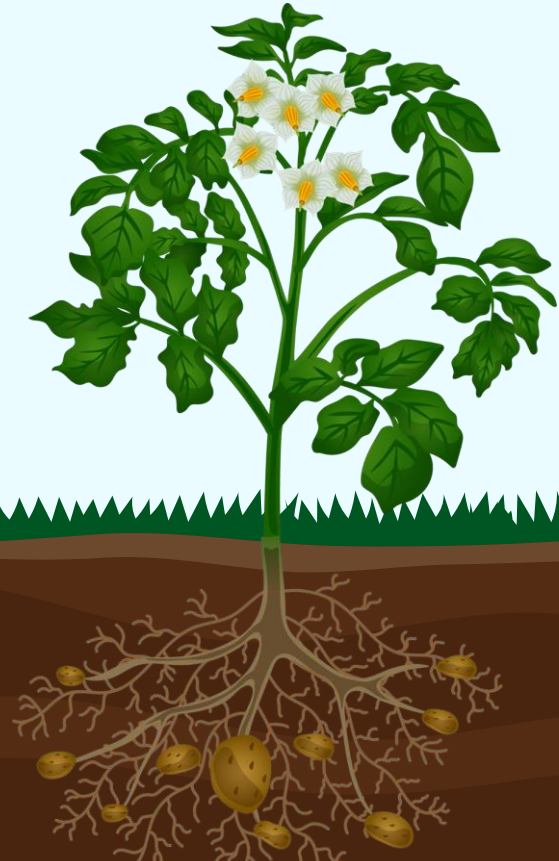


# Plant Disease Detection and Diagnosis System

Team members:

1. George Bebawy
2. Ahmed Hamdy
3. Youssef Ahmed



# Table of contents

**01**

**Intro &  
objectives**

**02**

**Goals of the  
Project**

**03**

**Dataset  
Overview**

**04**

**Data  
Preprocessing**

**05**

**Model  
Architecture  
& Summary**

**06**

**Deployment &  
Results**

01

# Intro & objectives



# Introduction

Our project, **Plant Disease Detection and Diagnosis System**, is designed to assist farmers, gardeners, and plant enthusiasts by leveraging AI to analyze images of plants and provide crucial insights about their health. The system functions in three key steps:

1. **Identifying the Plant Type:** Using image recognition, the system determines the species of the plant and provides a confidence percentage. For instance, it might predict with 90% certainty that the plant is an apple tree.
2. **Assessing Plant Health:** Once the plant type is identified, the system analyzes whether the plant is healthy or displaying symptoms of disease.
3. **Diagnosing the Disease:** If the plant is sick, the system identifies the specific disease and provides a detailed description of it.

By making plant health analysis quick and accessible, our goal is to help protect crops, reduce plant loss, and ensure healthier harvests.

# Objective

The primary objective of this project is to develop a system capable of:

1. **Identifying Plant Types:** Determine plant species from images with high confidence.
2. **Detecting Plant Health:** Quickly assess if a plant is healthy or sick.
3. **Diagnosing Plant Diseases:** Identify specific diseases and provide detailed information for timely treatment.
4. **Displaying Confidence Levels:** Offer percentage-based confidence in predictions to enhance accuracy and decision-making.

This solution aims to provide valuable insights, improve agricultural productivity, and contribute to sustainable farming practices.

02

# Goals of the Project



# Goals

## **User-Friendly Experience:**

Ensure that the platform is easy to use, providing farmers, gardeners, and plant lovers with a simple yet powerful tool for managing plant health.

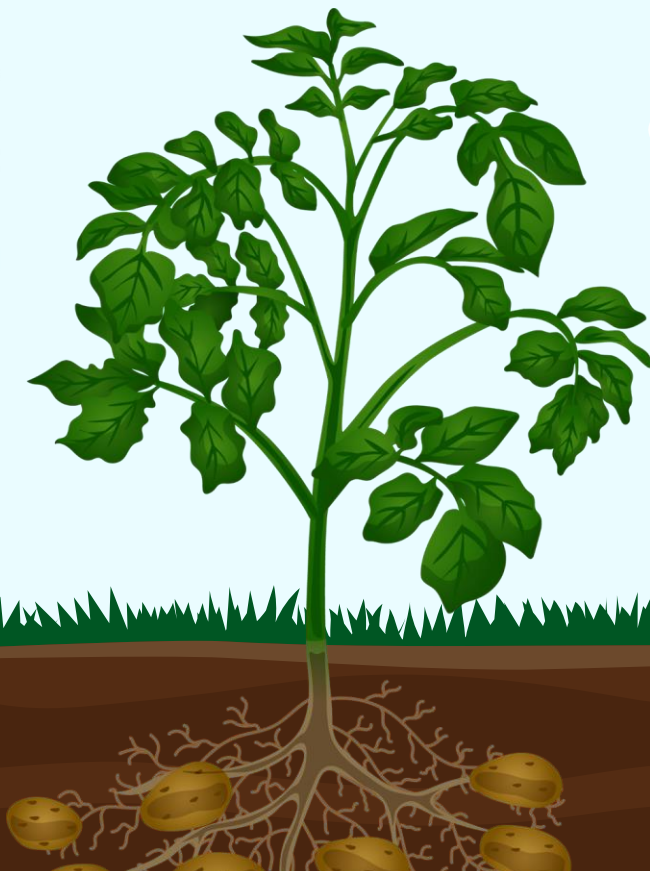
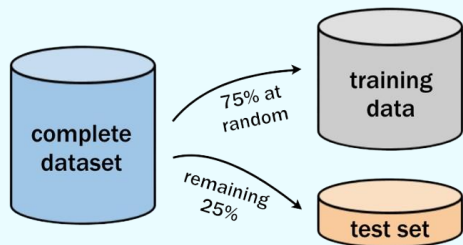
## **Facilitate Real-Time Disease Detection:**

Enable the system to provide real-time analysis so that users can receive instant feedback on plant health and potential diseases, allowing for quicker responses.



03

# Dataset Overview





# Dataset Overview

For the **Plant Disease Detection and Diagnosis System**, we have utilized the New Plant Diseases Dataset from Kaggle. This dataset plays a crucial role in training and testing the model's ability to accurately identify plant types and diagnose diseases.

Key features of the dataset include:

- **Total Images:** The dataset consists of over **87,000 images** of healthy and diseased plant leaves.
- **Plant Categories:** It covers **38 distinct classes**, representing both healthy and diseased plants from various species such as apple, cherry, corn, and grape.
- **Disease Types:** The dataset includes images of plants affected by various diseases like **Apple Scab, Black Rot, and Late Blight**, among others.
- **High-Quality Images:** All images are in **RGB format** and are of high resolution, which ensures that the model can extract fine details for better classification.
- **Diverse Conditions:** The dataset includes images captured in different conditions, adding variability in lighting, angles, and environmental factors, which helps the model generalize well across real-world scenarios.

