

Fertilizers Recommendation System for Disease Prediction

1. INTRODUCTION

1.1 Overview

Productivity, food security, sustainability, and environmental impact are handled using smart farming techniques. As the population level is increasing day by day food production plays a major role in it[1]. The quality of the food products is essential for good healthy human beings. For providing this healthy environment analysis of various factor that causes a productive loss in agriculture has to be done. Identification of diseased part, Image classification, Detection of an anomaly in plant leaves open wide area research named image processing in agriculture [2].

Identification of the plant diseases is the key to preventing the losses in the yield and quantity of the agricultural product. There are two main factors which cause disease in plants and they are pathogens and environmental condition. Pathogens are the main reason for causing diseases in the plant.

Virus is a living organism which has living cells in it and it affects the plant. The region affected by the virus in plant leaf and fruits are seen as yellow streaking, yellow spots, deformed leaves, and stunted growth [3]. The best way to prevent viral disease is the disposal of the viral affected region.

Fungi is also one of the main reason for the productivity loss in the plant. Ascomycetes and basidiomycetes are two main fungi mainly responsible for disease in the plant. Fungicides are widely used for controlling fungal infection in a plant [4]. Magnaporthe grisea which is commonly known as rice blast disease. Sclerotinia sclerotiorum is responsible for cotton rot. Oomycetes and Phytophthora are the fungal-like organisms which contain destructive pathogen in the plant.

When a plant is infected by microscopic living organisms then it is a bacterial infection. Bacteria is a one-celled organism. The bacterially infected regions in plants can be seen as wilts, spots, and scabs. The spots which are caused by the blights can spread rapidly in plants. Tropical plants and vegetables are affected mainly by wilts. The absorption water by the plant is blocked by the bacterial infection. Some of the bacterial plant pathogens are Burkholderia and Proteobacteria.

Health monitoring and disease detection on plant is very critical for sustainable agriculture. Crop diseases identification remains difficult in many parts of the world due to the lack of the necessary infrastructure. It is very difficult to monitor the plant diseases manually. It requires tremendous amount of work, expertise in the plant diseases, and also require the excessive processing time. Hence, image processing and deep learning techniques are used for the detection of plant diseases. An automated system is introduced to identify different diseases on

plants by checking the symptoms shown on the leaves of the plant. Deep learning techniques are used to identify the diseases and suggest the precautions that can be taken for those diseases.

1.2 Purpose

Plant disease, especially on leaves, is one of the major factors that reduce the yield in both quality and quantity of the food crops. Finding the leaf disease is an important role to preserve agriculture. Smart analysis and Comprehensive prediction model in agriculture helps the farmer to yield right crop at the right time. The main benefits of the proposed system are as follows: Yield right crop at the right time, Balancing the crop production, control plant disease, Economic growth, and planning to reduce the crop scarcity. Hence to Detect and recognize the plant diseases and to recommend fertilizer it is necessary to provide symptoms in identifying the disease at its earliest.

2 LITERATURE SURVEY

2.1 Existing problem

Azadbakht et al. [5] used Gaussian process regression, random forest regression, support vector regression and boosted regression tree to analyze the performance for identifying leaf rust of wheat at canopy scale.

Elavarasan et al. [6] presented the models based on both supervised and unsupervised machine learning algorithm that were used for analyzing several factors like crop disease analysis, yield prediction and crop improvement.

Johannes et al. [7] predicted the symptoms Rust, septoria and tan disease in wheat plant using image processing algorithms. The images used for this analysis were taken from the mobile devices. Image-based automatic identification of plant disease mostly fails under real field condition.

Phadikar et al. [8] proposed an autonomous system for classifying disease in rice crop. It mainly extracts an infected region from the rice plant leaf. Information loss reduction and selection of an important feature are done through Roush Set Theory (RST).

Karadag et al. [9] proposed a technique that helps in identifying pepper fusarium disease. Light reflection from plant helps in knowing information about plant through specter radiometer. Four types of leaves were used for it namely mycorrhizal fungus, fusarium diseased, healthy and mycorrhizal leaves.

Liang et al. [10] proposed a technique based on the robust image-based plant disease diagnosis and severity estimation network (PD2SE-Net). It mainly focuses on practical diagnosis. This proposed technique also helps in measuring the severity of the disease.

Sandesh Raut and Karthik Ingale [11]proposed fast and accurate method for detection and classification of plant diseases. The proposed algorithm was tested on main five diseases on the

plant they are Early Scorch, Cottony mold, Ashen Mold, Late scorch, Tiny Whiteness. The feature extracted was recognized through a pre-trained neural network.

Sanjay et al [12] proposed detection and classification of plant disease using image processing and artificial neural networks. The detection and classification of plant diseases was divided into three steps such as identification of infected object extraction of feature set of the infected leaf images and detection and classification the type of disease using ANN.

2.2 Proposed solution

The works specified in the literature have limitations in terms of accuracy, computation and resource management. Hence an automated and new fertilizers Recommendation System for crop disease prediction is introduced. This system identifies different diseases on plants by checking the symptoms shown on the leaves of the plant. Deep learning techniques are used to identify the diseases and suggest the precautions that can be taken for those diseases. A web Application is built where

- Farmers interact with the portal build
- Interacts with the user interface to upload images of diseased leaf
- Our model built analyses the Disease and suggests the farmer with fertilizers are to be used

3 THEORITICAL ANALYSIS

3.1 Block diagram

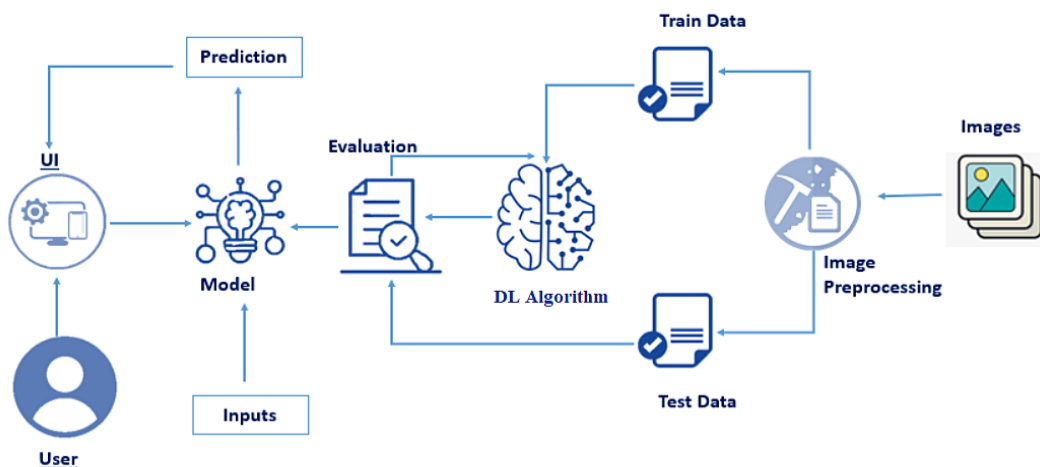


Figure 2: Architecture of the proposed system

The proposed approach is shown in Figure 2. The proposed approach consists of Data Collection,

Image Preprocessing, and Model Building for Fruit Disease Prediction, Model Building for Vegetable Disease Prediction, Testing of the Models, Application Building and Training the Model on IBM.

3.2 Hardware / Software designing

3.1 Hardware Requirements

- Processor : 2.5 gigahertz (GHz) frequency or above.
- RAM : A minimum of 4 GB of RAM.
- Hard disk : A minimum of 20 GB of available space.
- Input Device : High resolution camera
- Monitor : Minimum Resolution 1024 X 768.

3.2 Software Requirements

- Python, Python Web Frame Works,
- ANN, CNN, IBM Cloud,
- IBM Watson Studio, IBM Machine Learning,
- Python-Flask

4. EXPERIMENTAL INVESTIGATIONS

The first step in the implementation of the proposed approach is Data Collection. Dataset can be collected from the source given by the admins. Train and Test folders are created with each folder having subfolders with leaf images of different plant diseases. Two datasets are used, two models are created, one to detect vegetable leaf diseases like tomato, potato, and pepper plants and the second model is for fruits diseases like corn, peach, and apple.

Image Preprocessing:

Once the data are collected, this data is used to train the model. Before training the model the images are preprocessed and then feed on to the model for training. Keras ImageDataGenerator class for image preprocessing is used. The ImageDataGenerator accepts the original data, randomly transforms it, and returns only the new, transformed data. The next step is to build model for fruit leaf disease detection

Model Building for Fruit Disease Prediction:

The model building from the augmented images includes the following. Importing the model building Libraries, Initializing the model, Adding CNN Layers, Adding Hidden Layer, Adding Output Layer, Configuring the Learning Process, Training and testing the model and saving the model.

