

```
In [1]: import numpy as np
import tensorflow as tf
from keras.preprocessing.image import ImageDataGenerator
import matplotlib.pyplot as plt
```

loading model

```
In [2]: model = tf.keras.models.load_model('trained_model.keras')
```

```
In [3]: model.summary()

Model: "sequential"

Layer (type)                Output Shape                Param #
=====
conv2d (Conv2D)              (None, 128, 128, 32)        896
conv2d_1 (Conv2D)             (None, 126, 126, 32)        9248
max_pooling2d (MaxPooling2D) (None, 63, 63, 32)          0
conv2d_2 (Conv2D)             (None, 63, 63, 64)          18496
conv2d_3 (Conv2D)             (None, 61, 61, 64)          36928
max_pooling2d_1 (MaxPooling2D) (None, 30, 30, 64)          0
conv2d_4 (Conv2D)             (None, 30, 30, 128)         73856
conv2d_5 (Conv2D)             (None, 28, 28, 128)         147584
max_pooling2d_2 (MaxPooling2D) (None, 14, 14, 128)          0
conv2d_6 (Conv2D)             (None, 14, 14, 256)         295168
conv2d_7 (Conv2D)             (None, 12, 12, 256)         590080
max_pooling2d_3 (MaxPooling2D) (None, 6, 6, 256)           0
conv2d_8 (Conv2D)             (None, 6, 6, 512)          1180160
conv2d_9 (Conv2D)             (None, 4, 4, 512)           2359808
max_pooling2d_4 (MaxPooling2D) (None, 2, 2, 512)           0
dropout (Dropout)            (None, 2, 2, 512)           0
flatten (Flatten)             (None, 2048)                 0
dense (Dense)                 (None, 1500)                 3073500
dropout_1 (Dropout)           (None, 1500)                 0
dense_1 (Dense)               (None, 38)                   57038

Total params: 7,842,762
Trainable params: 7,842,762
Non-trainable params: 0
```

```
In [ ]:
```

Visualising images of test set

```
In [4]: import cv2
image_path = "test/test/CornCommonRust2.JPG"
#read img
img = cv2.imread(image_path)
#convert bgr to rgb
img = cv2.cvtColor(img,cv2.COLOR_BGR2RGB)

#Displaying images
plt.imshow(img)
plt.title("Test Image")
plt.xticks([])
plt.yticks([])
plt.show()
```



Testing Model

```
In [5]: image = tf.keras.preprocessing.image.load_img(image_path,target_size=(128,128))
input_arr =tf.keras.preprocessing.image.img_to_array(image)
input_arr =np.array([input_arr])#convert single image to a batch
print(input_arr.shape)
```

(1, 128, 128, 3)

```
In [6]: prediction = model.predict(input_arr)
prediction,prediction.shape
```

1/1 [=====] - 15s 15s/step

```
Out[6]: (array([[8.7829467e-18, 2.8660604e-19, 4.7152817e-21, 6.3119323e-22,
3.2426425e-21, 2.2499923e-18, 9.6758780e-19, 1.9149978e-13,
1.0000000e+00, 2.2833669e-19, 1.2302450e-17, 8.4352367e-26,
4.7415217e-23, 8.9849995e-23, 1.2010913e-24, 1.8906961e-20,
2.2671302e-17, 4.1544132e-23, 3.9089671e-16, 2.5080584e-17,
1.2075649e-12, 2.9245322e-24, 1.2035763e-18, 2.6732390e-24,
7.2733559e-24, 3.4162246e-20, 2.8513275e-19, 3.0768677e-24,
7.7857995e-26, 7.8720473e-19, 2.9442566e-16, 1.0691177e-22,
2.3213051e-21, 4.6419847e-28, 7.0633841e-25, 2.0332089e-27,
2.3638059e-27, 5.2839713e-20]], dtype=float32),

(1, 38))
```

```
In [7]: result_index=np.argmax(prediction)
result_index
```

```
Out[7]: 8
```

```
In [8]: class_name = ['Apple___Apple_scab',
'Apple___Black_rot',
'Apple___Cedar_apple_rust',
'Apple___healthy',
'Blueberry___healthy',
'Cherry_(including_sour)___Powdery_mildew',
'Cherry_(including_sour)___healthy',
'Corn_(maize)___Cercospora_leaf_spot Gray_leaf_spot',
'Corn_(maize)___Common_rust_',
'Corn_(maize)___Northern_Leaf_Blight',
'Corn_(maize)___healthy',
'Grape___Black_rot',
'Grape___Esca_(Black_Measles)',
'Grape___Leaf_blight_(Isariopsis_Leaf_Spot)',
'Grape___healthy',
'Orange___Haunglongbing_(Citrus_greening)',
'Peach___Bacterial_spot',
'Peach___healthy',
'Pepper,_bell___Bacterial_spot',
'Pepper,_bell___healthy',
'Potato___Early_blight',
'Potato___Late_blight',
'Potato___healthy',
'Raspberry___healthy',
'Soybean___healthy',
'Squash___Powdery_mildew',
'Strawberry___Leaf_scorch',
'Strawberry___healthy',
'Tomato___Bacterial_spot',
'Tomato___Early_blight',
'Tomato___Late_blight',
'Tomato___Leaf_Mold',
'Tomato___Septoria_leaf_spot',
'Tomato___Spider_mites Two-spotted_spider_mite',
'Tomato___Target_Spot',
'Tomato___Tomato_Yellow_Leaf_Curl_Virus',
'Tomato___Tomato_mosaic_virus',
'Tomato___healthy']
```

```
In [9]: #Displaying Result Of diseases predaction
model_prediction = class_name[result_index]
plt.imshow(img)
plt.title(f"Disease Name: {model_prediction}")
plt.xticks([])
plt.yticks([])
plt.show()
```



```
In [10]: model_prediction
```