# Dataset Overview



Apple\_\_Black\_rot



Apple\_\_Cedar\_apple\_rust



Apple\_\_healthy



Blueberry\_\_healthy



Cherry\_(including\_sour)\_\_healthy



Cherry\_(including\_sour)\_\_Powdery\_mildew



Corn\_(maize)\_\_Cercospora\_leaf\_spot  
Gray\_leaf\_spot



Peach\_\_healthy



Pepper,\_bell\_\_Bacterial\_spot



Pepper,\_bell\_\_healthy



Potato\_\_Early\_blight



Potato\_\_healthy



Potato\_\_Late\_blight



Raspberry\_\_healthy

# Dataset Diseases

Apple



Apple-scab



Black-rot



Cedar-apple-rust



healthy

Corn



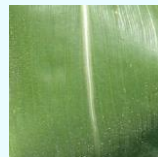
Common-rust



Gray-leaf-spot



Northern-Leaf-Blight



healthy

Tomato



Leaf-Mold



Target-Spot

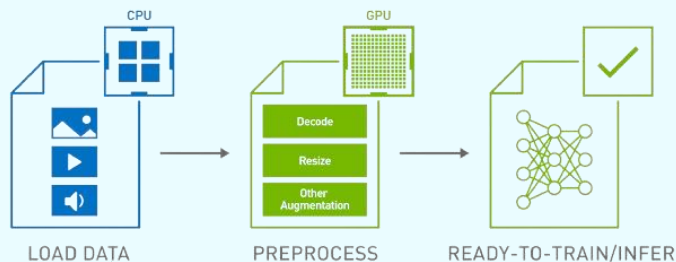


Septoria-leaf-spot



healthy

# Data Preprocessing



# Data Preprocessing

## Data Preprocessing

Effective data preprocessing is a critical step in ensuring that the model can accurately classify plant types and detect diseases.



## Dataset Splitting

The entire dataset is divided into three main subsets to ensure the model is trained, validated, and tested effectively  
Training -- Valid -- Test



# Dataset Splitting

```
import os
import shutil
from sklearn.model_selection import train_test_split

base_dir = '/kaggle/input/plantvillage-dataset/color'
train_dir = 'path_to_save/train'
valid_dir = 'path_to_save/valid'
test_dir = 'path_to_save/test'

os.makedirs(train_dir, exist_ok=True)
os.makedirs(valid_dir, exist_ok=True)
os.makedirs(test_dir, exist_ok=True)

classes = os.listdir(base_dir)

test_size_ratio = 33 / len(classes)
train_valid_classes, test_classes = train_test_split(classes, test_size=test_size_ratio, random_state=42)

valid_size_ratio = 0.2
train_classes, valid_classes = train_test_split(train_valid_classes, test_size=valid_size_ratio, random_state=42)

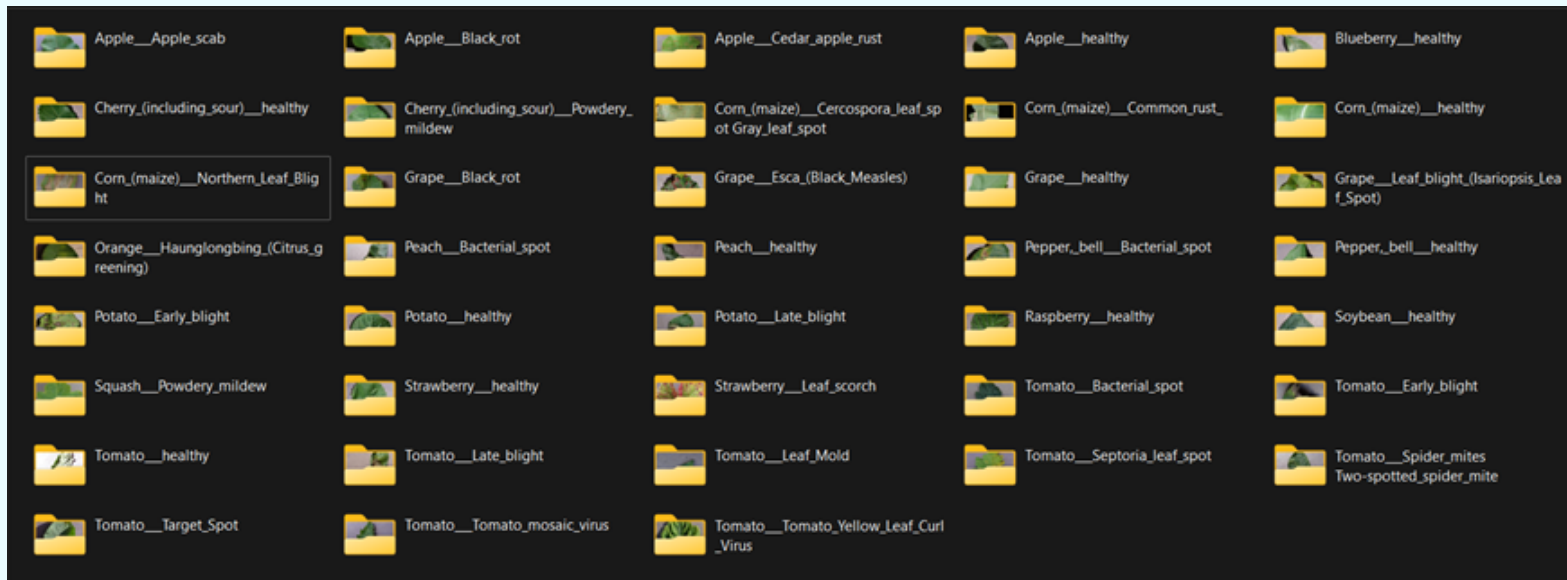
def copy_files(class_list, source_dir, dest_dir):
    for class_name in class_list:
        class_dir = os.path.join(source_dir, class_name)
        dest_class_dir = os.path.join(dest_dir, class_name)
        shutil.copytree(class_dir, dest_class_dir)

copy_files(train_classes, base_dir, train_dir)
copy_files(valid_classes, base_dir, valid_dir)
copy_files(test_classes, base_dir, test_dir)
```

- ▼ /kaggle/working
  - ▼ path\_to\_save
    - ▶ test
    - ▶ train
    - ▶ valid

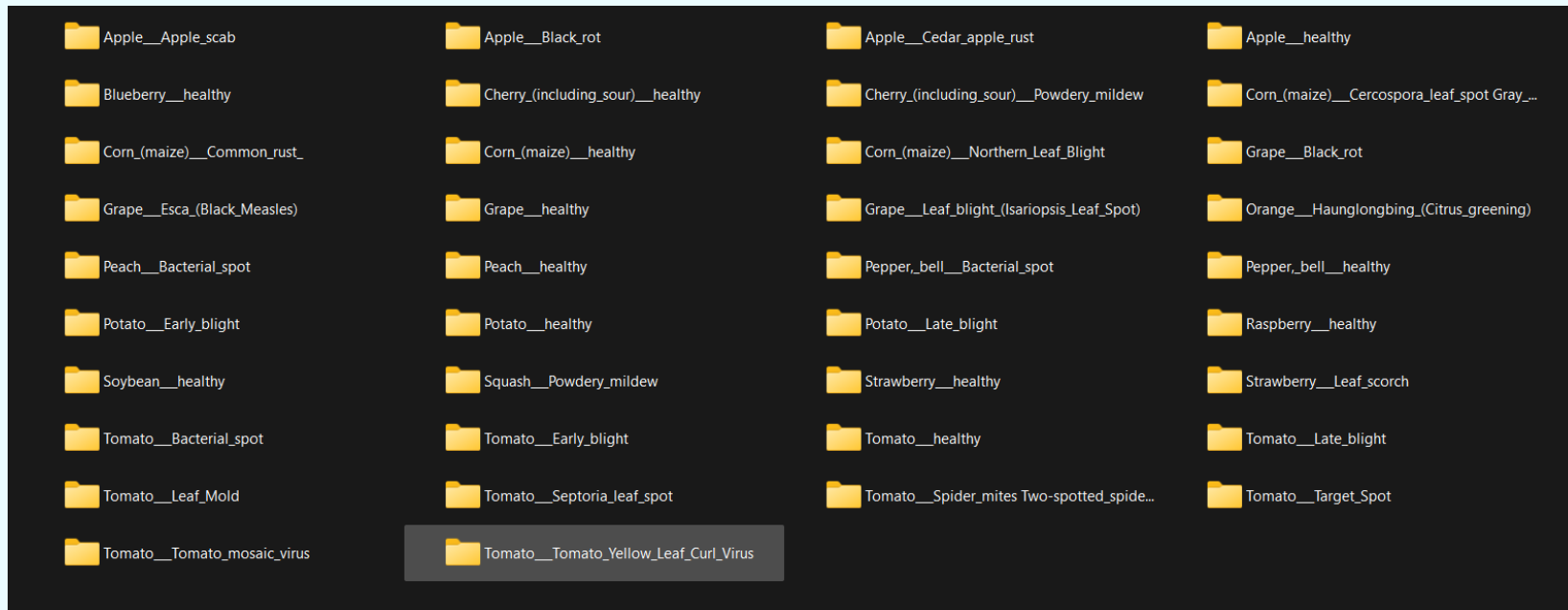
# Dataset Overview

38 different classes “Training data set”



