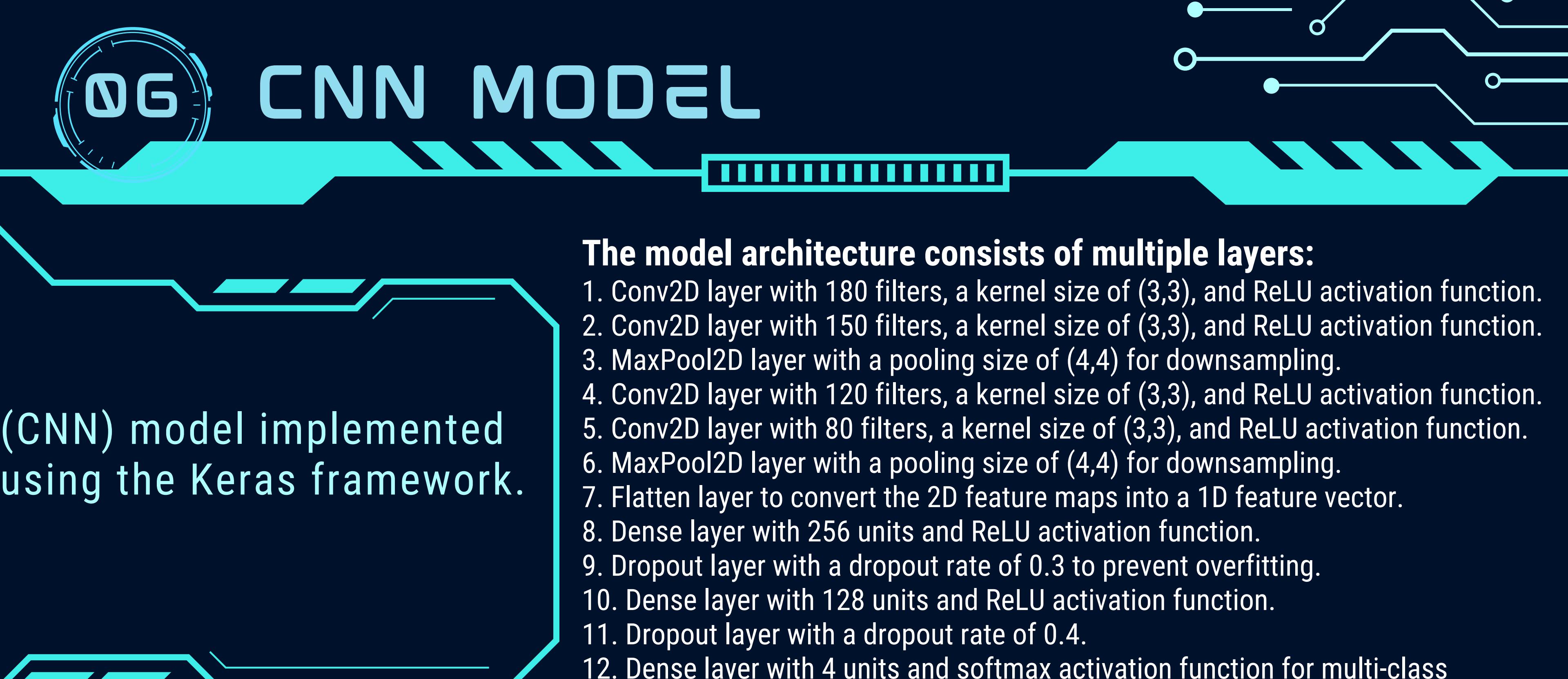


- preparing training data for a model by resizing the images to a consistent size (150x150) and storing them with their respective class labels.
- Separate labels and inputs
- divide data to train and test



CNN and TL Models





The model is compiled with the Adam optimizer, sparse categorical cross-entropy loss function, and accuracy as the evaluation metric. This model is suitable for our image classification tasks with four output classes.