Three layers are added for CNN

- Convolution layer
- Pooling layer
- Flattening layer

Convolution Layer:

The first layer of the neural network model, the convolution layer is added. The convolution layer, takes a number of feature detectors, feature detector size, expected input shape of the image, and activation function as arguments. This layer applies feature detectors on the input image and returns a feature map (features from the image). 32 3 x 3 convolution filters are used in this work.

Activation Function:

These are the functions that help to decide if the node need to be activated or not. These functions introduce non-linearity in the networks. Relu activation in used in the initial layers and softmax in final layer.

Pooling layer:

Max Pooling selects the maximum element from the region of the feature map covered by the filter. Thus, the output after the max-pooling layer would be a feature map containing the most prominent features of the previous feature map.

Flatten layer:

The flatten layer is used to convert n-dimensional arrays to 1-dimensional arrays. This 1D array will be given as input to ANN fully connected layers.

Dense layers:

The name suggests that layers are fully connected (dense) by the neurons in a network layer. Each neuron in a layer receives input from all the neurons present in the previous layer. Dense is used to add the layers.

Compile the model:

After adding all the required layers, the model is compiled. For this step, loss function, optimizer and metrics for evaluation are passed as arguments.

Fit and save the model:

Fit the neural network model with the train and test set, number of epochs and validation steps. Steps per epoch is determined by number of training images/batch size, for validation steps number of validation images/batch size.

Accuracy, Loss:

Loss value implies how poorly or well a model behaves after each iteration of optimization. An accuracy metric is used to measure the algorithm's performance in an interpretable way. The accuracy of a model is usually determined after the model parameters and is calculated in the form of a percentage.

The weights are to be saved for future use. The weights are saved in as .h5.

Application Building:

After the model is built, the model is integrated into a web application so that normal users can also use it. The new users need to initially register in the portal. After registration users can log in to browse the images to detect the disease.

In this section, you have to build

- HTML pages - front end
- Python script - Server-side script

Python script is created using spyder IDE

6. RESULT

The screen shots of various steps of the automated system are shown below.

Figure 6.1-6.3 shows the training and testing of fruits disease classification using IBM Watson.

```
model.add(Dense(20, activation='relu'))
model.add(Dense(6, activation='softmax'))

In [14]: # Compile the model
model.compile(loss='categorical_crossentropy', optimizer='adam', metrics=['accuracy'])

In [*]: # Fit the model
model.fit_generator(x_train, steps_per_epoch=168, validation_data=x_test, validation_steps=52, epochs=3)
/tmp/ususer/ipykernel_164/876280831.py:2: UserWarning: 'Model.fit_generator' is deprecated and will be removed in a future version. Please use 'Model.fit', which supports generators.
model.fit_generator(x_train, steps_per_epoch=168, validation_data=x_test, validation_steps=52, epochs=3)
Epoch 1/3
123/168 [=====] - ETA: 48s - loss: 0.9772 - accuracy: 0.6938

In [*]: model.save('fruit.h5')

In [24]: ls
fruit-dataset/ fruit.h5

In [26]: !tar -zcvf FruitDiseaseClassification.tgz fruit.h5
fruit.h5

In [27]: ls
fruit-dataset/ FruitDiseaseClassification.tgz fruit.h5

Test

In [16]: import numpy as np
from tensorflow.keras.models import load_model
from tensorflow.keras.preprocessing import image

In [31]: # Load the model
model = load_model('fruit.h5')

In [32]: img = image.load_img(n"/home/ususer/work/fruit-dataset/test/Apple___Black_rot/2a4cb085-e831-4ec4-8acf-49cea51eeb5f___3R_Frg6.5_2900.39G", target_size=(128,128))
img
```

Figure 6.1: Fruit Leaf Disease model training

```
model.add(Dense(20, activation='relu'))
model.add(Dense(6, activation='softmax'))

In [14]: # Compile the model
model.compile(loss='categorical_crossentropy', optimizer='adam', metrics=['accuracy'])

In [*]: # Fit the model
model.fit_generator(x_train, steps_per_epoch=168, validation_data=x_test, validation_steps=52, epochs=3)
/tmp/ususer/ipykernel_164/876280831.py:2: UserWarning: 'Model.fit_generator' is deprecated and will be removed in a future version. Please use 'Model.fit', which supports generators.
model.fit_generator(x_train, steps_per_epoch=168, validation_data=x_test, validation_steps=52, epochs=3)
Epoch 1/3
168/168 [=====] - 203s 1s/step - loss: 0.8118 - accuracy: 0.7438 - val_loss: 56.1117 - val_accuracy: 0.8257
Epoch 2/3
123/168 [=====] - ETA: 46s - loss: 0.2901 - accuracy: 0.8994

In [*]: model.save('fruit.h5')

In [24]: ls
fruit-dataset/ fruit.h5

In [26]: !tar -zcvf FruitDiseaseClassification.tgz fruit.h5
fruit.h5

In [27]: ls
fruit-dataset/ FruitDiseaseClassification.tgz fruit.h5

Test

In [16]: import numpy as np
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In [31]: # Load the model
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img
```