# Dataset Overview

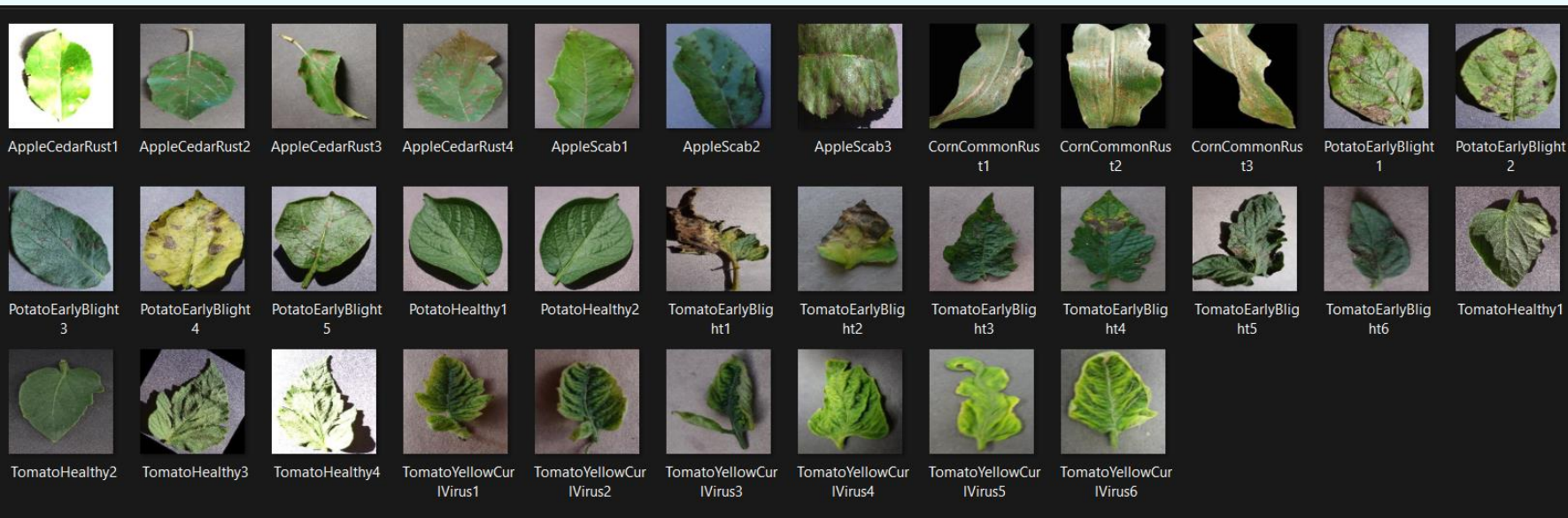
38 different classes “Validation data set”





# Dataset Overview

## Testing Images



05

# Model Architecture



# Model Architecture

## Import library

```
import tensorflow as tf
import matplotlib.pyplot as plt
import pandas as pd
import seaborn as sns
import tensorflow as tf
from tensorflow.keras import layers, models
```

## Data Preprocessing

```
# first We Load training dataset #
```

```
training_set = tf.keras.utils.image_dataset_from_directory(
    'train',
    labels="inferred",
    label_mode="categorical",
    class_names=None,
    color_mode="rgb",
    batch_size=32,
    image_size=(128, 128),
    shuffle=True,
    seed=None,
    validation_split=None,
    subset=None,
    interpolation="bilinear",
    follow_links=False,
    crop_to_aspect_ratio=False
)
print(f'Found {training_set.cardinality().numpy()} batches of training data')
```

```
Found 78295 files belonging to 38 classes.
Found 2197 batches of training data
```

# Model Architecture

```
# second we load validation dataset #
```

```
validation_set = tf.keras.utils.image_dataset_from_directory(
    'valid',
    labels="inferred",
    label_mode="categorical",
    class_names=None,
    color_mode="rgb",
    batch_size=32,
    image_size=(128, 128),
    shuffle=True,
    seed=None,
    validation_split=None,
    subset=None,
    interpolation="bilinear",
    follow_links=False,
    crop_to_aspect_ratio=False
)
print(f'Found {validation_set.cardinality().numpy()} batches of validation data')
```

Found 17572 files belonging to 38 classes.

Found 550 batches of validation data

## Attention Mechanism

```
# Squeeze-and-Excitation block definition #
```

```
# Squeeze-and-Excitation block definition
def se_block(input_tensor, ratio=16):
    filters = input_tensor.shape[-1] # Number of filters in the input tensor
    se_shape = (1, 1, filters)

    se = layers.GlobalAveragePooling2D()(input_tensor)
    se = layers.Reshape(se_shape)(se)
    se = layers.Dense(filters // ratio, activation='relu', use_bias=False)(se)
    se = layers.Dense(filters, activation='sigmoid', use_bias=False)(se)
    x = layers.multiply([input_tensor, se])
    return x
```

# Model Architecture

## CNN

### CNN Architecture

```
# Input Layer
inputs = layers.Input(shape=(128, 128, 3))

# First Convolutional Layer Block
x = layers.Conv2D(filters=32, kernel_size=3, padding='same', activation='relu')(inputs)
x = layers.Conv2D(filters=32, kernel_size=3, activation='relu')(x)
x = layers.MaxPooling2D(pool_size=2, strides=2)(x)

# Attention block after the first convolutional Layer
x = se_block(x)

# Second Convolutional Layer Block
x = layers.Conv2D(filters=64, kernel_size=3, padding='same', activation='relu')(x)
x = layers.Conv2D(filters=64, kernel_size=3, activation='relu')(x)
x = layers.MaxPooling2D(pool_size=2, strides=2)(x)

# Attention block after the second convolutional Layer
x = se_block(x)

# Third Convolutional Layer Block
x = layers.Conv2D(filters=128, kernel_size=3, padding='same', activation='relu')(x)
x = layers.Conv2D(filters=128, kernel_size=3, activation='relu')(x)
x = layers.MaxPooling2D(pool_size=2, strides=2)(x)

# Fourth Convolutional Layer Block
x = layers.Conv2D(filters=256, kernel_size=3, padding='same', activation='relu')(x)
x = layers.Conv2D(filters=256, kernel_size=3, activation='relu')(x)
x = layers.MaxPooling2D(pool_size=2, strides=2)(x)

# Fifth Convolutional Layer Block
x = layers.Conv2D(filters=512, kernel_size=3, padding='same', activation='relu')(x)
x = layers.Conv2D(filters=512, kernel_size=3, activation='relu')(x)
x = layers.MaxPooling2D(pool_size=2, strides=2)(x)
```