MODELS ACCURACY

CNN



TRAINING
ACCURACY



95.6%

TRAINING
LOSS



11.73%

EVALUATE
ACCURACY



88.06%

EVALUATE
LOSS



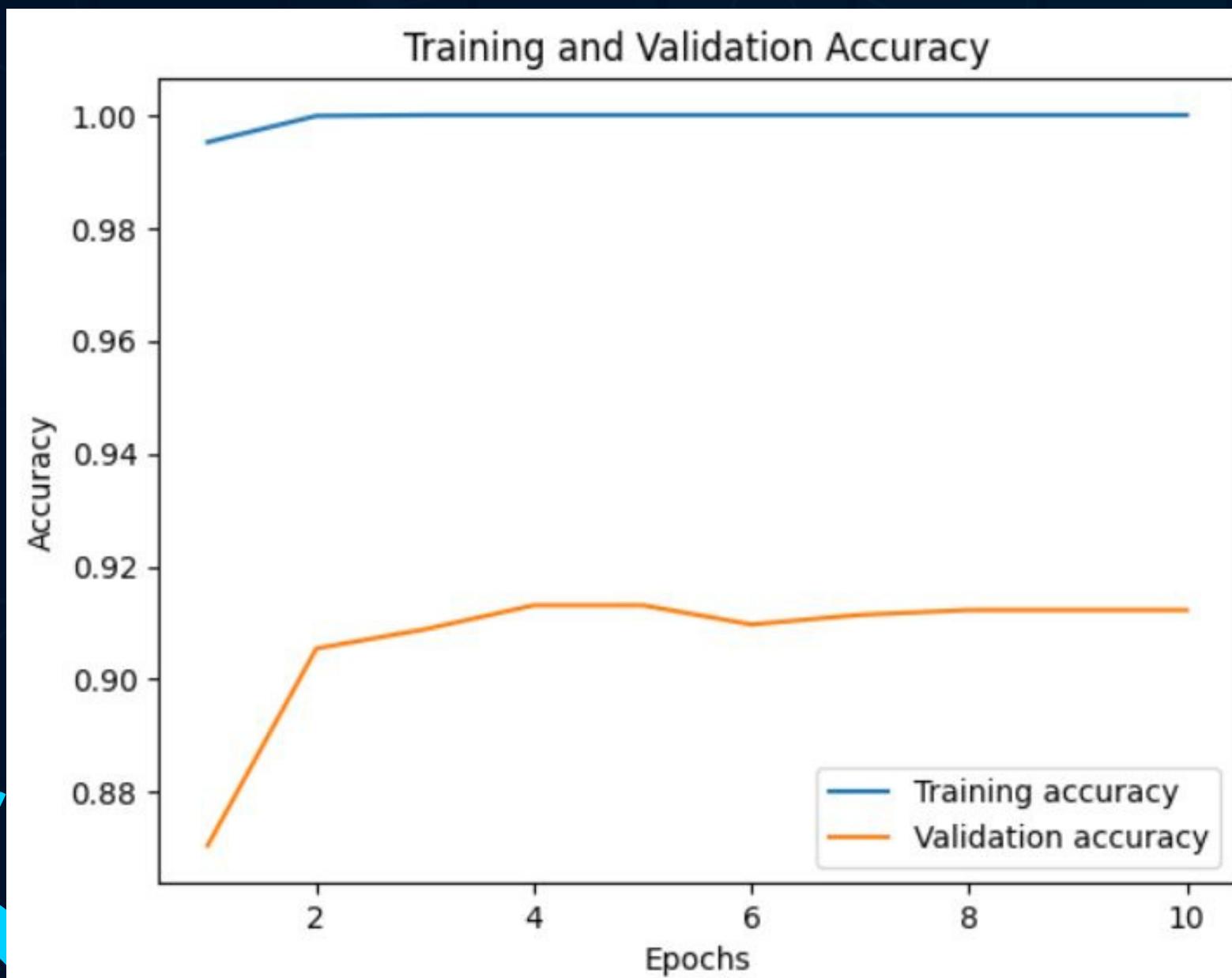
44.54%



07

MODELS ACCURACY DENSENET121

GAP BETWEEN TRAINING ACCURACY
AND VALIDATION ACCURACY



GAP BETWEEN TRAINING LOSS
AND VALIDATION LOSS



From epoch 11

07

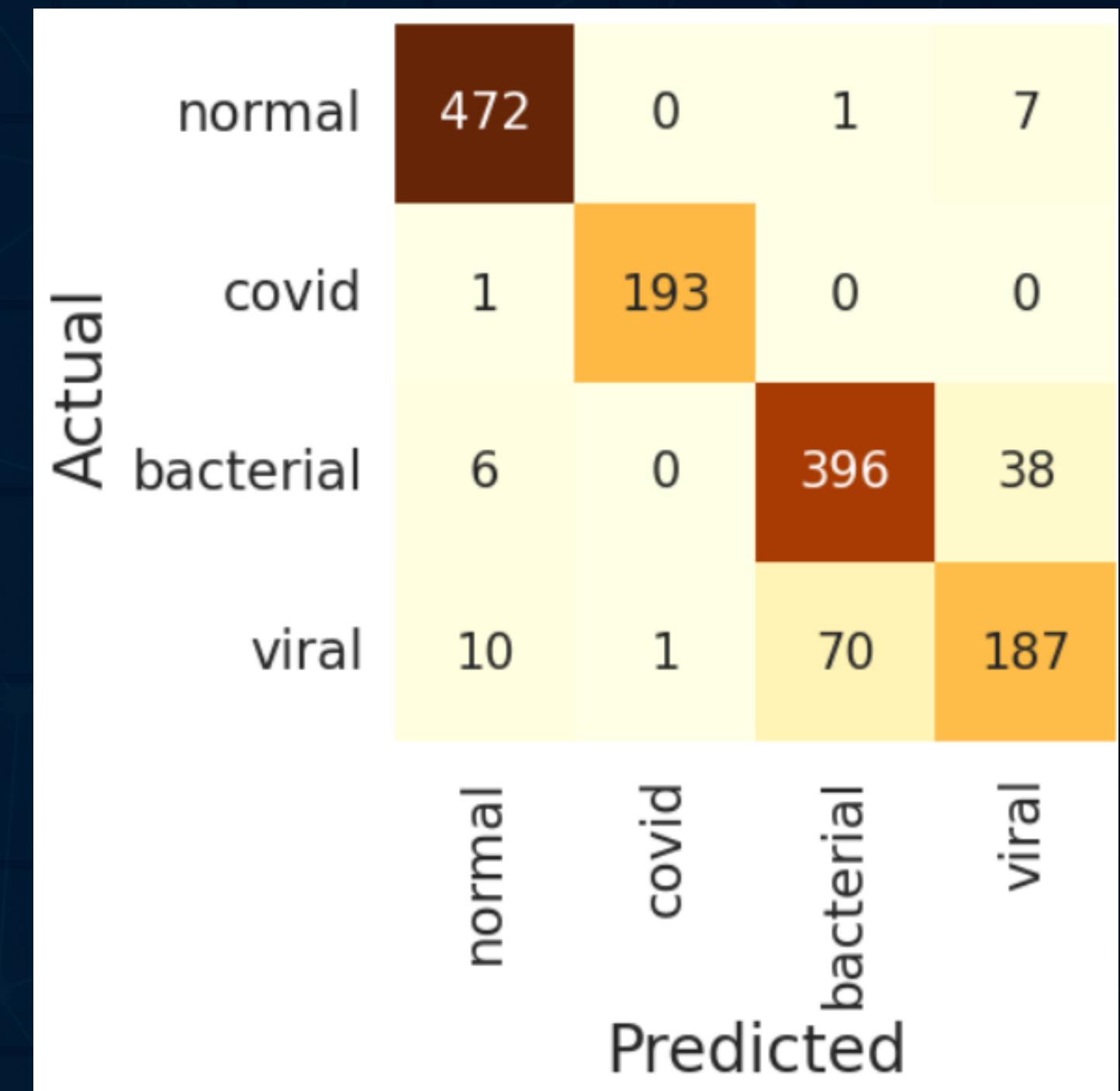
MODELS ACCURACY

DENSENET121

CLASSIFICATION REPORT

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| normal | 0.9652 | 0.9833 | 0.9742 | 480 |
| covid | 0.9948 | 0.9948 | 0.9948 | 194 |
| bacterial | 0.8480 | 0.9000 | 0.8732 | 440 |
| viral | 0.8060 | 0.6978 | 0.7480 | 268 |
| accuracy | | | 0.9030 | 1382 |
| macro avg | 0.9035 | 0.8940 | 0.8976 | 1382 |
| weighted avg | 0.9012 | 0.9030 | 0.9011 | 1382 |