Figure 6.2 : Training process of the model

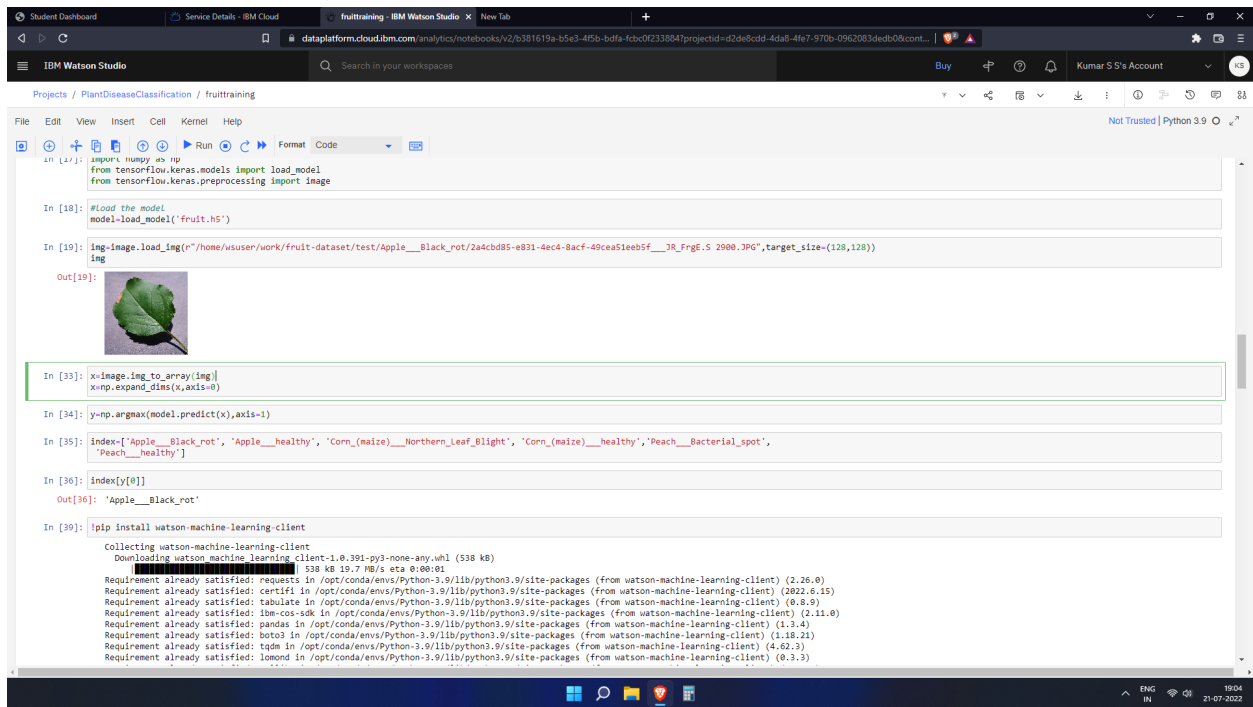


Figure 6.3 Testing of the trained model with an image from the test dataset

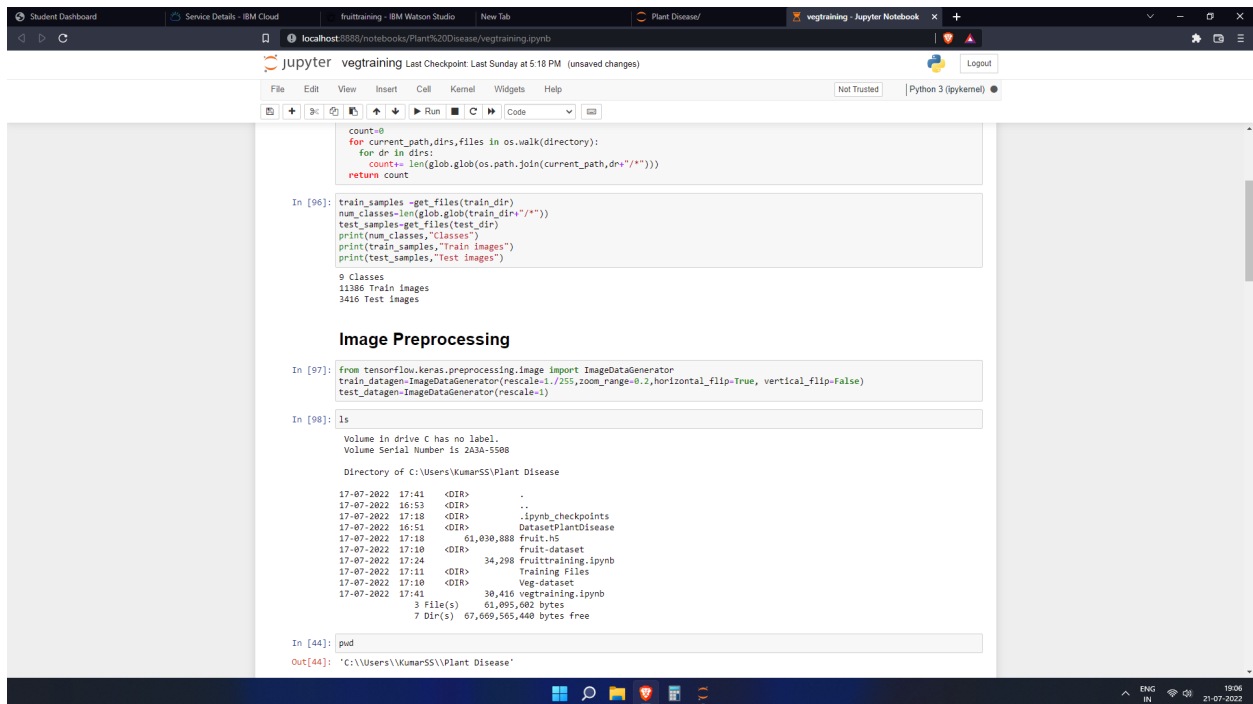


Figure 6.4 Implementation of vegetable disease classification model using Jupyter

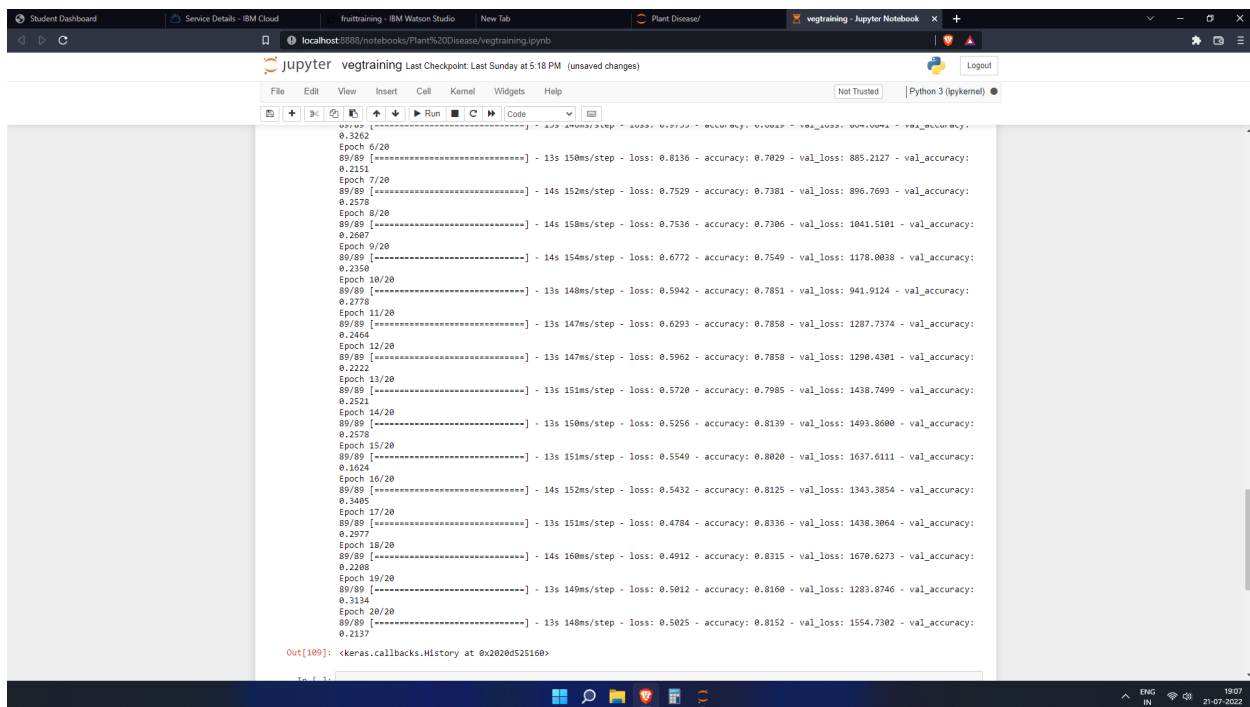


Figure 6.5 Training of vegetable disease classification model using Jupyter

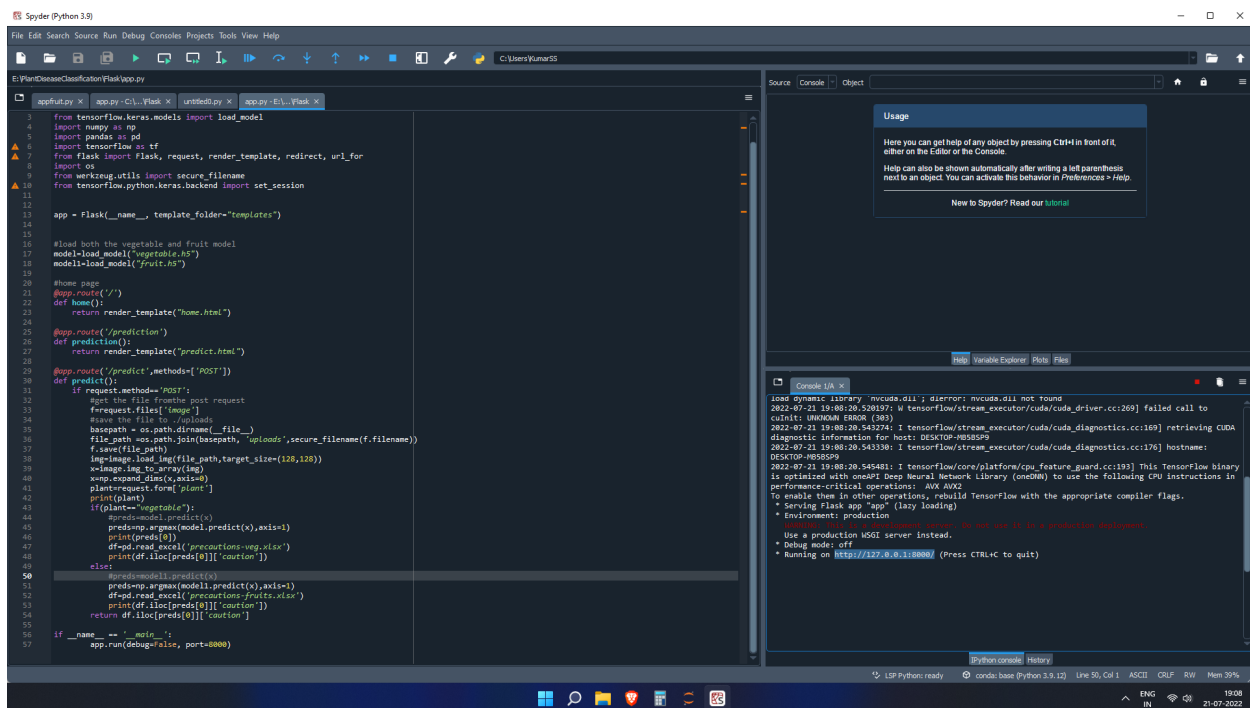


Figure 6.6 Creating web application using Spyder IDE

The trained and saved model fruits.h5 and vegetables.h5 was used for creating a web application using Flask. The process is shown in Figure 6.6.