# Model Architecture

# DNN

## DNN Block

```
# Flattening
x = layers.Dropout(0.25)(x)
x = layers.Flatten()(x)

# DNN Block (Dense Layer)
x = layers.Dense(units=1500, activation='relu')(x)
x = layers.Dropout(0.4)(x)

# Output Layer
outputs = layers.Dense(units=38, activation='softmax')(x)
```

## Model Compilation and Summary

```
# Compiling the model with the current Adam optimizer
model.compile(optimizer=tf.keras.optimizers.Adam(learning_rate=0.0001),
              loss='categorical_crossentropy',
              metrics=['accuracy'])

# Model summary
model.summary()
```

# Model Summary

Layer (type)	Output Shape	Param #	Connected to
input_layer_1 (InputLayer)	(None, 128, 128, 3)	0	-
conv2d_2 (Conv2D)	(None, 128, 128, 32)	896	input_layer_1[0]...
conv2d_3 (Conv2D)	(None, 126, 126, 32)	9,248	conv2d_2[0][0]
max_pooling2d_1 (MaxPooling2D)	(None, 63, 63, 32)	0	conv2d_3[0][0]
global_average_pooling2d_1 (GlobalAveragePooling2D)	(None, 32)	0	max_pooling2d_1[0]...
reshape_1 (Reshape)	(None, 1, 1, 32)	0	global_average_pooling2d_1[0]...
dense_2 (Dense)	(None, 1, 1, 2)	64	reshape_1[0][0]
dense_3 (Dense)	(None, 1, 1, 32)	64	dense_2[0][0]
multiply_1 (Multiply)	(None, 63, 63, 32)	0	max_pooling2d_1[0]...
conv2d_4 (Conv2D)	(None, 63, 63, 64)	18,496	multiply_1[0][0]
conv2d_5 (Conv2D)	(None, 61, 61, 64)	36,928	conv2d_4[0][0]
max_pooling2d_2 (MaxPooling2D)	(None, 30, 30, 64)	0	conv2d_5[0][0]
global_average_pooling2d_2 (GlobalAveragePooling2D)	(None, 64)	0	max_pooling2d_2[0]...
reshape_2 (Reshape)	(None, 1, 1, 64)	0	global_average_pooling2d_2[0]...
dense_4 (Dense)	(None, 1, 1, 4)	256	reshape_2[0][0]
dense_5 (Dense)	(None, 1, 1, 64)	256	dense_4[0][0]
multiply_2 (Multiply)	(None, 30, 30, 64)	0	max_pooling2d_2[0]...

conv2d_6 (Conv2D)	(None, 30, 30, 128)	73,856	multiply_2[0][0]
conv2d_7 (Conv2D)	(None, 28, 28, 128)	147,584	conv2d_6[0][0]
max_pooling2d_3 (MaxPooling2D)	(None, 14, 14, 128)	0	conv2d_7[0][0]
conv2d_8 (Conv2D)	(None, 14, 14, 256)	295,168	max_pooling2d_3[0]...
conv2d_9 (Conv2D)	(None, 12, 12, 256)	590,080	conv2d_8[0][0]
max_pooling2d_4 (MaxPooling2D)	(None, 6, 6, 256)	0	conv2d_9[0][0]
conv2d_10 (Conv2D)	(None, 6, 6, 512)	1,180,160	max_pooling2d_4[0]...
conv2d_11 (Conv2D)	(None, 4, 4, 512)	2,359,808	conv2d_10[0][0]
max_pooling2d_5 (MaxPooling2D)	(None, 2, 2, 512)	0	conv2d_11[0][0]
dropout (Dropout)	(None, 2, 2, 512)	0	max_pooling2d_5[0]...
flatten (Flatten)	(None, 2048)	0	dropout[0][0]
dense_6 (Dense)	(None, 1500)	3,073,500	flatten[0][0]
dropout_1 (Dropout)	(None, 1500)	0	dense_6[0][0]
dense_7 (Dense)	(None, 38)	57,038	dropout_1[0][0]

Total params: 7,843,402 (29.92 MB)

Trainable params: 7,843,402 (29.92 MB)

Non-trainable params: 0 (0.00 B)

# Model Summary

```
# Training the model
training_history = model.fit(x=training_set,
                             validation_data=validation_set,
                             epochs=10)
```

```
Epoch 1/10
2197/2197 ----- 960s 436ms/step - accuracy: 0.3804 - loss: 2.1861 - val_accuracy: 0.8352 - val_loss: 0.5128
Epoch 2/10
2197/2197 ----- 929s 423ms/step - accuracy: 0.8321 - loss: 0.5318 - val_accuracy: 0.9066 - val_loss: 0.2836
Epoch 3/10
2197/2197 ----- 929s 423ms/step - accuracy: 0.9004 - loss: 0.3047 - val_accuracy: 0.9190 - val_loss: 0.2482
Epoch 4/10
2197/2197 ----- 940s 428ms/step - accuracy: 0.9306 - loss: 0.2078 - val_accuracy: 0.9495 - val_loss: 0.1617
Epoch 5/10
2197/2197 ----- 936s 426ms/step - accuracy: 0.9491 - loss: 0.1535 - val_accuracy: 0.9544 - val_loss: 0.1416
Epoch 6/10
2197/2197 ----- 933s 425ms/step - accuracy: 0.9621 - loss: 0.1162 - val_accuracy: 0.9556 - val_loss: 0.1332
Epoch 7/10
2197/2197 ----- 928s 422ms/step - accuracy: 0.9673 - loss: 0.0983 - val_accuracy: 0.9374 - val_loss: 0.2075
Epoch 8/10
2197/2197 ----- 927s 422ms/step - accuracy: 0.9720 - loss: 0.0842 - val_accuracy: 0.9657 - val_loss: 0.1117
Epoch 9/10
2197/2197 ----- 926s 421ms/step - accuracy: 0.9788 - loss: 0.0670 - val_accuracy: 0.9548 - val_loss: 0.1558
Epoch 10/10
2197/2197 ----- 946s 431ms/step - accuracy: 0.9804 - loss: 0.0615 - val_accuracy: 0.9627 - val_loss: 0.1187
```

## Cheak The accuracy

```
# Evaluate Training Set Accuracy
train_loss, train_acc = model.evaluate(training_set)
print('Training accuracy:', train_acc)
```

```
# Evaluate Validation Set Accuracy
val_loss, val_acc = model.evaluate(validation_set)
print('Validation accuracy:', val_acc)
```

```
2197/2197 ----- 228s 101ms/step - accuracy: 0.9880 - loss: 0.0335
Training accuracy: 0.9892311096191406
550/550 ----- 57s 103ms/step - accuracy: 0.9629 - loss: 0.1235
Validation accuracy: 0.9626678824424744
```

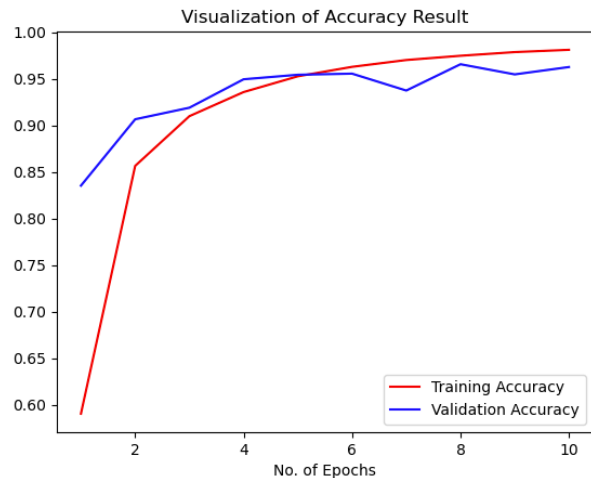
Training accuracy: 0.9892311096191406  
550/550 ----- 57s 103ms/s  
Validation accuracy: 0.9626678824424744