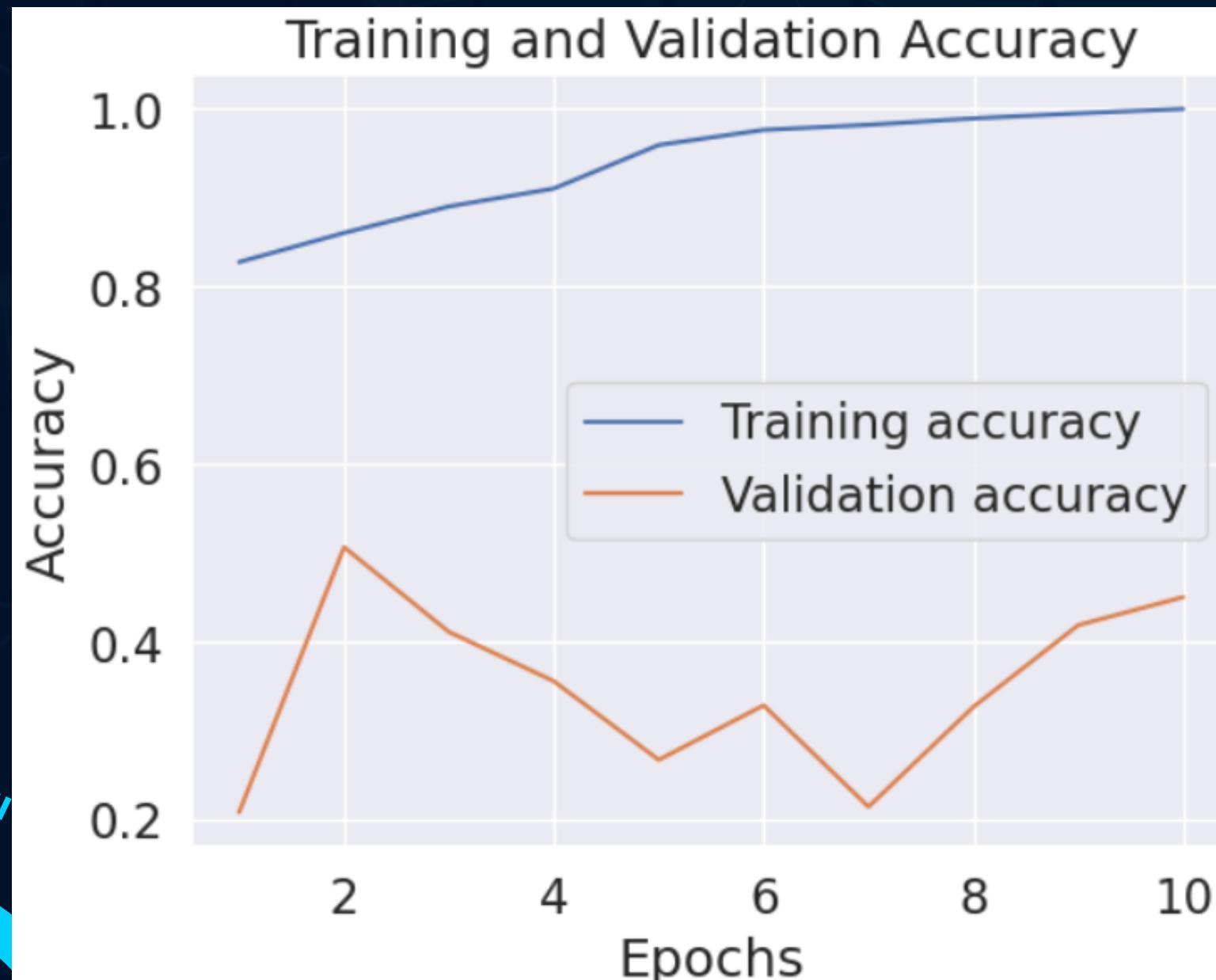
CONFUSION MATRIX



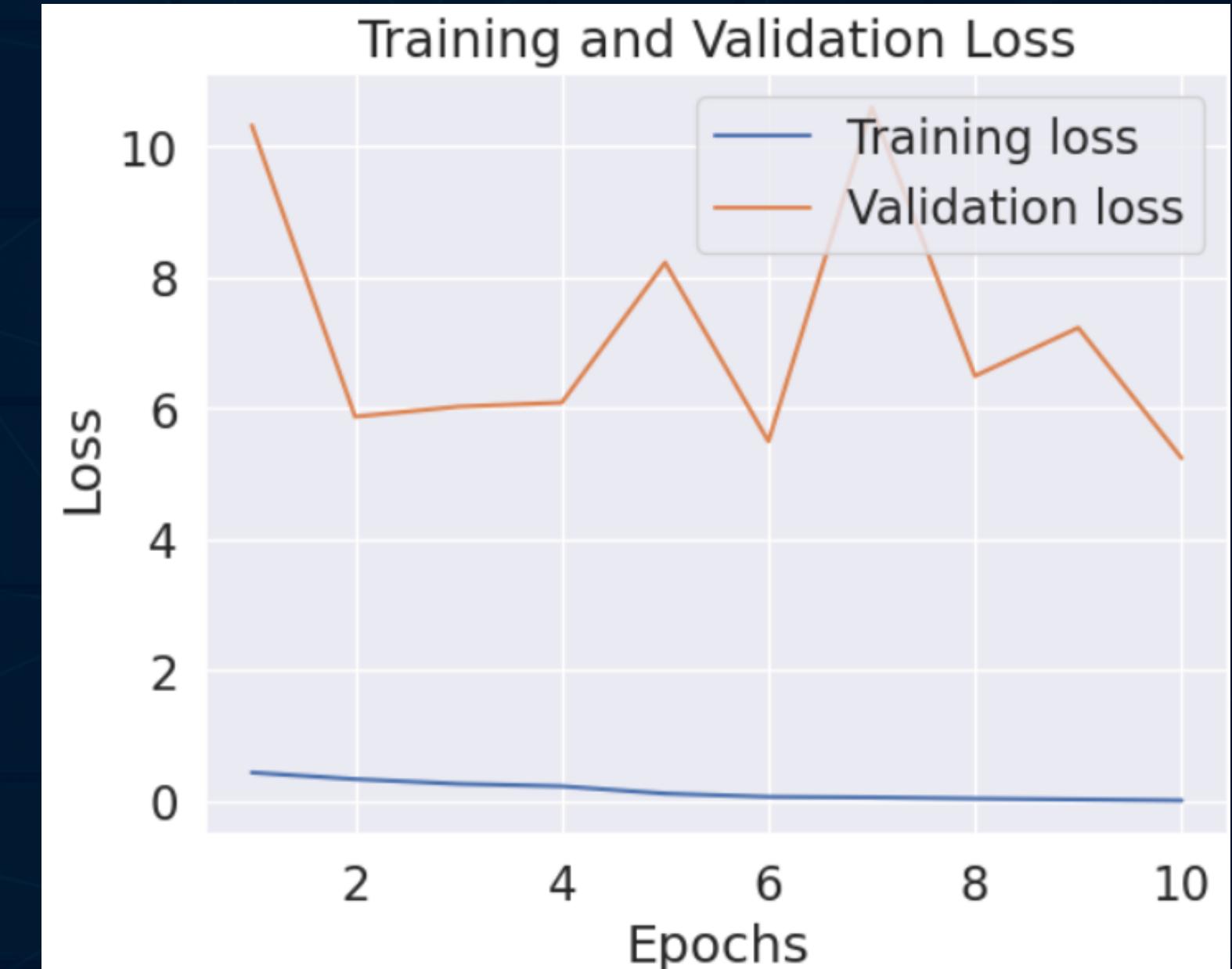
07

MODELS ACCURACY MOBILENET

GAP BETWEEN TRAINING ACCURACY
AND VALIDATION ACCURACY



GAP BETWEEN TRAINING LOSS
AND VALIDATION LOSS



07

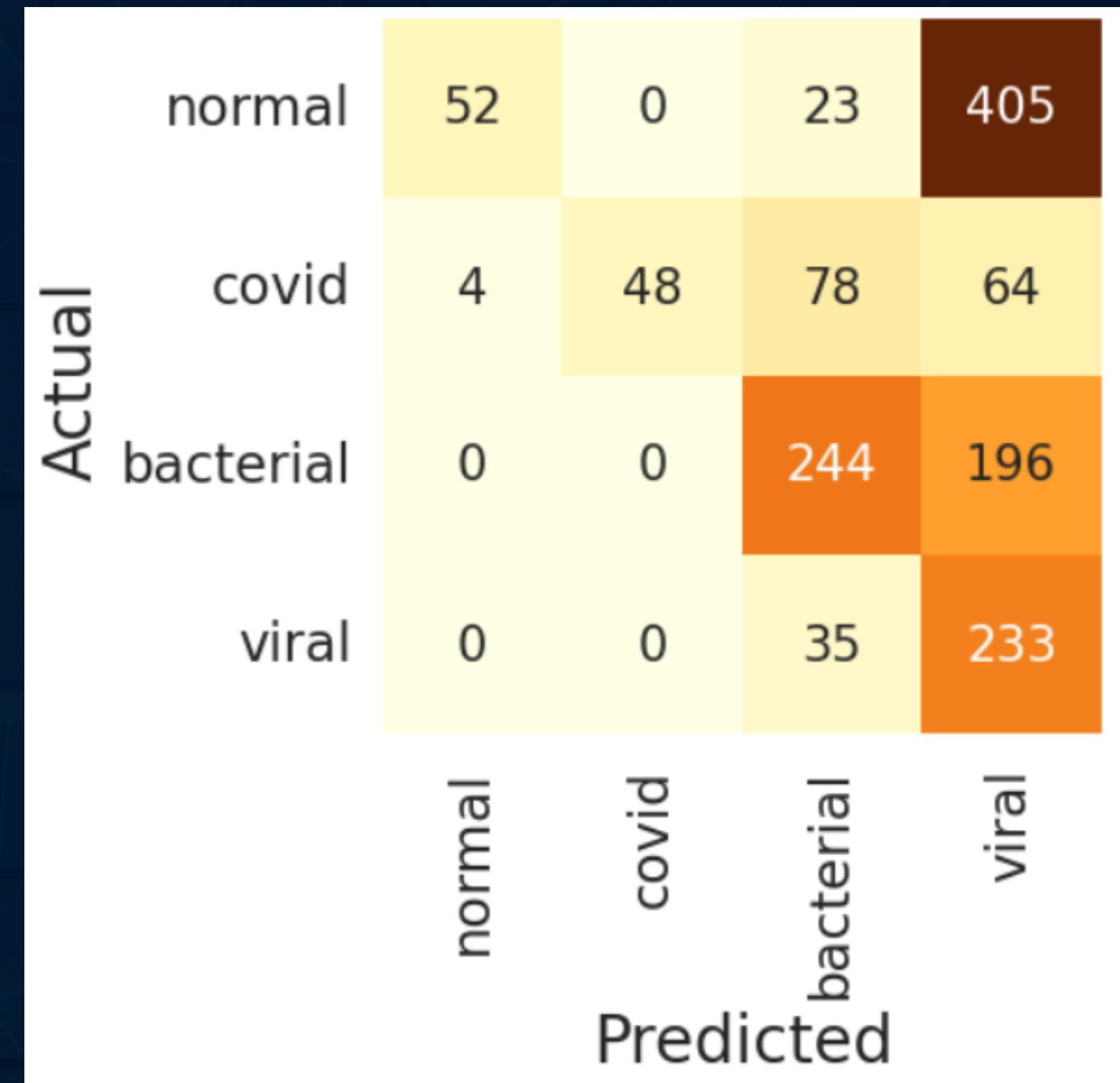
MODELS ACCURACY

MOBILENET

CLASSIFICATION REPORT

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| normal | 0.9286 | 0.1083 | 0.1940 | 480 |
| covid | 1.0000 | 0.2474 | 0.3967 | 194 |
| bacterial | 0.6421 | 0.5545 | 0.5951 | 440 |
| viral | 0.2595 | 0.8694 | 0.3997 | 268 |
| accuracy | | | 0.4175 | 1382 |
| macro avg | 0.7075 | 0.4449 | 0.3964 | 1382 |
| weighted avg | 0.7176 | 0.4175 | 0.3901 | 1382 |