Figure 6.7: Demo page of the web app

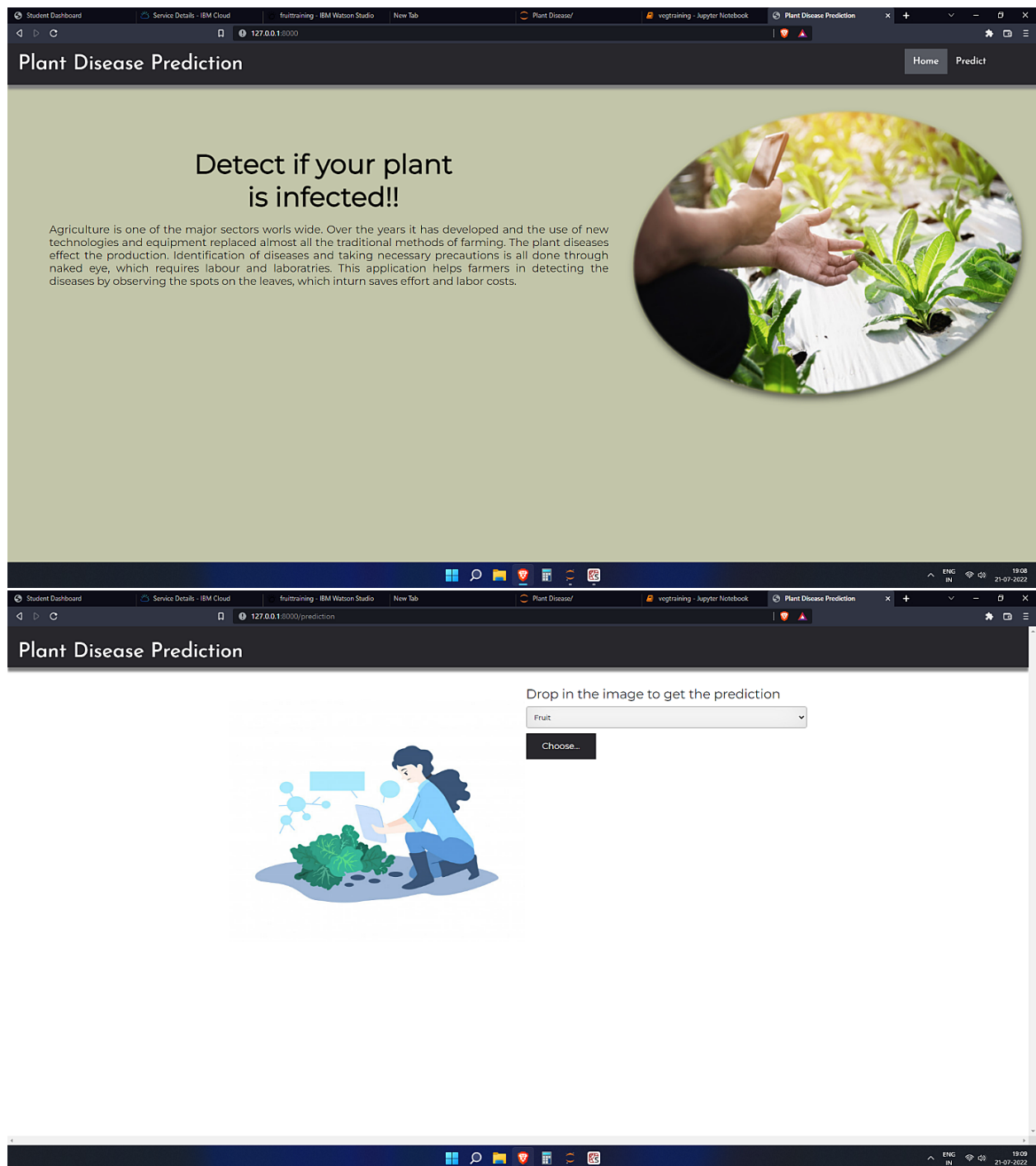


Figure 6.8: Web app page asking to choose one type from vegetable and fruit

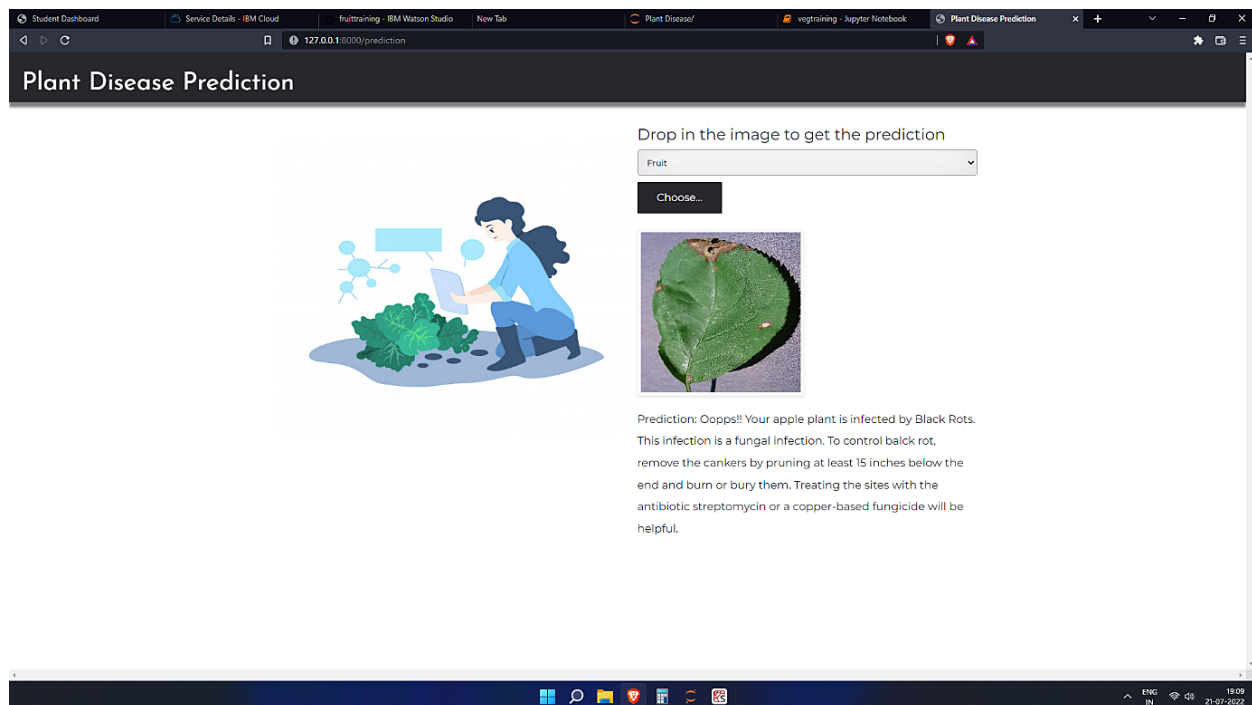


Figure 6.9: Web app page predicting the disease and recommending the suitable fertilizer for the test data.

7 ADVANTAGES & DISADVANTAGES

The proposed system can help farmers to identify crop diseases at an early stage. The system also suggests the combination of fertilizers to be used by the farmers, based on the analysis done by the system. This can promote crop production and thereby increase the productivity. This can also protect the quality of soil, so that future cultivations can yield good productivity.

The limitation of the system is, there are cases where the system couldn't identify the diseases correctly. Also the types of diseases considered are minimal. If the user interface allows to connect to the camera directly, it would be helpful to all the farmers.

8 APPLICATIONS

The proposed system can be applied in different areas,

1. Automated Disease Prediction
2. Automatic fertilizer recommendation
3. Disease prediction by mobile phones.
4. Smart Agriculture Solutions

9 CONCLUSION

Monitoring the health and identifying the disease on plants is essential for sustainable agriculture. An automated system using image processing and deep learning techniques was introduced to identify different diseases on plants by checking the symptoms shown on the leaves of the plant and suggest the precautions that can be taken for those diseases. Two datasets i) vegetable leaf diseases like tomato, potato, and pepper plants and the ii) fruits diseases like corn, peach, and apple was studied in this work. The system was trained and modelled using deep learning CNN. The trained models were used to create a web app through which farmers can easily identify the disease of the plant and get suggestions for the use of Fertilizers.

10 FUTURE SCOPE

The system can be further improved by using some recent networks. Transfer learning techniques can be applied to improve the accuracy. The system can also be extended to more number of disease prediction.