# Model Summary

## Plotting the accuracy results

```
epochs = [i for i in range(1,11)]  
plt.plot(epochs,training_history.history['accuracy'],color='red',label='Training Accuracy')  
plt.plot(epochs,training_history.history['val_accuracy'],color='blue',label='Validation Accuracy')  
plt.xlabel('No. of Epochs')  
plt.title('Visualization of Accuracy Result')  
plt.legend()  
plt.show()
```



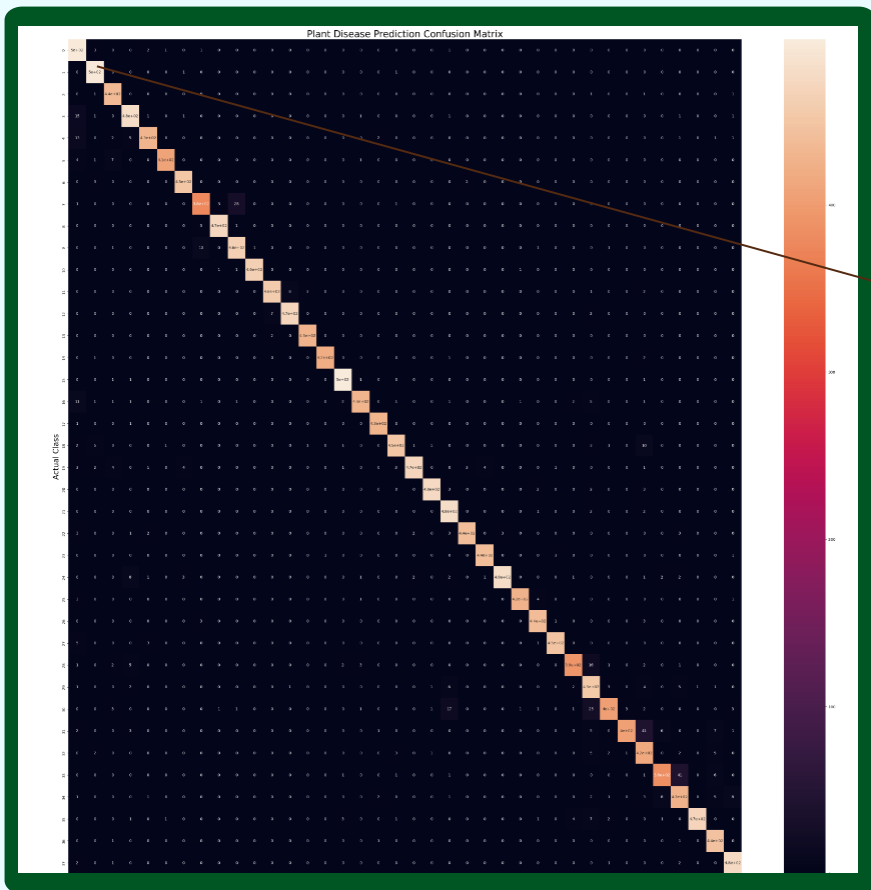
# Model Summary

	precision	recall	f1-score	support
Apple__Apple_scab	0.88	0.98	0.93	504
Apple__Black_rot	0.96	1.00	0.98	497
Apple__Cedar_apple_rust	0.95	0.99	0.97	448
Apple__healthy	0.94	0.96	0.95	502
Blueberry__healthy	0.97	0.94	0.96	454
Cherry_(including_sour)__Powdery_mildew	0.99	0.97	0.98	421
Cherry_(including_sour)__healthy	0.98	0.99	0.98	456
Corn_(maize)__Cercospora_leaf_spot_Gray_leaf_spot	0.95	0.92	0.94	418
Corn_(maize)__Common_rust	0.99	0.99	0.99	477
Corn_(maize)__Northern_Leaf_Blight	0.94	0.97	0.95	477
Corn_(maize)__healthy	1.00	0.99	1.00	465
Grape__Black_rot	0.98	0.97	0.98	472
Grape__Esca_(Black_Measles)	0.98	0.98	0.98	480
Grape__Leaf_blight_(Isariopsis_Leaf_Spot)	0.99	1.00	0.99	438
Grape__healthy	1.00	0.99	0.99	423
Orange__Haunglongbing_(Citrus_greening)	0.98	0.99	0.99	503
Peach__Bacterial_spot	0.98	0.94	0.96	459
Peach__healthy	0.99	1.00	0.99	432
Pepper,_bell__Bacterial_spot	0.98	0.95	0.96	478
Pepper,_bell__healthy	0.99	0.95	0.97	497
Potato__Early_blight	0.99	0.98	0.99	485
Potato__Late_blight	0.93	0.99	0.96	485
Potato__healthy	0.99	0.96	0.98	456
Raspberry__healthy	0.99	0.99	0.99	445
Soybean__healthy	0.99	0.95	0.97	505
Squash__Powdery_mildew	0.99	0.98	0.99	434
Strawberry__Leaf_scorch	0.98	0.98	0.98	444
Strawberry__healthy	0.99	0.98	0.98	456
Tomato__Bacterial_spot	0.98	0.92	0.95	425
Tomato__Early_blight	0.86	0.94	0.90	488
Tomato__Late_blight	0.97	0.87	0.92	463
Tomato__Leaf_Mold	0.99	0.86	0.92	478
Tomato__Septoria_leaf_spot	0.84	0.95	0.89	436
Tomato__Spider_mites_Two-spotted_spider_mite	0.96	0.88	0.92	435
Tomato__Target_Spot	0.88	0.93	0.90	457
Tomato__Tomato_Yellow_Leaf_Curl_Virus	1.00	0.96	0.98	498
Tomato__Tomato_mosaic_virus	0.95	0.99	0.97	448
Tomato__healthy	0.97	0.99	0.98	481
accuracy			0.96	17572
macro avg	0.96	0.96	0.96	17572
weighted avg	0.96	0.96	0.96	17572

	precision	recall	f1-score	support
Apple__Apple_scab	0.88	0.98	0.93	504

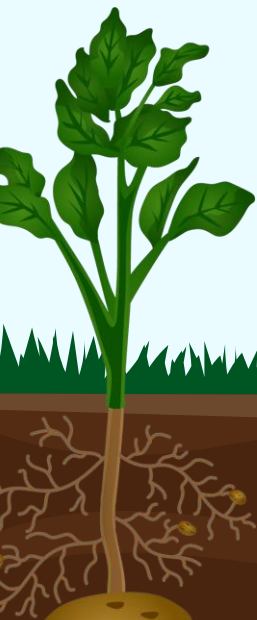
# Model Summary

C.M



	Predicted 0	Predicted 1	Predicted 2	Predicted 3
Actual 0	500	3	0	2
Actual 1	0	497	0	0
Actual 2	0	1	440	0
Actual 3	15	1	0	459