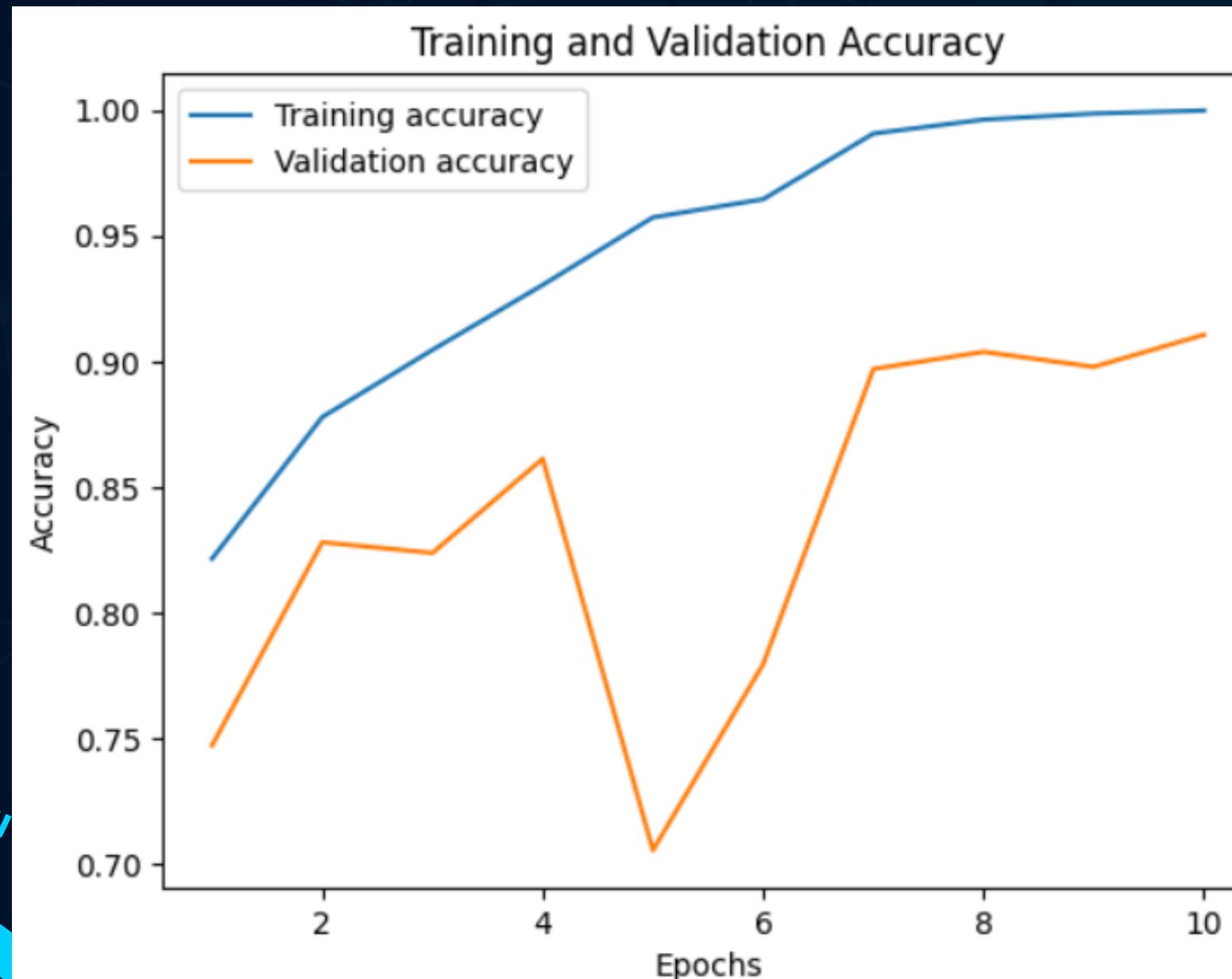
CONFUSION MATRIX



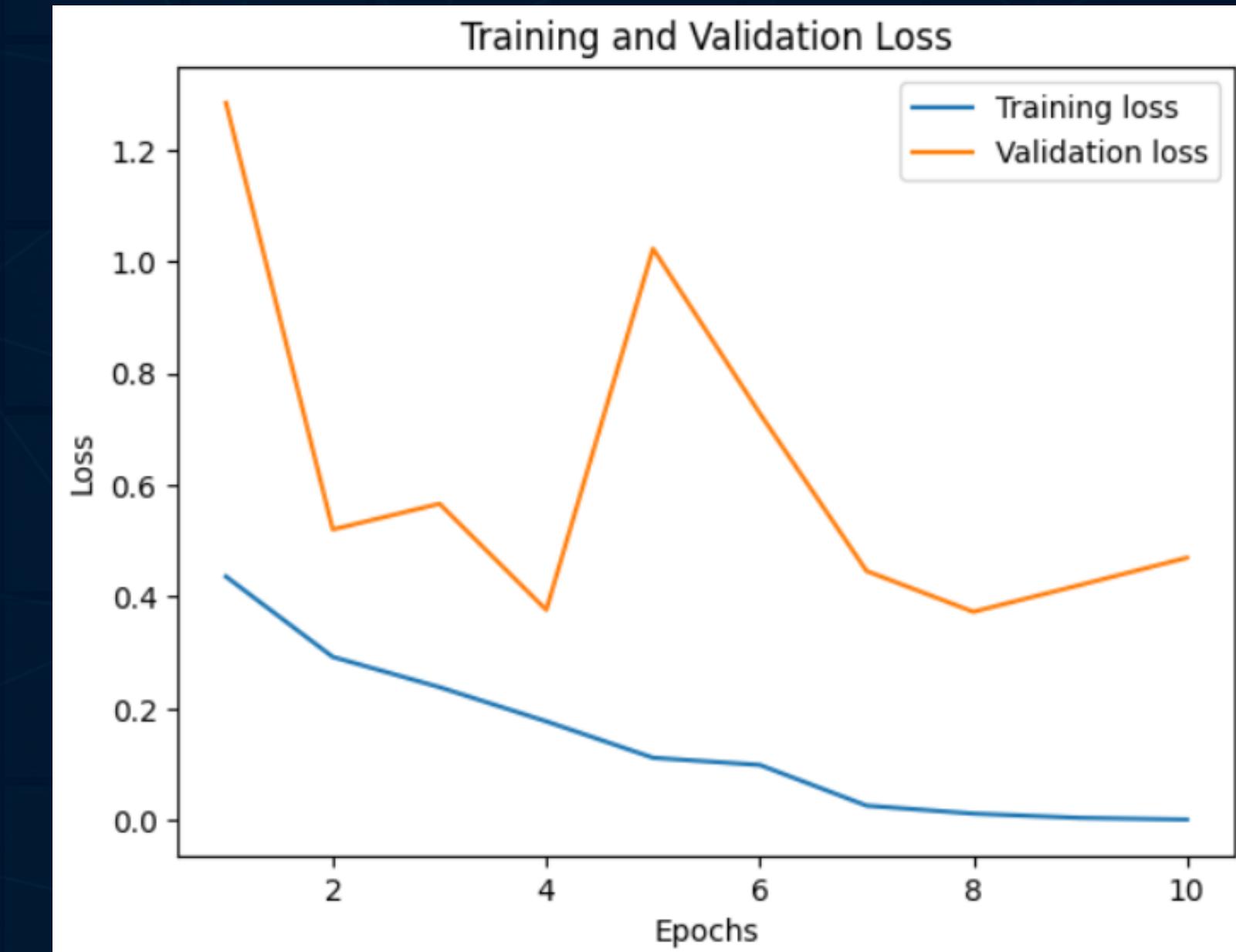
07

MODELS ACCURACY XCEPTION

GAP BETWEEN TRAINING ACCURACY
AND VALIDATION ACCURACY



GAP BETWEEN TRAINING LOSS
AND VALIDATION LOSS



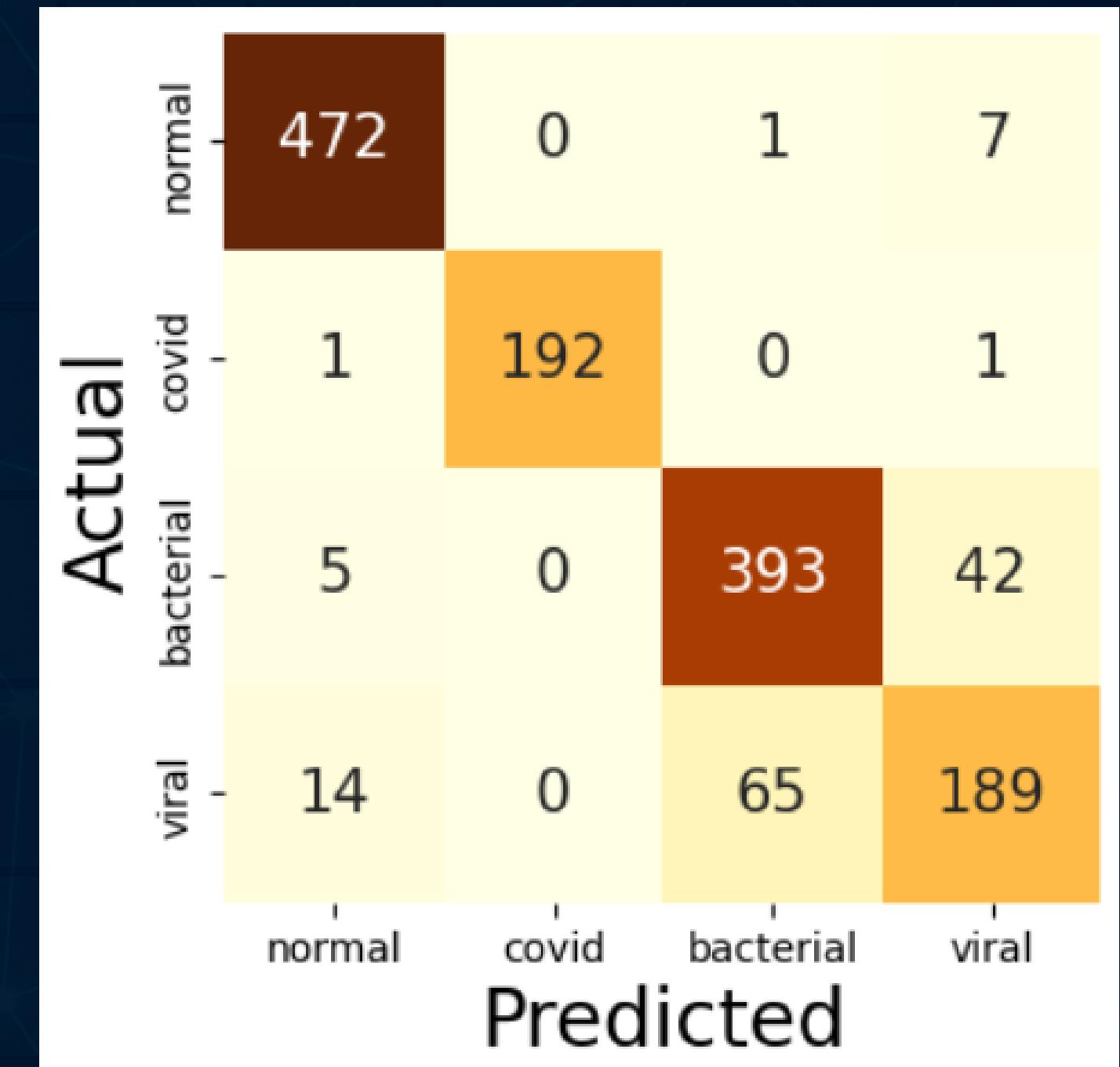
07

MODELS ACCURACY XCEPTION

CLASSIFICATION REPORT

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| normal | 0.9593 | 0.9833 | 0.9712 | 480 |
| covid | 1.0000 | 0.9897 | 0.9948 | 194 |
| bacterial | 0.8562 | 0.8932 | 0.8743 | 440 |
| viral | 0.7908 | 0.7052 | 0.7456 | 268 |
| accuracy | | | 0.9016 | 1382 |
| macro avg | 0.9016 | 0.8929 | 0.8965 | 1382 |
| weighted avg | 0.8995 | 0.9016 | 0.8999 | 1382 |