11 BIBILOGRAPHY

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APPENDIX

A. Source Code

Fruit disease training Model

```
import os, types
import pandas as pd
from botocore.client import Config
import ibm_boto3

def __iter__(self): return 0
```

```

# @hidden_cell
# The following code accesses a file in your IBM Cloud Object Storage. It
includes your credentials.
# You might want to remove those credentials before you share the notebook.
client_5081f81d79d14bc98585e4995f36ea4f = ibm_boto3.client(service_name='s3',
    ibm_api_key_id='ayQ-QLPt1QO3ZO_iHtnBUv8N2Lh6qRL08LZ4_Fi-p3KG',
    ibm_auth_endpoint="https://iam.cloud.ibm.com/oidc/token",
    config=Config(signature_version='oauth'),
    endpoint_url='https://s3.private.us.cloud-object-storage.appdomain.cloud')

streaming_body_2 =
client_5081f81d79d14bc98585e4995f36ea4f.get_object(Bucket='plantdiseaseclassifi
cation-donotdelete-pr-3jizmpkeywlkkv', Key='fruit-dataset.zip')['Body']

# Your data file was loaded into a botocore.response.StreamingBody object.
# Please read the documentation of ibm_boto3 and pandas to learn more about
the possibilities to load the data.
# ibm_boto3 documentation: https://ibm.github.io/ibm-cos-sdk-python/
# pandas documentation: http://pandas.pydata.org/

```

In [2]:

```

from io import BytesIO
import zipfile
unzip=zipfile.ZipFile(BytesIO(streaming_body_2.read()), 'r')
file_paths=unzip.namelist()
for path in file_paths:
    unzip.extract(path)

```

In [3]:

```

from tensorflow.keras.preprocessing.image import ImageDataGenerator

```

In [7]:

```

train_datagen=ImageDataGenerator(rescale=1./255, zoom_range=0.2, horizontal_flip=
True, vertical_flip=False)
test_datagen=ImageDataGenerator(rescale=1)

```

In [4]:

```

ls
fruit-dataset/

```

In [5]:

```

pwd

```

Out[5]:

```

'/home/wsuser/work'

```

In [8]:

```
x_train=train_datagen.flow_from_directory(r"/home/wsuser/work/fruit-  
dataset/train",target_size=(128,128),class_mode='categorical',batch_size=32)  
x_test=test_datagen.flow_from_directory(r"/home/wsuser/work/fruit-  
dataset/test",target_size=(128,128),class_mode='categorical',batch_size=32)  
Found 5384 images belonging to 6 classes.  
Found 1686 images belonging to 6 classes.
```

In [9]:

```
x_train.class_indices
```

Out[9]:

```
{'Apple___Black_rot': 0,  
 'Apple___healthy': 1,  
 'Corn_(maize)___Northern_Leaf_Blight': 2,  
 'Corn_(maize)___healthy': 3,  
 'Peach___Bacterial_spot': 4,  
 'Peach___healthy': 5}
```

In [10]:

```
#Import keras models and layers
```

```
from keras.models import Sequential  
from keras.layers import Dense  
from keras.layers import Convolution2D  
from keras.layers import MaxPooling2D  
from keras.layers import Flatten
```

In [11]:

```
model=Sequential()  
model.add(Convolution2D(32,(3,3), input_shape=(128,128,3),activation='relu'))  
model.add(MaxPooling2D(pool_size=(2,2)))  
model.add(Flatten())
```

In [12]:

```
model.summary()  
Model: "sequential"
```

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 126, 126, 32)	896
max_pooling2d (MaxPooling2D)	(None, 63, 63, 32)	0
flatten (Flatten)	(None, 127008)	0

```
Total params: 896
Trainable params: 896
Non-trainable params: 0
```

In [13]:

```
model.add(Dense(40, activation='relu'))
model.add(Dense(20, activation='relu'))
model.add(Dense(6, activation='softmax'))
```

In [14]:

```
# Compile the model
model.compile(loss='categorical_crossentropy', optimizer='adam',
metrics=['accuracy'])
```

In [15]:

```
#Fit the model
model.fit_generator(x_train, steps_per_epoch=168, validation_data=x_test,
validation_steps=52, epochs=3)
/tmp/wsuser/ipykernel_164/876289031.py:2: UserWarning: `Model.fit_generator` is
deprecated and will be removed in a future version. Please use `Model.fit`,
which supports generators.
    model.fit_generator(x_train, steps_per_epoch=168, validation_data=x_test,
validation_steps=52, epochs=3)
Epoch 1/3
168/168 [=====] - 203s 1s/step - loss: 0.8118 -
accuracy: 0.7438 - val_loss: 56.1117 - val_accuracy: 0.8257
Epoch 2/3
168/168 [=====] - 201s 1s/step - loss: 0.2739 -
accuracy: 0.9049 - val_loss: 344.3285 - val_accuracy: 0.5150
Epoch 3/3
168/168 [=====] - 202s 1s/step - loss: 0.2159 -
accuracy: 0.9277 - val_loss: 390.2771 - val_accuracy: 0.5571
```

Out[15]:

```
<keras.callbacks.History at 0x7f8e92092700>
```

In [16]:

```
model.save('fruit.h5')
```

In [24]:

```
ls
fruit-dataset/  fruit.h5
```

In [26]:

```
!tar -zcvf FruitDiseaseClassification.tgz fruit.h5
fruit.h5
```

In [27]:


```
ls
fruit-dataset/  FruitDiseaseClassification.tgz  fruit.h5
```

Test

In [17]:

```
import numpy as np
from tensorflow.keras.models import load_model
from tensorflow.keras.preprocessing import image
```

In [18]:

```
#load the model
model=load_model('fruit.h5')
```

In [19]:

```
img=image.load_img(r"/home/wsuser/work/fruit-
dataset/test/Apple___Black_rot/2a4cbd85-e831-4ec4-8acf-
49cea51eeb5f___JR_FrgE.S 2900.JPG",target_size=(128,128))
img
```

Out[19]:

In [33]:

```
x=image.img_to_array(img)
x=np.expand_dims(x,axis=0)
```

In [34]:

```
y=np.argmax(model.predict(x),axis=1)
```

In [35]:

```
index=['Apple___Black_rot', 'Apple___healthy',
'Corn_(maize)___Northern_Leaf_Blight',
'Corn_(maize)___healthy', 'Peach___Bacterial_spot',
'Peach___healthy']
```

In [36]:

```
index[y[0]]
```

Out[36]:

```
'Apple___Black_rot'
```

In [39]:

```
!pip install watson-machine-learning-client
Collecting watson-machine-learning-client
  Downloading watson_machine_learning_client-1.0.391-py3-none-any.whl (538 kB)
    |████████████████████████████████████████| 538 kB 19.7 MB/s eta 0:00:01
Requirement already satisfied: requests in /opt/conda/envs/Python-
```

3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (2.26.0)
Requirement already satisfied: certifi in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from watson-machine-learning-client)
(2022.6.15)
Requirement already satisfied: tabulate in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (0.8.9)
Requirement already satisfied: ibm-cos-sdk in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (2.11.0)
Requirement already satisfied: pandas in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (1.3.4)
Requirement already satisfied: boto3 in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from watson-machine-learning-client)
(1.18.21)
Requirement already satisfied: tqdm in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (4.62.3)
Requirement already satisfied: lomond in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (0.3.3)
Requirement already satisfied: urllib3 in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (1.26.7)
Requirement already satisfied: s3transfer<0.6.0,>=0.5.0 in
/opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from boto3->watson-
machine-learning-client) (0.5.0)
Requirement already satisfied: jmespath<1.0.0,>=0.7.1 in
/opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from boto3->watson-
machine-learning-client) (0.10.0)
Requirement already satisfied: botocore<1.22.0,>=1.21.21 in
/opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from boto3->watson-
machine-learning-client) (1.21.41)
Requirement already satisfied: python-dateutil<3.0.0,>=2.1 in
/opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from
botocore<1.22.0,>=1.21.21->boto3->watson-machine-learning-client) (2.8.2)
Requirement already satisfied: six>=1.5 in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from python-dateutil<3.0.0,>=2.1-
>botocore<1.22.0,>=1.21.21->boto3->watson-machine-learning-client) (1.15.0)
Requirement already satisfied: ibm-cos-sdk-s3transfer==2.11.0 in
/opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from ibm-cos-sdk-
>watson-machine-learning-client) (2.11.0)
Requirement already satisfied: ibm-cos-sdk-core==2.11.0 in
/opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from ibm-cos-sdk-
>watson-machine-learning-client) (2.11.0)
Requirement already satisfied: charset-normalizer~=2.0.0 in
/opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from requests->watson-
machine-learning-client) (2.0.4)
Requirement already satisfied: idna<4,>=2.5 in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from requests->watson-machine-learning-

```
client) (3.3)
Requirement already satisfied: pytz>=2017.3 in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from pandas->watson-machine-learning-client)
(2021.3)
Requirement already satisfied: numpy>=1.17.3 in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from pandas->watson-machine-learning-client)
(1.20.3)
Installing collected packages: watson-machine-learning-client
Successfully installed watson-machine-learning-client-1.0.391
```

In [37]:

```
from ibm_watson_machine_learning import APIClient
wml_credentials={
    "url":"https://us-south.ml.cloud.ibm.com",
    "apikey":"79usIO-TltcICWjvzF9sxfShUpnN9UWqIn0B960NTJlE"
}
```

In [38]:

```
client=APIClient(wml_credentials)
```

In [39]:

```
client
```

Out[39]:

```
<ibm_watson_machine_learning.client.APIClient at 0x7fbfb58dc430>
```

In [40]:

```
def guid_space_name(client, PlantDiseaseDeploy):
    space=client.spaces.get_details()
    return (next(item for item in space ['resources'] if item
    ['entity']['name']==PlantDiseaseDeploy)['metadata']['id'])
```

In [41]:

```
space_uid=guid_space_name(client, 'PlantDiseaseDeploy')
print(space_uid)
0a27f7f4-c72b-418f-a896-a0f57f6b1154
```