# Model Deployment



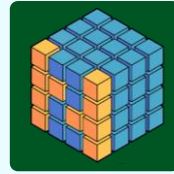
# Model Deployment



**Streamlit**



**Tensor Flow**

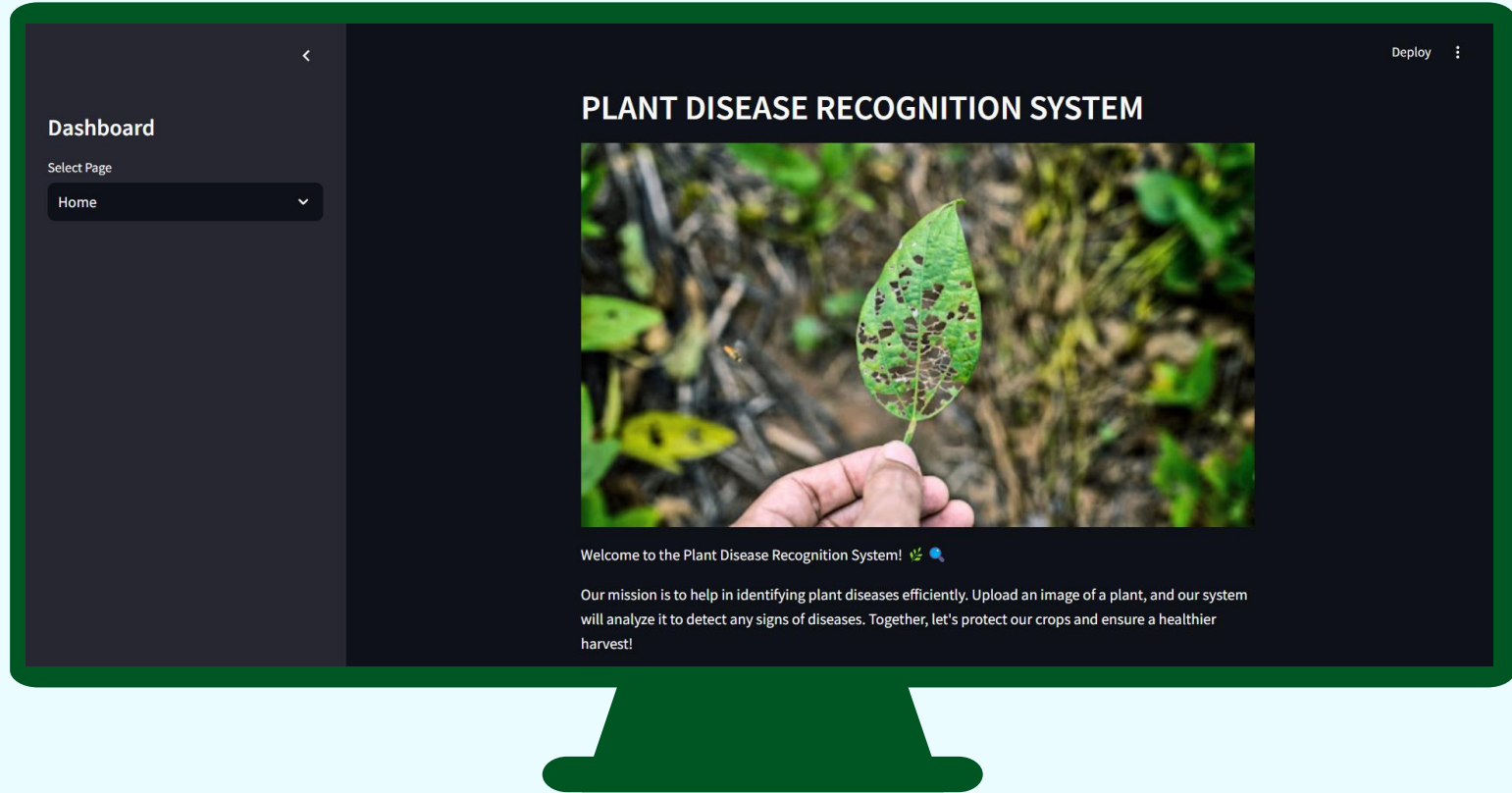


**NumPy**

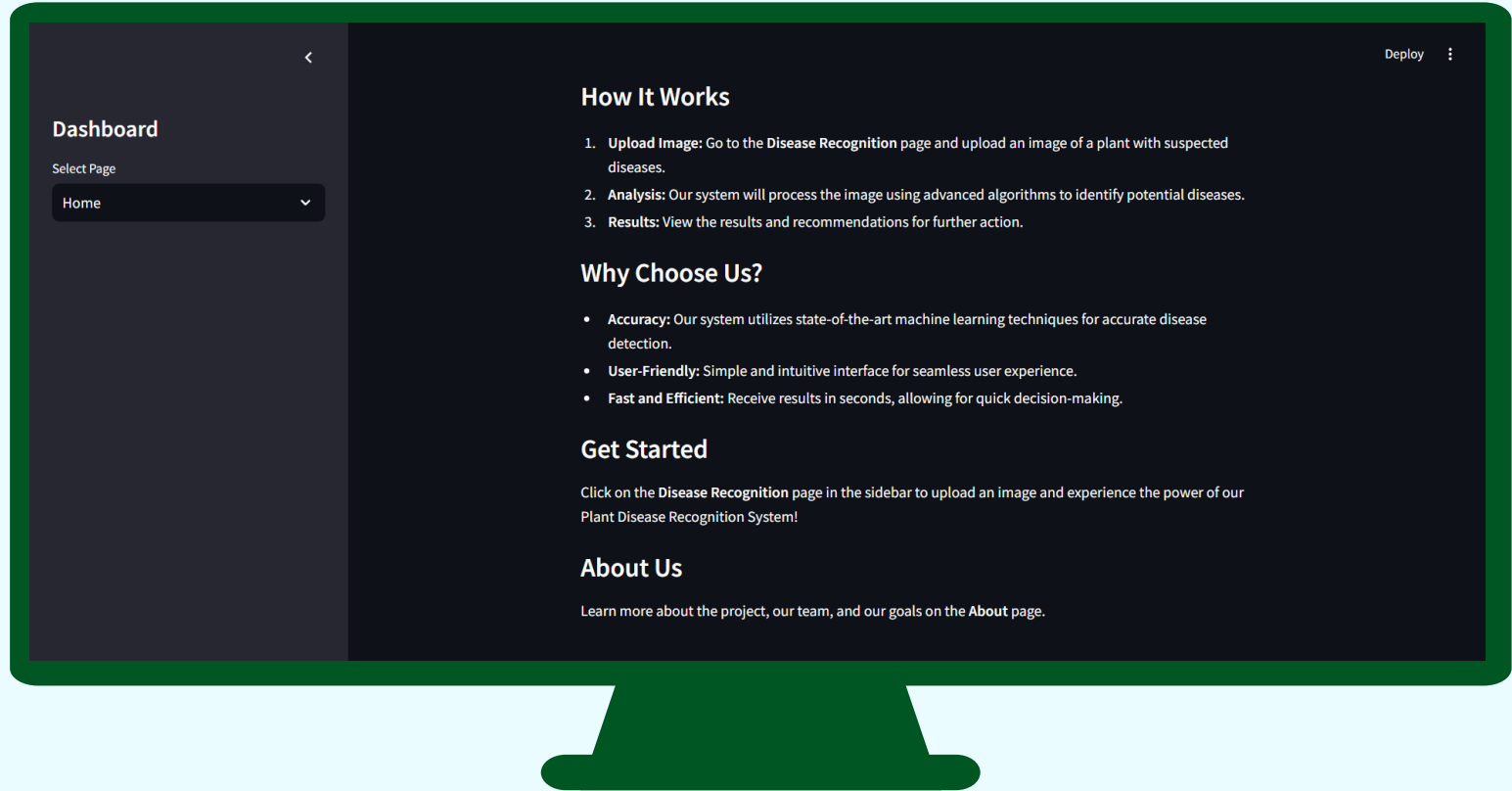