CONFUSION MATRIX

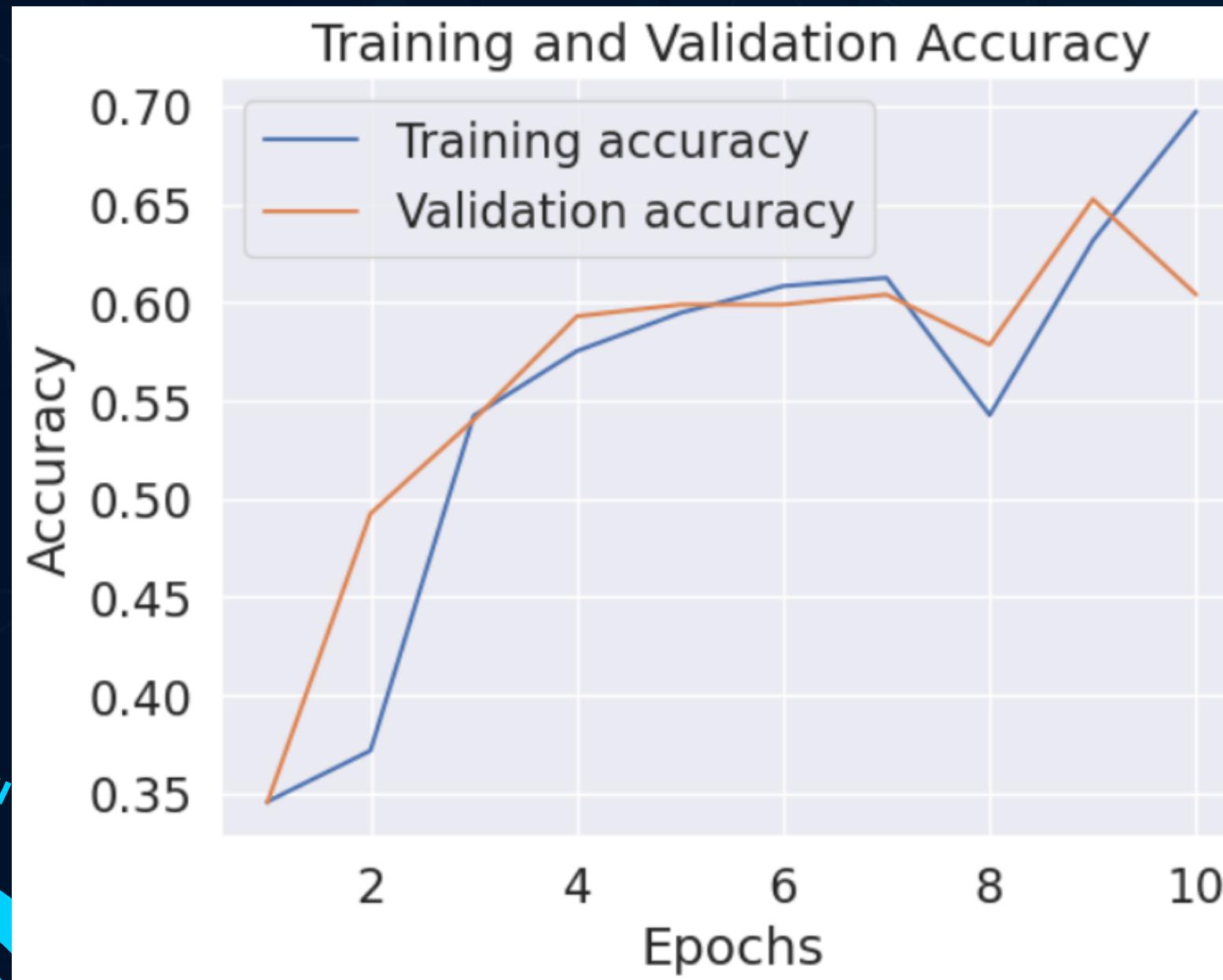


07

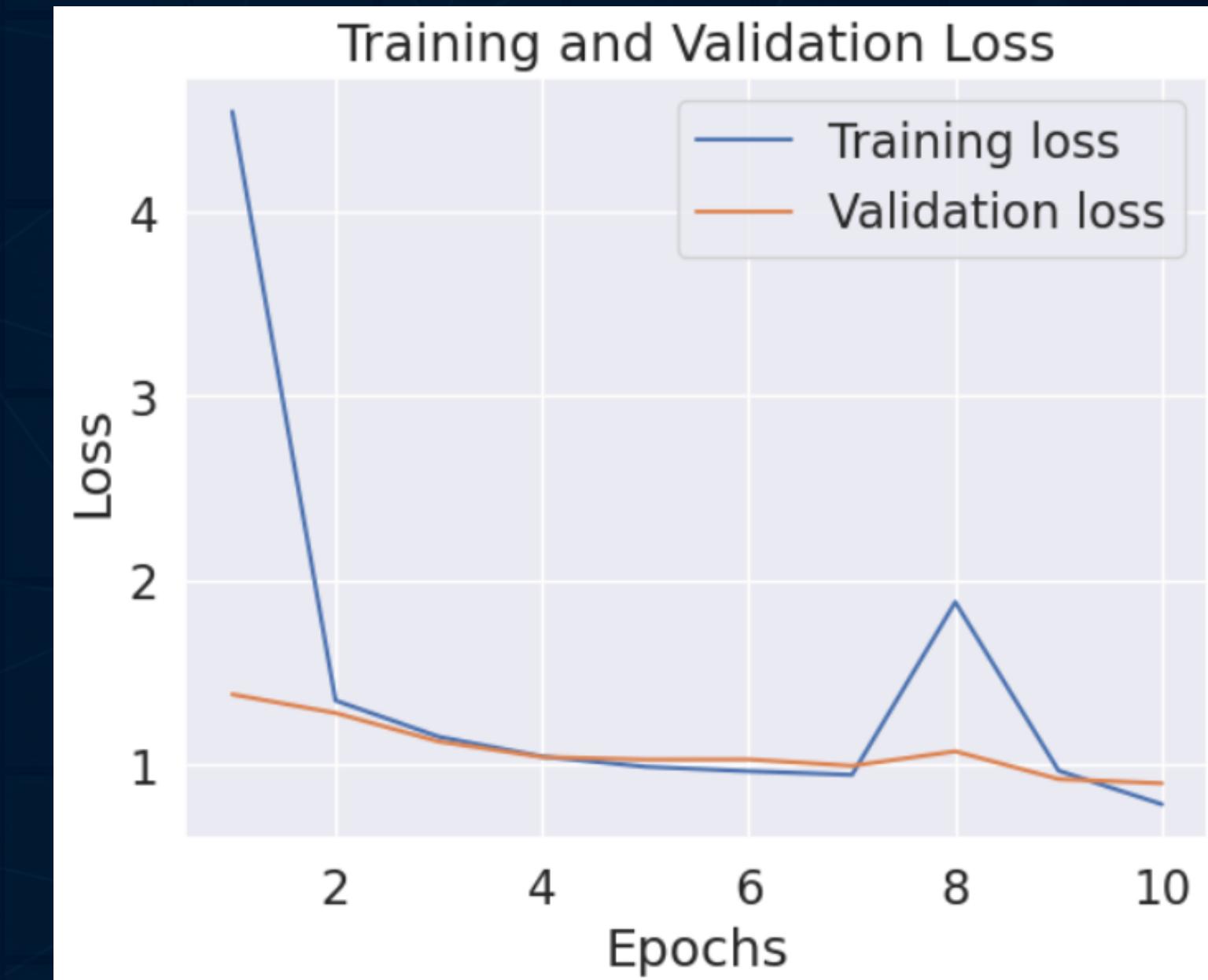
MODELS ACCURACY

VGG16

GAP BETWEEN TRAINING ACCURACY
AND VALIDATION ACCURACY



GAP BETWEEN TRAINING LOSS
AND VALIDATION LOSS



07

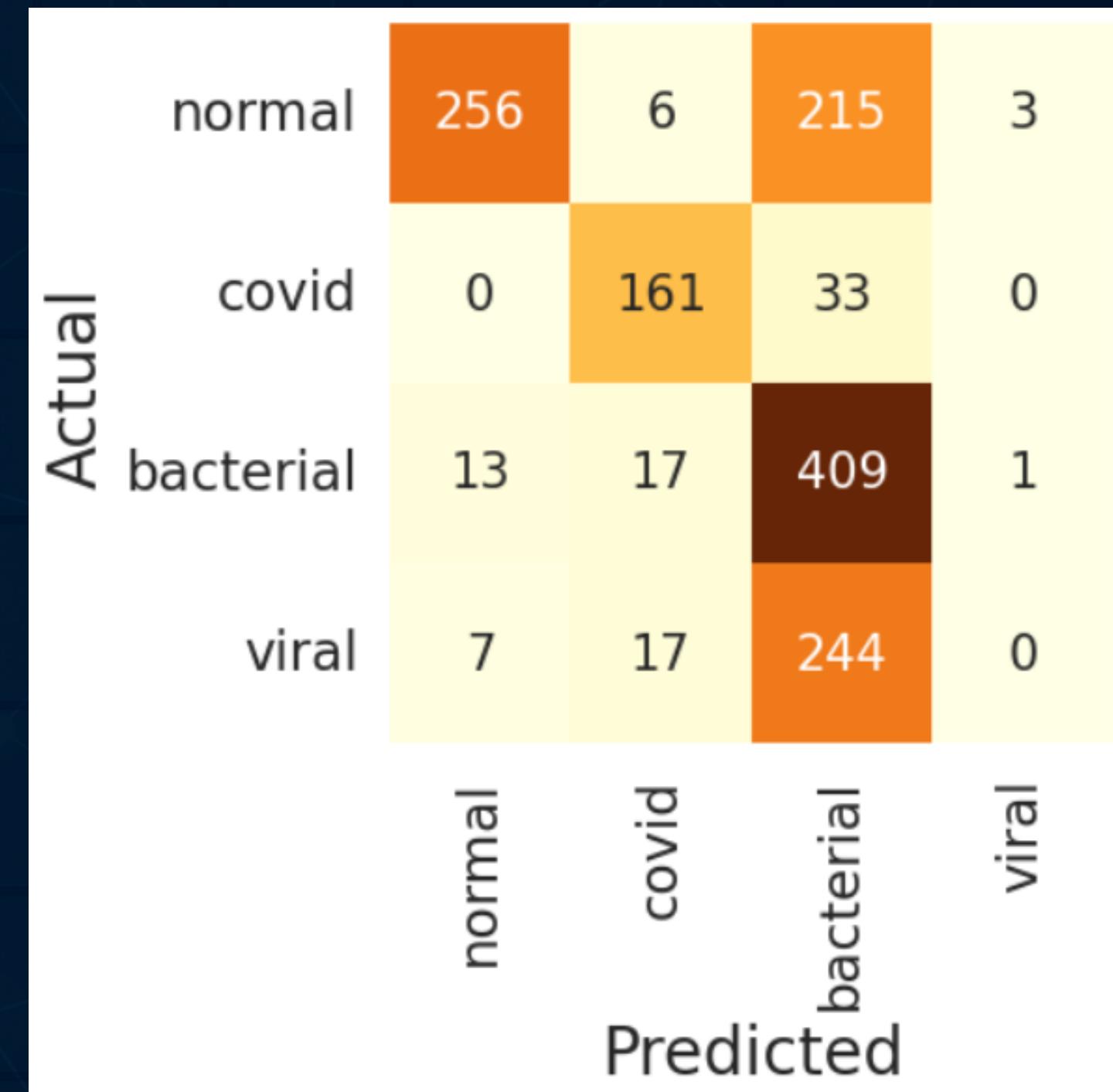
MODELS ACCURACY

VGG16

CLASSIFICATION REPORT

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| normal | 0.9275 | 0.5333 | 0.6772 | 480 |
| covid | 0.8010 | 0.8299 | 0.8152 | 194 |
| bacterial | 0.4539 | 0.9295 | 0.6100 | 440 |
| viral | 0.0000 | 0.0000 | 0.0000 | 268 |
| accuracy | | | 0.5977 | 1382 |
| macro avg | 0.5456 | 0.5732 | 0.5256 | 1382 |
| weighted avg | 0.5791 | 0.5977 | 0.5439 | 1382 |