In [42]:

```
client.set.default_space(space_uid)
```

Out[42]:

```
'SUCCESS'
```

In [44]:

```
pip install tensorflow
Requirement already satisfied: tensorflow in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (2.7.2)
Requirement already satisfied: protobuf>=3.9.2 in /opt/conda/envs/Python-
3.9/lib/python3.9/site-packages (from tensorflow) (3.19.1)
```

Requirement already satisfied: gast<0.5.0,>=0.2.1 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (0.4.0)

Requirement already satisfied: termcolor>=1.1.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (1.1.0)

Requirement already satisfied: astunparse>=1.6.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (1.6.3)

Requirement already satisfied: typing-extensions>=3.6.6 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (3.7.4.3)

Requirement already satisfied: h5py>=2.9.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (3.2.1)

Requirement already satisfied: grpcio<2.0,>=1.24.3 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (1.42.0)

Requirement already satisfied: keras<2.8,>=2.7.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (2.7.0)

Requirement already satisfied: flatbuffers<3.0,>=1.12 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (2.0)

Requirement already satisfied: tensorboard~=2.7 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (2.7.0)

Requirement already satisfied: tensorflow-estimator<2.8,~=2.7.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (2.7.0)

Requirement already satisfied: absl-py>=0.4.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (0.12.0)

Requirement already satisfied: opt-einsum>=2.3.2 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (3.3.0)

Requirement already satisfied: keras-preprocessing>=1.1.1 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (1.1.2)

Requirement already satisfied: wheel<1.0,>=0.32.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (0.37.0)

Requirement already satisfied: google-pasta>=0.1.1 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (0.2.0)

Requirement already satisfied: six>=1.12.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (1.15.0)

Requirement already satisfied: wrapt>=1.11.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (1.12.1)

Requirement already satisfied: tensorflow-io-gcs-filesystem>=0.21.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (0.23.1)

Requirement already satisfied: numpy>=1.14.5 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow) (1.20.3)

Requirement already satisfied: google-auth-oauthlib<0.5,>=0.4.1 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~=2.7->tensorflow) (0.4.4)

Requirement already satisfied: google-auth<3,>=1.6.3 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~=2.7->tensorflow) (1.23.0)

Requirement already satisfied: markdown>=2.6.8 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~=2.7->tensorflow) (3.3.3)

Requirement already satisfied: tensorboard-plugin-wit>=1.6.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~=2.7->tensorflow) (1.6.0)

Requirement already satisfied: werkzeug>=0.11.15 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~=2.7->tensorflow) (2.0.2)

Requirement already satisfied: setuptools>=41.0.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~=2.7->tensorflow) (58.0.4)

Requirement already satisfied: tensorboard-data-server<0.7.0,>=0.6.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~=2.7->tensorflow) (0.6.1)

Requirement already satisfied: requests<3,>=2.21.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~=2.7->tensorflow) (2.26.0)

Requirement already satisfied: rsa<5,>=3.1.4 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from google-auth<3,>=1.6.3->tensorboard~=2.7->tensorflow) (4.7.2)

Requirement already satisfied: cachetools<5.0,>=2.0.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from google-auth<3,>=1.6.3->tensorboard~=2.7->tensorflow) (4.2.2)

Requirement already satisfied: pyasn1-modules>=0.2.1 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from google-auth<3,>=1.6.3->tensorboard~=2.7->tensorflow) (0.2.8)

Requirement already satisfied: requests-oauthlib>=0.7.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from google-auth-oauthlib<0.5,>=0.4.1->tensorboard~=2.7->tensorflow) (1.3.0)

Requirement already satisfied: pyasn1<0.5.0,>=0.4.6 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from pyasn1-modules>=0.2.1->google-auth<3,>=1.6.3->tensorboard~=2.7->tensorflow) (0.4.8)

Requirement already satisfied: idna<4,>=2.5 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from requests<3,>=2.21.0->tensorboard~=2.7->tensorflow) (3.3)

Requirement already satisfied: urllib3<1.27,>=1.21.1 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from requests<3,>=2.21.0->tensorboard~=2.7->tensorflow) (1.26.7)

Requirement already satisfied: certifi>=2017.4.17 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from requests<3,>=2.21.0->tensorboard~=2.7->tensorflow) (2022.6.15)

Requirement already satisfied: charset-normalizer~=2.0.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from requests<3,>=2.21.0->tensorboard~=2.7->tensorflow) (2.0.4)

Requirement already satisfied: oauthlib>=3.0.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from requests-oauthlib>=0.7.0->google-auth-

oauthlib<0.5,>=0.4.1->tensorboard~=2.7->tensorflow) (3.2.0)

Note: you may need to restart the kernel to use updated packages.

In [48]:

```
client.software_specifications.list(100)
```

NAME	ASSET_ID	TYPE
default_py3.6	0062b8c9-8b7d-44a0-a9b9-46c416adcbd9	base
kernel-spark3.2-scala2.12	020d69ce-7ac1-5e68-ac1a-31189867356a	base
pytorch-onnx_1.3-py3.7-edt	069ea134-3346-5748-b513-49120e15d288	base
scikit-learn_0.20-py3.6	09c5a1d0-9c1e-4473-a344-eb7b665ff687	base
spark-mllib_3.0-scala_2.12	09f4cff0-90a7-5899-b9ed-1ef348aebdee	base
pytorch-onnx_rt22.1-py3.9	0b848dd4-e681-5599-be41-b5f6fccc6471	base
ai-function_0.1-py3.6	0cdb0f1e-5376-4f4d-92dd-da3b69aa9bda	base
shiny-r3.6	0e6e79df-875e-4f24-8ae9-62dcc2148306	base
tensorflow_2.4-py3.7-horovod	1092590a-307d-563d-9b62-4eb7d64b3f22	base
pytorch_1.1-py3.6	10ac12d6-6b30-4ccd-8392-3e922c096a92	base
tensorflow_1.15-py3.6-ddl	111e41b3-de2d-5422-a4d6-bf776828c4b7	base
runtime-22.1-py3.9	12b83a17-24d8-5082-900f-0ab31fbfd3cb	base
scikit-learn_0.22-py3.6	154010fa-5b3b-4ac1-82af-4d5ee5abbc85	base
default_r3.6	1b70aec3-ab34-4b87-8aa0-a4a3c8296a36	base
pytorch-onnx_1.3-py3.6	1bc6029a-cc97-56da-b8e0-39c3880dbbe7	base
pytorch-onnx_rt22.1-py3.9-edt	1d362186-7ad5-5b59-8b6c-9d0880bde37f	base
tensorflow_2.1-py3.6	1eb25b84-d6ed-5dde-b6a5-3fbdf1665666	base
spark-mllib_3.2	20047f72-0a98-58c7-9ff5-a77b012eb8f5	base
tensorflow_2.4-py3.8-horovod	217c16f6-178f-56bf-824a-b19f20564c49	base
runtime-22.1-py3.9-cuda	26215f05-08c3-5a41-a1b0-da66306ce658	base
do_py3.8	295addb5-9ef9-547e-9bf4-92ae3563e720	base
autoai-ts_3.8-py3.8	2aa0c932-798f-5ae9-abd6-15e0c2402fb5	base
tensorflow_1.15-py3.6	2b73a275-7cbf-420b-a912-eae7f436e0bc	base
pytorch_1.2-py3.6	2c8ef57d-2687-4b7d-acce-01f94976dac1	base
spark-mllib_2.3	2e51f700-bca0-4b0d-88dc-5c6791338875	base
pytorch-onnx_1.1-py3.6-edt	32983cea-3f32-4400-8965-dde874a8d67e	base
spark-mllib_3.0-py37	36507ebe-8770-55ba-ab2a-eafe787600e9	base
spark-mllib_2.4	390d21f8-e58b-4fac-9c55-d7ceda621326	base
xgboost_0.82-py3.6	39e31acd-5f30-41dc-ae44-60233c80306e	base
pytorch-onnx_1.2-py3.6-edt	40589d0e-7019-4e28-8daa-fb03b6f4fe12	base
default_r36py38	41c247d3-45f8-5a71-b065-8580229facf0	base
autoai-ts_rt22.1-py3.9	4269d26e-07ba-5d40-8f66-2d495b0c71f7	base
autoai-obm_3.0	42b92e18-d9ab-567f-988a-4240baled5f7	base
pmml-3.0_4.3	493bcb95-16f1-5bc5-bee8-81b8af80e9c7	base
spark-mllib_2.4-r_3.6	49403dff-92e9-4c87-a3d7-a42d0021c095	base
xgboost_0.90-py3.6	4ff8d6c2-1343-4c18-85e1-689c965304d3	base
pytorch-onnx_1.1-py3.6	50f95b2a-bc16-43bb-bc94-b0bed208c60b	base
autoai-ts_3.9-py3.8	52c57136-80fa-572e-8728-a5e7cbb42cde	base