**Keras**

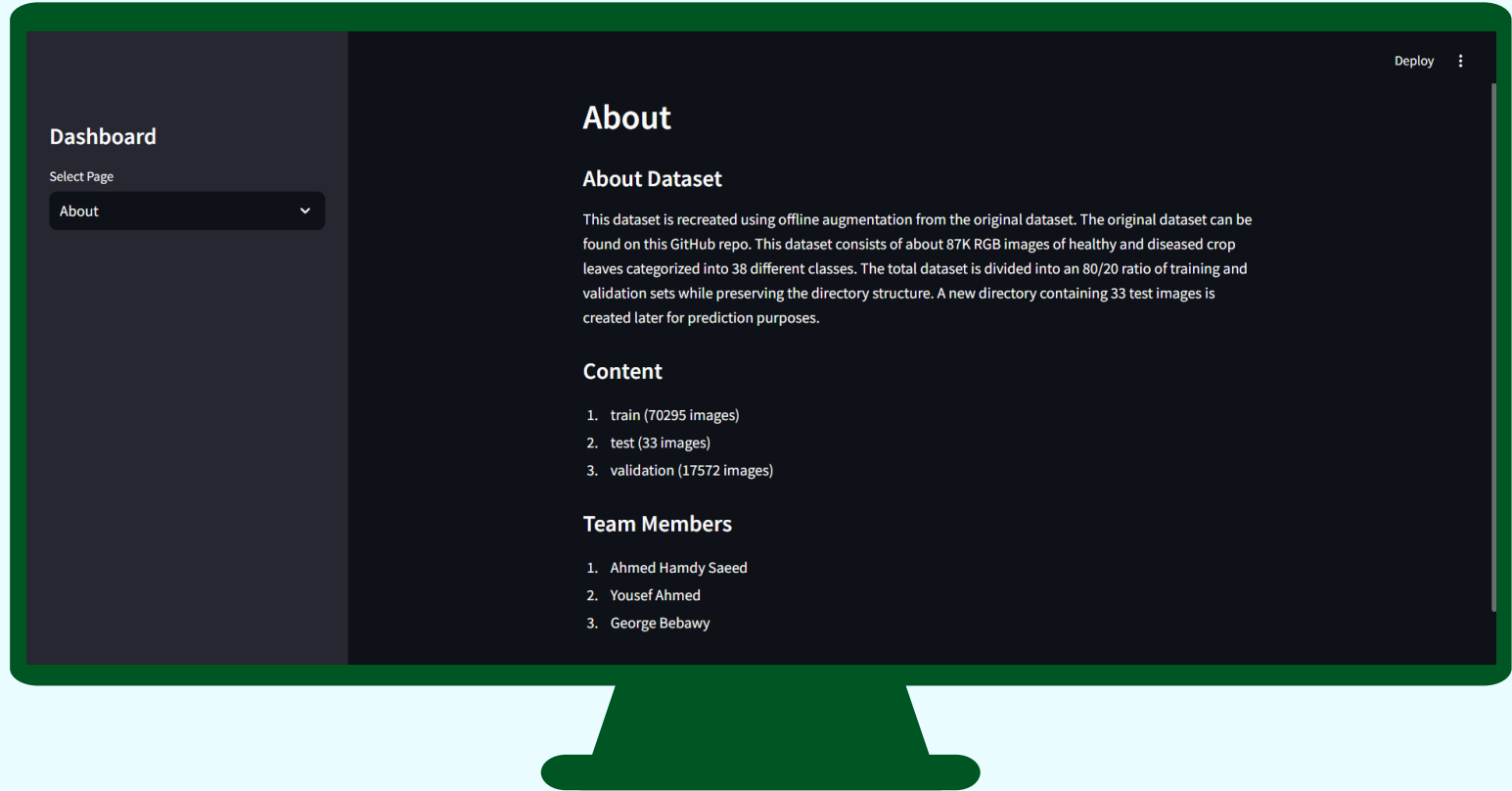
# Model Deployment



# Model Deployment

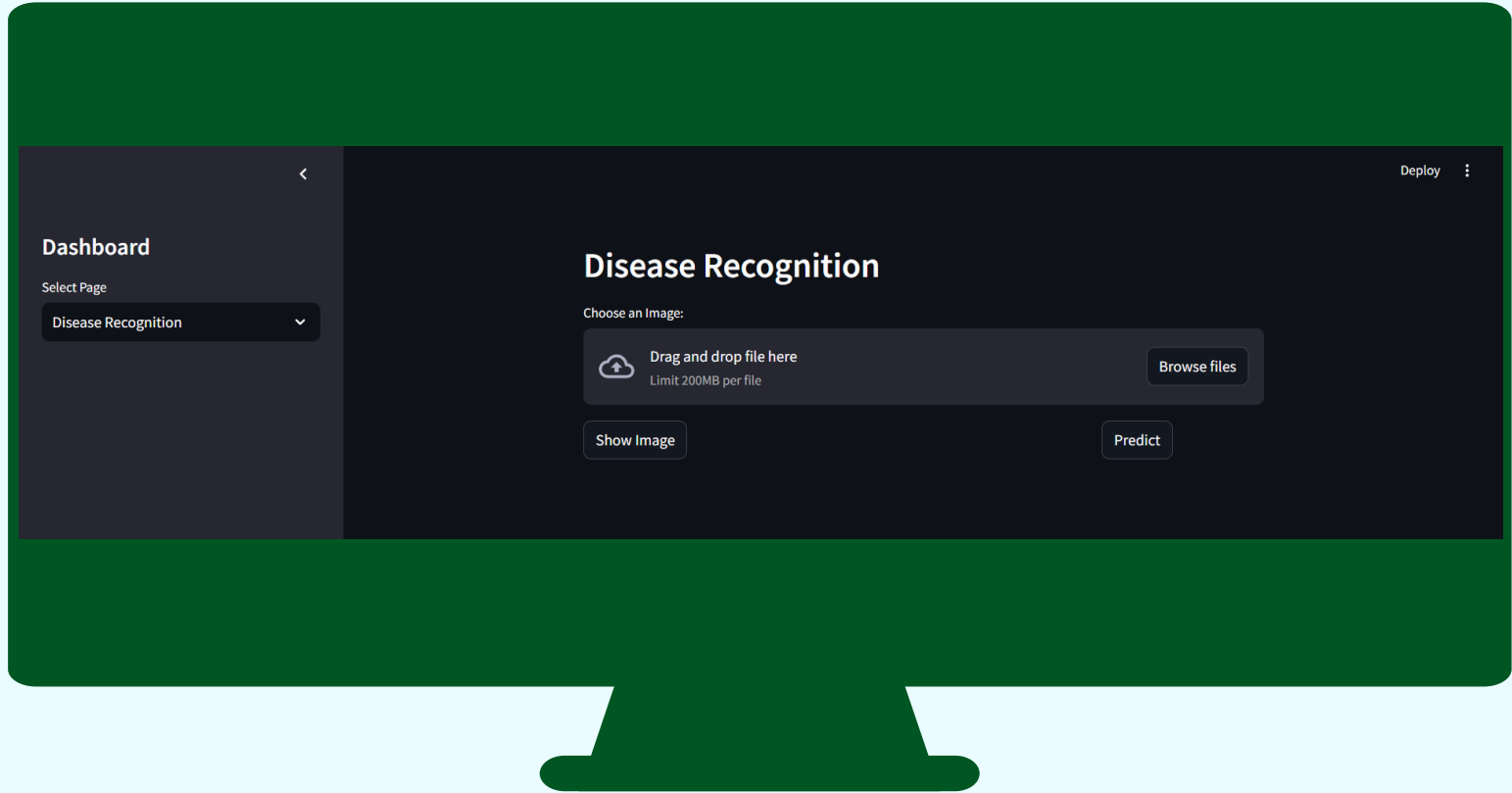


# Model Deployment

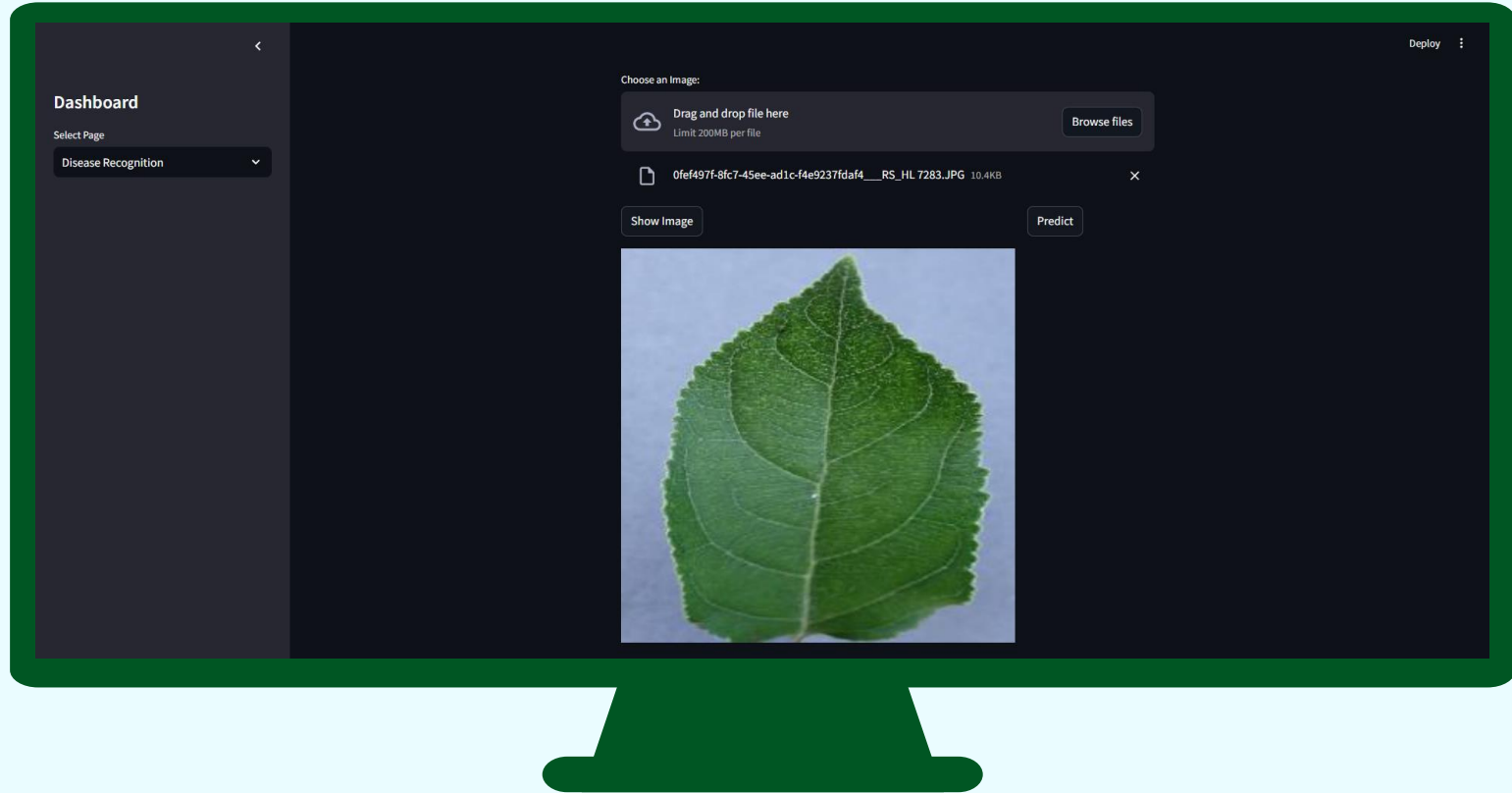