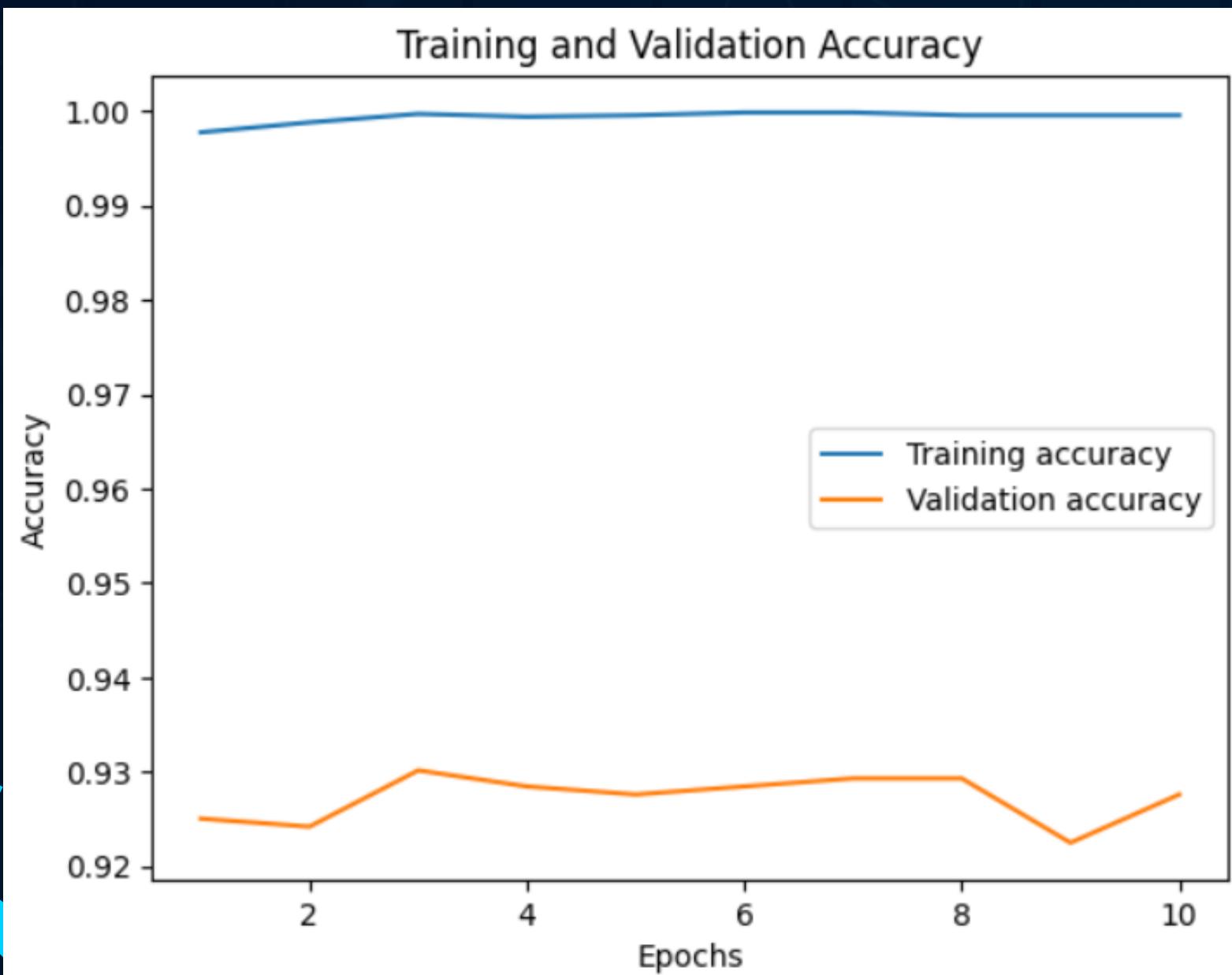
CONFUSION MATRIX



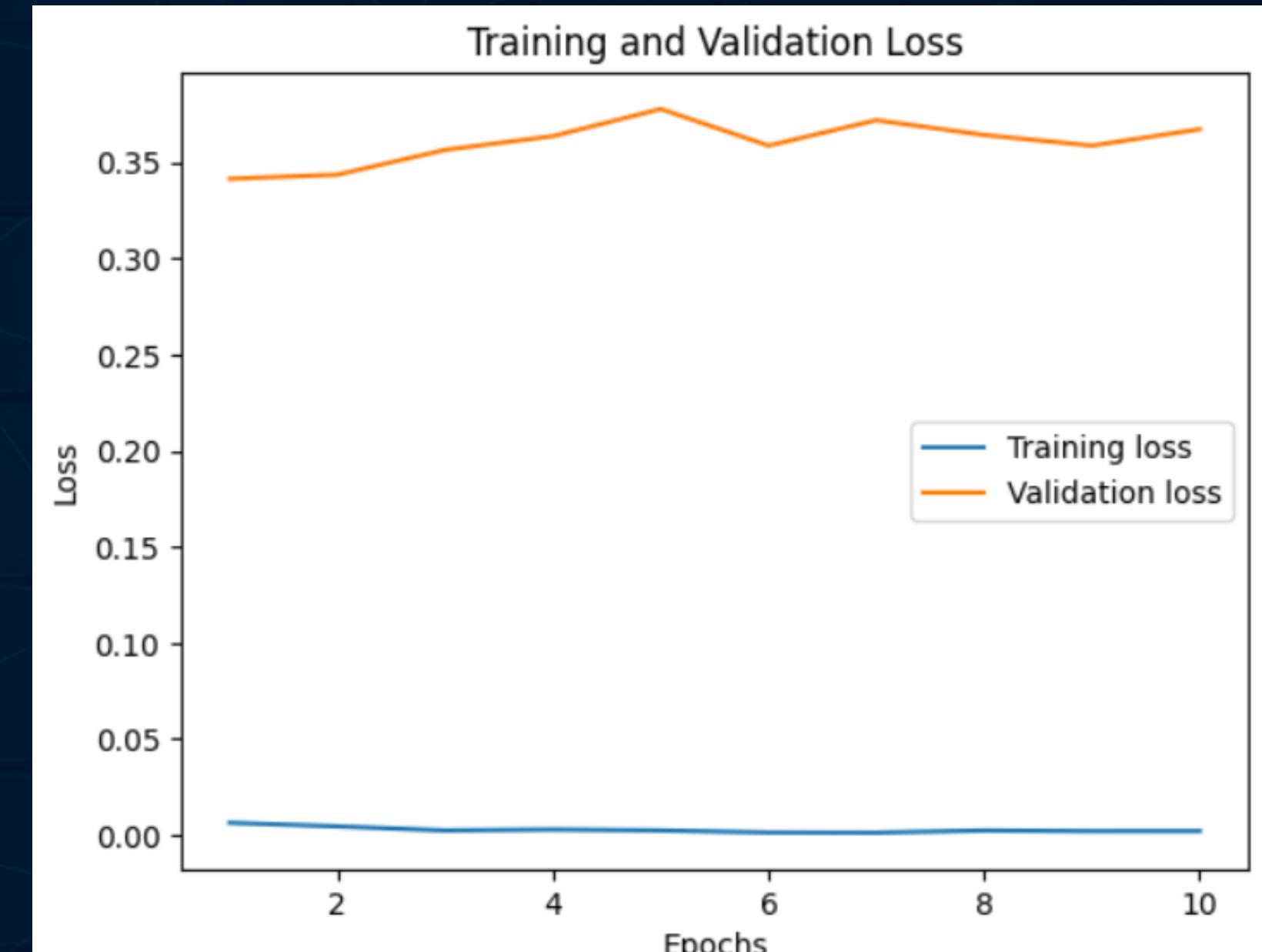
07

MODELS ACCURACY EFFICIENTNETV2B1

GAP BETWEEN TRAINING ACCURACY
AND VALIDATION ACCURACY



GAP BETWEEN TRAINING LOSS
AND VALIDATION LOSS



07

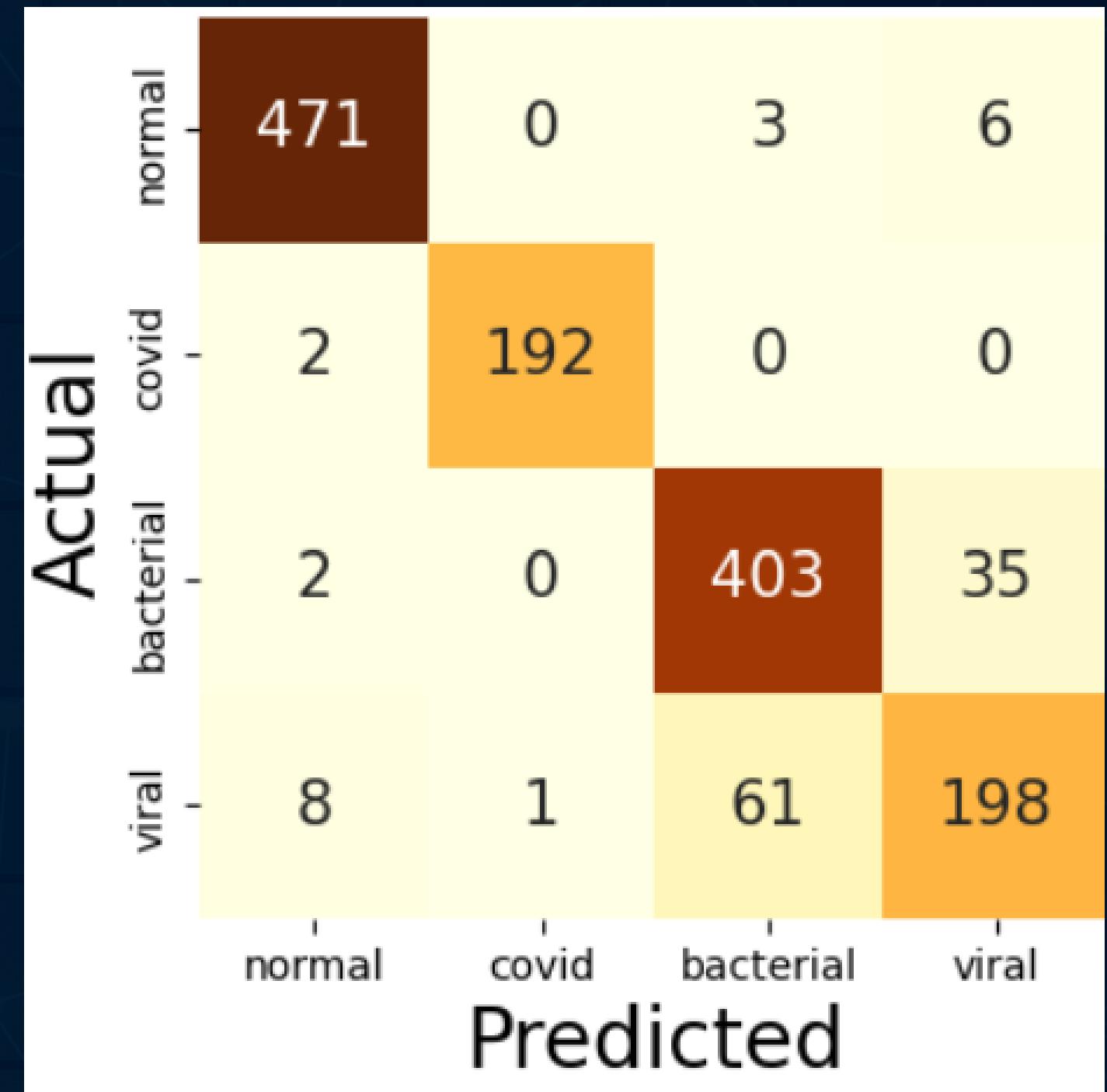
MODELS ACCURACY

EFFICIENTNETV2B1

CLASSIFICATION REPORT

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| normal | 0.9752 | 0.9812 | 0.9782 | 480 |
| covid | 0.9948 | 0.9897 | 0.9922 | 194 |
| bacterial | 0.8630 | 0.9159 | 0.8886 | 440 |
| viral | 0.8285 | 0.7388 | 0.7811 | 268 |
| accuracy | | | 0.9146 | 1382 |
| macro avg | 0.9153 | 0.9064 | 0.9100 | 1382 |
| weighted avg | 0.9137 | 0.9146 | 0.9134 | 1382 |