spark-mllib_2.4-scala_2.11	55a70f99-7320-4be5-9fb9-9edb5a443af5	base
spark-mllib_3.0	5c1b0ca2-4977-5c2e-9439-ffd44ea8ffe9	base
autoai-obm_2.0	5c2e37fa-80b8-5e77-840f-d912469614ee	base
spss-modeler_18.1	5c3cad7e-507f-4b2a-a9a3-ab53a21dee8b	base
cuda-py3.8	5d3232bf-c86b-5df4-a2cd-7bb870a1cd4e	base
autoai-kb_3.1-py3.7	632d4b22-10aa-5180-88f0-f52dfb6444d7	base
pytorch-onnx_1.7-py3.8	634d3cdc-b562-5bf9-a2d4-ea90a478456b	base
spark-mllib_2.3-r_3.6	6586b9e3-ccd6-4f92-900f-0f8cb2bd6f0c	base
tensorflow_2.4-py3.7	65e171d7-72d1-55d9-8ebb-f813d620c9bb	base
spss-modeler_18.2	687eddc9-028a-4117-b9dd-e57b36f1efa5	base
pytorch-onnx_1.2-py3.6	692a6a4d-2c4d-45ff-a1ed-b167ee55469a	base
spark-mllib_2.3-scala_2.11	7963efe5-bbec-417e-92cf-0574e21b4e8d	base
spark-mllib_2.4-py37	7abc992b-b685-532b-a122-a396a3cdbaab	base
caffe_1.0-py3.6	7bb3dbe2-da6e-4145-918d-b6d84aa93b6b	base
pytorch-onnx_1.7-py3.7	812c6631-42b7-5613-982b-02098e6c909c	base
cuda-py3.6	82c79ece-4d12-40e6-8787-a7b9e0f62770	base
tensorflow_1.15-py3.6-horovod	8964680e-d5e4-5bb8-919b-8342c6c0dfd8	base
hybrid_0.1	8c1a58c6-62b5-4dc4-987a-df751c2756b6	base
pytorch-onnx_1.3-py3.7	8d5d8a87-a912-54cf-81ec-3914adaa988d	base
caffe-ibm_1.0-py3.6	8d863266-7927-4d1e-97d7-56a7f4c0a19b	base
spss-modeler_17.1	902d0051-84bd-4af6-ab6b-8f6aa6fdeabb	base
do_12.10	9100fd72-8159-4eb9-8a0b-a87e12eefa36	base
do_py3.7	9447fa8b-2051-4d24-9eef-5acb0e3c59f8	base
spark-mllib_3.0-r_3.6	94bb6052-c837-589d-83f1-f4142f219e32	base
cuda-py3.7-opence	94e9652b-7f2d-59d5-ba5a-23a414ea488f	base
nlp-py3.8	96e60351-99d4-5a1c-9cc0-473ac1b5a864	base
cuda-py3.7	9a44990c-1aa1-4c7d-baf8-c4099011741c	base
hybrid_0.2	9b3f9040-9cee-4ead-8d7a-780600f542f7	base
spark-mllib_3.0-py38	9f7a8fc1-4d3c-5e65-ab90-41fa8de2d418	base
autoai-kb_3.3-py3.7	a545cca3-02df-5c61-9e88-998b09dc79af	base
spark-mllib_3.0-py39	a6082a27-5acc-5163-b02c-6b96916eb5e0	base
runtime-22.1-py3.9-do	a7e7dbf1-1d03-5544-994d-e5ec845ce99a	base
default_py3.8	ab9e1b80-f2ce-592c-a7d2-4f2344f77194	base
tensorflow_rt22.1-py3.9	acd9c798-6974-5d2f-a657-ce06e986df4d	base
kernel-spark3.2-py3.9	ad7033ee-794e-58cf-812e-a95f4b64b207	base
autoai-obm_2.0 with Spark 3.0	af10f35f-69fa-5d66-9bf5-acb58434263a	base
default_py3.7_opence	c2057dd4-f42c-5f77-a02f-72bdbd3282c9	base
tensorflow_2.1-py3.7	c4032338-2a40-500a-beef-b01ab2667e27	base
do_py3.7_opence	cc8f8976-b74a-551a-bb66-6377f8d865b4	base
autoai-kb_3.0-py3.6	d139f196-e04b-5d8b-9140-9a10ca1fa91a	base
spark-mllib_3.0-py36	d82546d5-dd78-5fbb-9131-2ec309bc56ed	base
autoai-kb_3.4-py3.8	da9b39c3-758c-5a4f-9cfd-457dd4d8c395	base
kernel-spark3.2-r3.6	db2fe4d6-d641-5d05-9972-73c654c60e0a	base
autoai-kb_rt22.1-py3.9	db6afe93-665f-5910-b117-d879897404d9	base
tensorflow_rt22.1-py3.9-horovod	dda170cc-ca67-5da7-9b7a-cf84c6987fae	base

autoai-ts_1.0-py3.7	deef04f0-0c42-5147-9711-89f9904299db	base
tensorflow_2.1-py3.7-horovod	e384fce5-fdd1-53f8-bc71-11326c9c635f	base
default_py3.7	e4429883-c883-42b6-87a8-f419d64088cd	base
do_22.1	e51999ba-6452-5f1f-8287-17228b88b652	base
autoai-obm_3.2	eae86aab-da30-5229-a6a6-1d0d4e368983	base
do_20.1	f686cdd9-7904-5f9d-a732-01b0d6b10dc5	base
scikit-learn_0.19-py3.6	f963fa9d-4bb7-5652-9c5d-8d9289ef6ad9	base
tensorflow_2.4-py3.8	fe185c44-9a99-5425-986b-59bd1d2eda46	base

In [49]:

```
software_space_uid=client.software_specifications.get_uid_by_name('tensorflow_rt
22.1-py3.9')
software_space_uid
```

Out[49]:

```
'acd9c798-6974-5d2f-a657-ce06e986df4d'
```

In [52]:

```
model_details =
client.repository.store_model(model='FruitDiseaseClassification.tgz',
    meta_props={
        client.repository.ModelMetaNames.NAME: 'PlantDiseaseDeploy',
        client.repository.ModelMetaNames.TYPE: 'tensorflow_2.7',

client.repository.ModelMetaNames.SOFTWARE_SPEC_UID: software_space_uid } )
```

In [53]:

```
model_id=client.repository.get_model_id(model_details)
```

In [54]:

```
model_id
```

Out[54]:

```
'13fa8062-da8c-4f6b-b6bb-d3f1e385af17'
```

In [57]:

```
client.repository.download(model_id, 'FruitDisease.tar.gz')
Successfully saved model content to file: 'FruitDisease.tar.gz'
```

Out[57]:

```
'/home/wsuser/work/FruitDisease.tar.gz'
```

In [58]:

```
ls
fruit-dataset/      FruitDisease.tag.gb  fruit.h5
FruitDiseaseClassification.tgz  FruitDisease.tar.gz
```

In []:

Vegetable disease training model

```
# Import Libraries
import os
import glob
import matplotlib.pyplot as plt
import numpy as np

# Keras API
import keras
from keras.models import Sequential
from keras.layers import Dense, Dropout, Flatten
from keras.layers import Conv2D, MaxPooling2D, Activation, AveragePooling2D, BatchNormalization
from keras.preprocessing.image import ImageDataGenerator
```

In [94]:

```
train_dir = "Veg-dataset/train_set/"
test_dir = "Veg-dataset/test_set/"
```

In [95]:

```
# function to get count of images
def get_files(directory):
    if not os.path.exists(directory):
        return 0
    count=0
    for current_path, dirs, files in os.walk(directory):
        for dr in dirs:
            count+= len(glob.glob(os.path.join(current_path, dr+"/*")))
    return count
```

In [96]:

```
train_samples = get_files(train_dir)
num_classes = len(glob.glob(train_dir+"/*"))
test_samples = get_files(test_dir)
print(num_classes, "Classes")
print(train_samples, "Train images")
print(test_samples, "Test images")
9 Classes
11386 Train images
3416 Test images
```

Image Preprocessing

In [97]:

```

from tensorflow.keras.preprocessing.image import ImageDataGenerator
train_datagen=ImageDataGenerator(rescale=1./255, zoom_range=0.2, horizontal_flip=
True, vertical_flip=False)
test_datagen=ImageDataGenerator(rescale=1)

```

In [98]:

```

ls
Volume in drive C has no label.
Volume Serial Number is 2A3A-5508

Directory of C:\Users\KumarSS\Plant Disease

```

```

17-07-2022  17:41    <DIR>          .
17-07-2022  16:53    <DIR>          ..
17-07-2022  17:18    <DIR>          .ipynb_checkpoints
17-07-2022  16:51    <DIR>          DatasetPlantDisease
17-07-2022  17:18             61,030,888 fruit.h5
17-07-2022  17:10    <DIR>          fruit-dataset
17-07-2022  17:24             34,298 fruittraining.ipynb
17-07-2022  17:11    <DIR>          Training Files
17-07-2022  17:10    <DIR>          Veg-dataset
17-07-2022  17:41             30,416 vegtraining.ipynb
               3 File(s)          61,095,602 bytes
               7 Dir(s)  67,669,565,440 bytes free

```

In [44]:

```
pwd
```

Out[44]:

```
'C:\\Users\\KumarSS\\Plant Disease'
```

In [99]:

```

x_train=train_datagen.flow_from_directory('Veg-
dataset/train_set',target_size=(128,128),class_mode='categorical',batch_size=1
6)
Found 11386 images belonging to 9 classes.