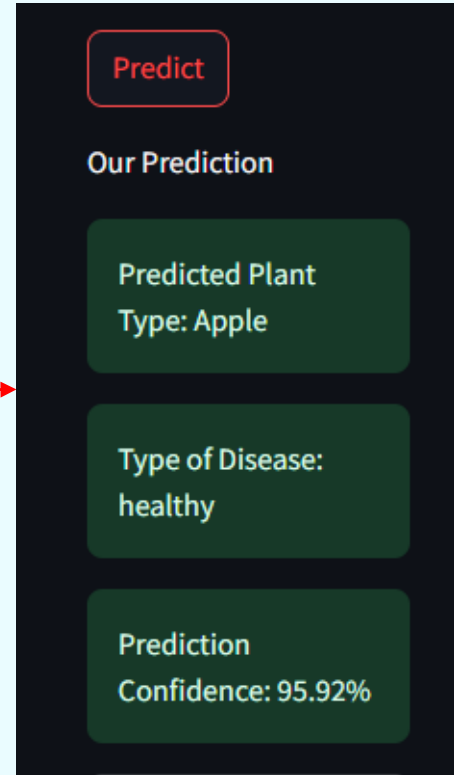
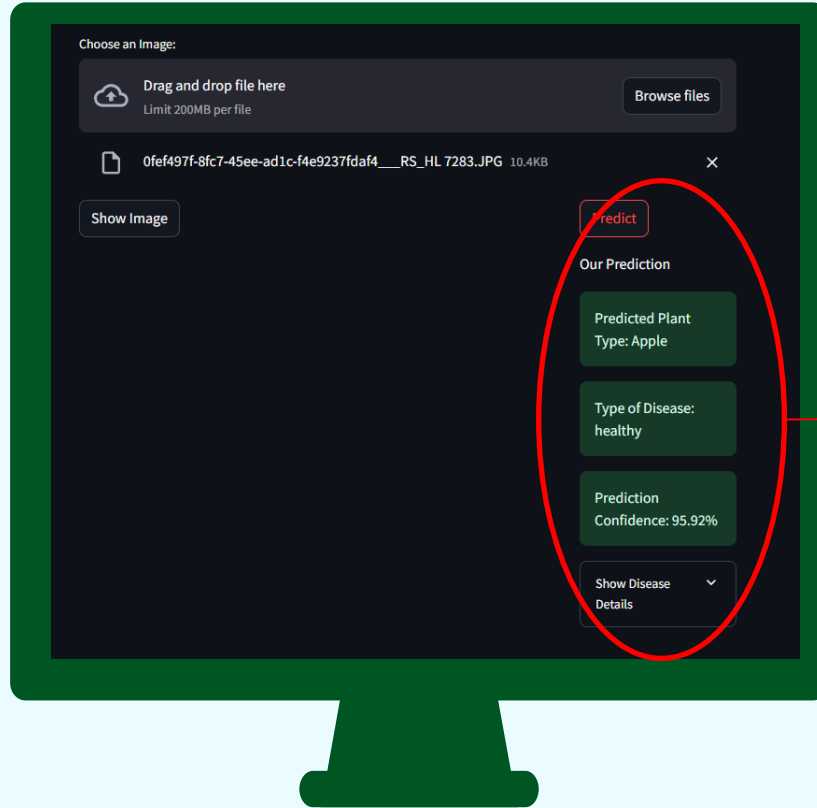
# Model Deployment



# Model Deployment



# Model Deployment



# Thanks!

Do you have any questions?