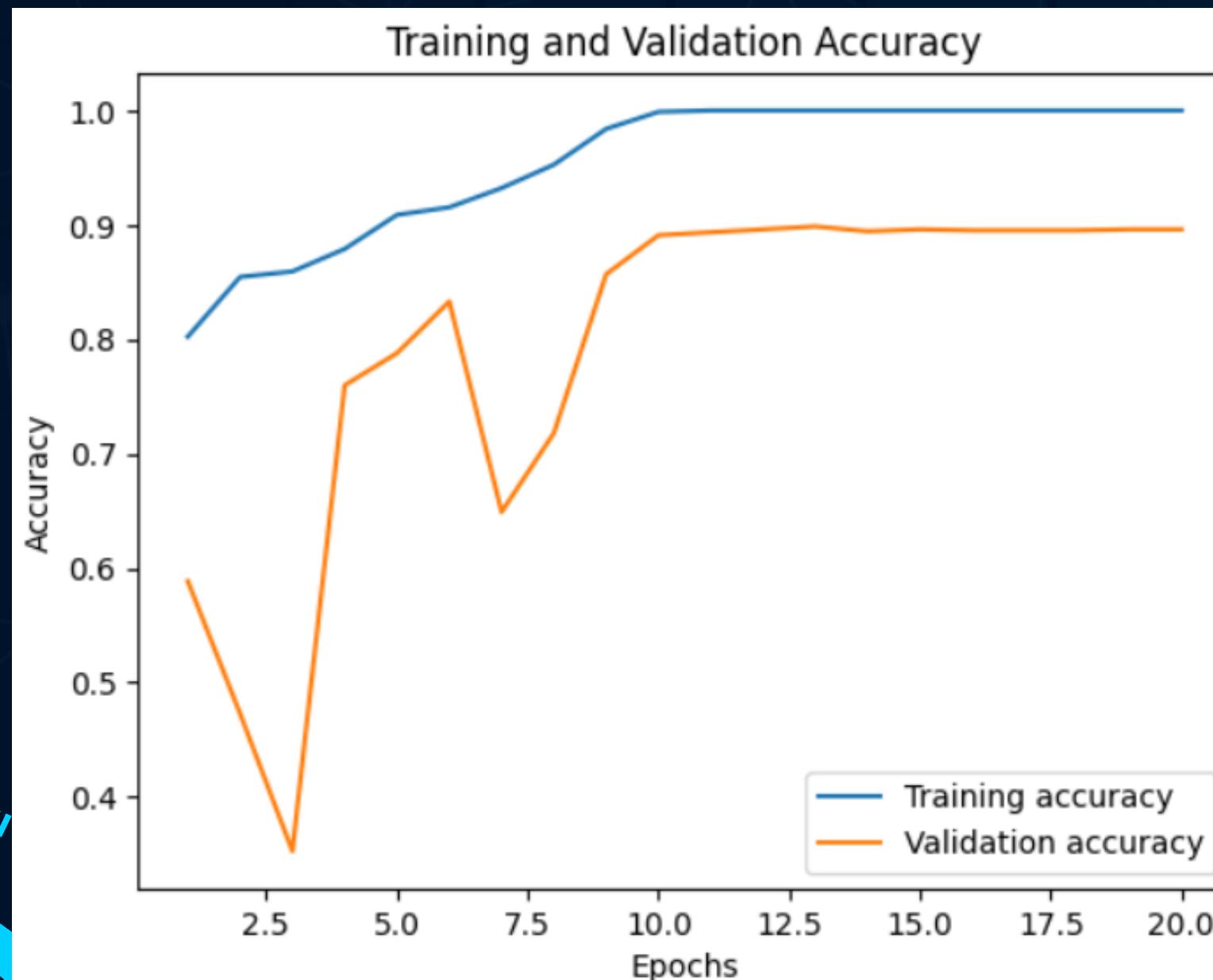
CONFUSION MATRIX



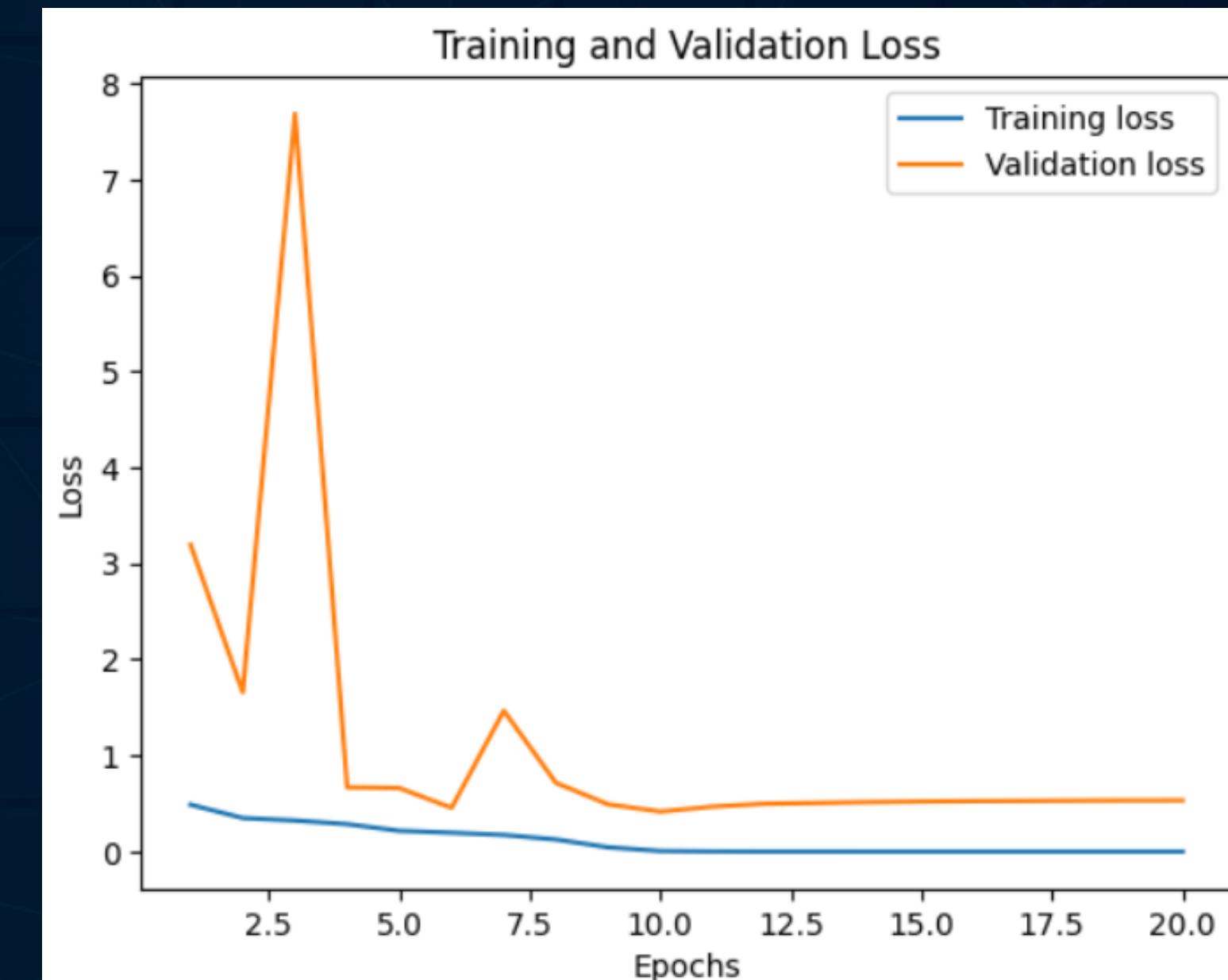
07

MODELS ACCURACY RESNET50V2

GAP BETWEEN TRAINING ACCURACY
AND VALIDATION ACCURACY



GAP BETWEEN TRAINING LOSS
AND VALIDATION LOSS



07

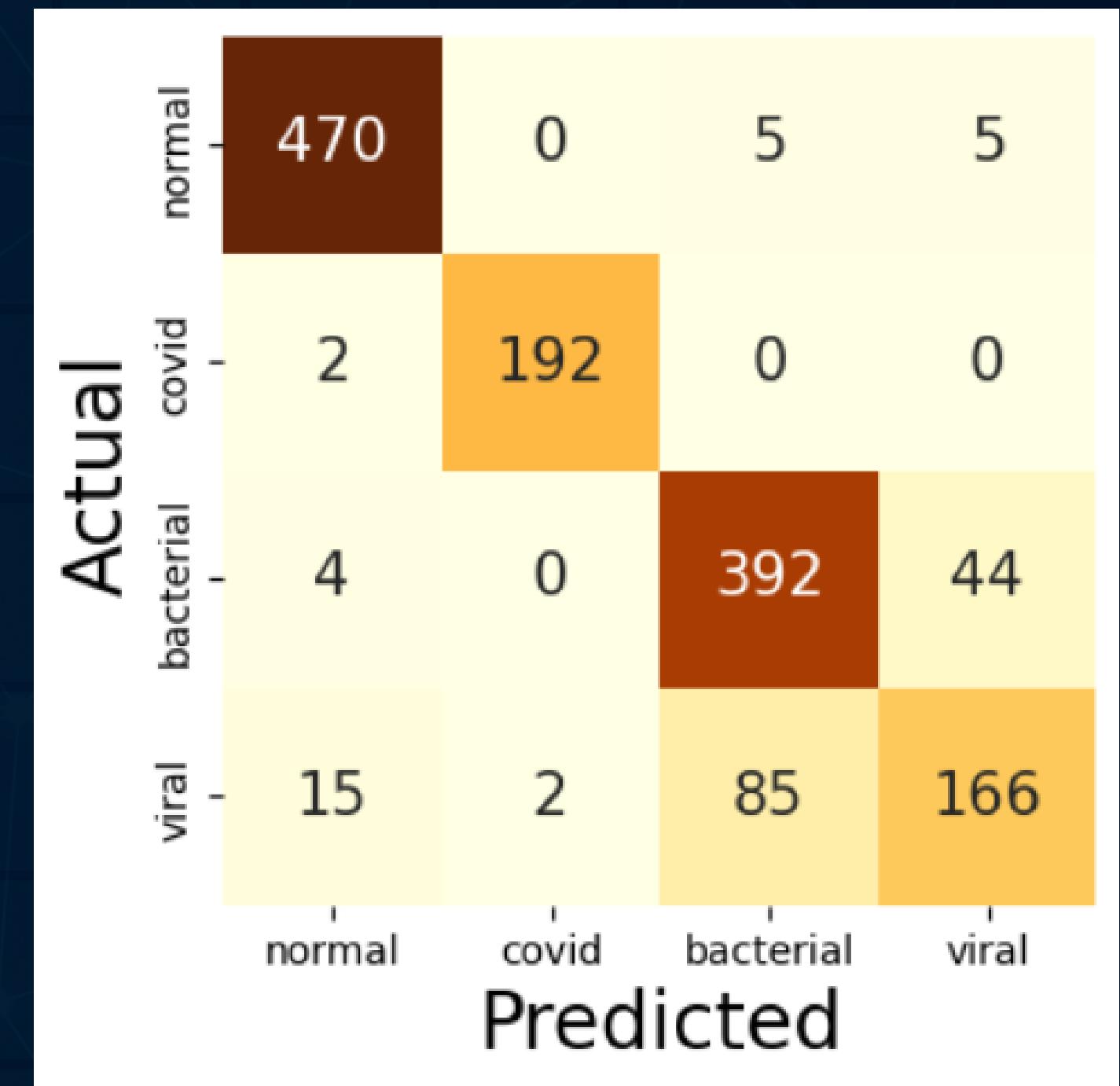
MODELS ACCURACY

RESNET50V2

CLASSIFICATION REPORT

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| normal | 0.9572 | 0.9792 | 0.9681 | 480 |
| covid | 0.9897 | 0.9897 | 0.9897 | 194 |
| bacterial | 0.8133 | 0.8909 | 0.8503 | 440 |
| viral | 0.7721 | 0.6194 | 0.6874 | 268 |
| accuracy | | | 0.8828 | 1382 |
| macro avg | 0.8831 | 0.8698 | 0.8739 | 1382 |
| weighted avg | 0.8801 | 0.8828 | 0.8792 | 1382 |

CONFUSION MATRIX





DEPLOYMENT

Pneumonia Predictor

Select image



R

Predict

Prediction : Pneumonia-Bacterial

Percentage : 1.0

The screenshot shows a user interface titled "Pneumonia Predictor". At the top right is a "Select image" button. Below it is a chest X-ray image with a "R" marker in the top left corner. A "Predict" button is located below the image. The bottom section displays the prediction results: "Prediction : Pneumonia-Bacterial" and "Percentage : 1.0". The background of the slide features a dark blue gradient with a subtle grid pattern and decorative blue circular icons in the corners.



Our Team

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THANKS!

Do you have questions?
Google it 😊