```

In [100]:

```

x_test=test_datagen.flow_from_directory('Veg-
dataset/test_set',target_size=(128,128),class_mode='categorical',batch_size=2
6)
Found 3416 images belonging to 9 classes.

```

In [101]:

```
x_train.class_indices
```

Out[101]:

```
{'Pepper,_bell___Bacterial_spot': 0,
```

```
'Pepper___bell___healthy': 1,
'Potato___Early_blight': 2,
'Potato___Late_blight': 3,
'Potato___healthy': 4,
'Tomato___Bacterial_spot': 5,
'Tomato___Late_blight': 6,
'Tomato___Leaf_Mold': 7,
'Tomato___Septoria_leaf_spot': 8}
```

CNN Implementation

In [102]:

```
#Import keras models and layers
```

```
from keras.models import Sequential
from keras.layers import Dense
from keras.layers import Convolution2D
from keras.layers import MaxPooling2D
from keras.layers import Flatten
```

In [103]:

```
model=Sequential()
model.add(Convolution2D(32, (3, 3), input_shape=(128, 128, 3), activation='relu'))
model.add(MaxPooling2D(pool_size=(2, 2)))
model.add(Flatten())
```

In [104]:

```
model.summary()
Model: "sequential_4"
```

Layer (type)	Output Shape	Param #
conv2d_4 (Conv2D)	(None, 126, 126, 32)	896
max_pooling2d_4 (MaxPooling 2D)	(None, 63, 63, 32)	0
flatten_4 (Flatten)	(None, 127008)	0
Total params: 896		
Trainable params: 896		
Non-trainable params: 0		

Hidden layers

In [105]:

```
model.add(Dense(300, activation='relu'))
model.add(Dense(150, activation='relu'))
model.add(Dense(75, activation='relu'))
model.add(Dense(9, activation='softmax'))
```

In [106]:

```
# Compile the model
model.compile(loss='categorical_crossentropy', optimizer='adam',
metrics=['accuracy'])
```

In [107]:

```
len(x_train)
```

Out[107]:

```
712
```

In [108]:

```
len(x_test)
```

Out[108]:

```
132
```

In [109]:

```
#Fit the model
model.fit_generator(x_train, steps_per_epoch=89, validation_data=x_test,
validation_steps=27, epochs=20)
#model.fit(x_train, steps_per_epoch=len(x_train), validation_data=x_test,
validation_steps=len(x_test), epochs=3)
Epoch 1/20
C:\Users\KumarSS\AppData\Local\Temp\ipykernel_11188\4058285867.py:2:
UserWarning: `Model.fit_generator` is deprecated and will be removed in a
future version. Please use `Model.fit`, which supports generators.
    model.fit_generator(x_train, steps_per_epoch=89, validation_data=x_test,
validation_steps=27, epochs=20)
89/89 [=====] - 14s 154ms/step - loss: 2.4896 -
accuracy: 0.3111 - val_loss: 212.3326 - val_accuracy: 0.4188
Epoch 2/20
89/89 [=====] - 13s 148ms/step - loss: 1.4210 -
accuracy: 0.5070 - val_loss: 255.8210 - val_accuracy: 0.4117
Epoch 3/20
89/89 [=====] - 13s 147ms/step - loss: 1.1745 -
accuracy: 0.5997 - val_loss: 351.9586 - val_accuracy: 0.4103
Epoch 4/20
```

89/89 [=====] - 13s 145ms/step - loss: 0.9475 -
accuracy: 0.6685 - val_loss: 797.9529 - val_accuracy: 0.2664
Epoch 5/20
89/89 [=====] - 13s 146ms/step - loss: 0.9755 -
accuracy: 0.6819 - val_loss: 864.6841 - val_accuracy: 0.3262
Epoch 6/20
89/89 [=====] - 13s 150ms/step - loss: 0.8136 -
accuracy: 0.7029 - val_loss: 885.2127 - val_accuracy: 0.2151
Epoch 7/20
89/89 [=====] - 14s 152ms/step - loss: 0.7529 -
accuracy: 0.7381 - val_loss: 896.7693 - val_accuracy: 0.2578
Epoch 8/20
89/89 [=====] - 14s 158ms/step - loss: 0.7536 -
accuracy: 0.7306 - val_loss: 1041.5101 - val_accuracy: 0.2607
Epoch 9/20
89/89 [=====] - 14s 154ms/step - loss: 0.6772 -
accuracy: 0.7549 - val_loss: 1178.0038 - val_accuracy: 0.2350
Epoch 10/20
89/89 [=====] - 13s 148ms/step - loss: 0.5942 -
accuracy: 0.7851 - val_loss: 941.9124 - val_accuracy: 0.2778
Epoch 11/20
89/89 [=====] - 13s 147ms/step - loss: 0.6293 -
accuracy: 0.7858 - val_loss: 1287.7374 - val_accuracy: 0.2464
Epoch 12/20
89/89 [=====] - 13s 147ms/step - loss: 0.5962 -
accuracy: 0.7858 - val_loss: 1290.4301 - val_accuracy: 0.2222
Epoch 13/20
89/89 [=====] - 13s 151ms/step - loss: 0.5720 -
accuracy: 0.7985 - val_loss: 1438.7499 - val_accuracy: 0.2521
Epoch 14/20
89/89 [=====] - 13s 150ms/step - loss: 0.5256 -
accuracy: 0.8139 - val_loss: 1493.8600 - val_accuracy: 0.2578
Epoch 15/20
89/89 [=====] - 13s 151ms/step - loss: 0.5549 -
accuracy: 0.8020 - val_loss: 1637.6111 - val_accuracy: 0.1624
Epoch 16/20
89/89 [=====] - 14s 152ms/step - loss: 0.5432 -
accuracy: 0.8125 - val_loss: 1343.3854 - val_accuracy: 0.3405
Epoch 17/20
89/89 [=====] - 13s 151ms/step - loss: 0.4784 -
accuracy: 0.8336 - val_loss: 1438.3064 - val_accuracy: 0.2977
Epoch 18/20
89/89 [=====] - 14s 160ms/step - loss: 0.4912 -
accuracy: 0.8315 - val_loss: 1670.6273 - val_accuracy: 0.2208
Epoch 19/20

```
89/89 [=====] - 13s 149ms/step - loss: 0.5012 -  
accuracy: 0.8160 - val_loss: 1283.8746 - val_accuracy: 0.3134  
Epoch 20/20  
89/89 [=====] - 13s 148ms/step - loss: 0.5025 -  
accuracy: 0.8152 - val_loss: 1554.7302 - val_accuracy: 0.2137
```

Out[109]:

```
<keras.callbacks.History at 0x2020d525160>
```

In []:

Save the model

In [110]:

```
model.save('vegetable.h5')
```

In [111]:

```
ls  
Volume in drive C has no label.  
Volume Serial Number is 2A3A-5508  
  
Directory of C:\Users\KumarSS\Plant Disease  
  
17-07-2022  17:47    <DIR>          .  
17-07-2022  16:53    <DIR>          ..  
17-07-2022  17:18    <DIR>          .ipynb_checkpoints  
17-07-2022  16:51    <DIR>          DatasetPlantDisease  
17-07-2022  17:18             61,030,888 fruit.h5  
17-07-2022  17:10    <DIR>          fruit-dataset  
17-07-2022  17:24             34,298 fruittraining.ipynb  
17-07-2022  17:11    <DIR>          Training Files  
17-07-2022  17:10    <DIR>          Veg-dataset  
17-07-2022  17:47             457,981,136 vegetable.h5  
17-07-2022  17:47             15,570 vegtraining.ipynb  
          4 File(s)      519,061,892 bytes  
          7 Dir(s)   67,206,922,240 bytes free
```

In []:

Testing the model

Loading [MathJax]/jax/output/CommonHTML/fonts/TeX/fontdata.js

Import Models

In [25]:

```
import numpy as np
import keras
from tensorflow.keras.models import load_model
from tensorflow.keras.preprocessing.image import img_to_array
from keras.preprocessing import image
import tensorflow as tf
Load the saved model (fruit)
```

In [40]:

```
model=load_model('fruit.h5')
```

In [41]:

```
img=keras.utils.load_img("fruit-dataset/test/Apple___Black_rot/2a4cbd85-e831-4ec4-8acf-49cea51eeb5f___JR_FrgE.S 2900.JPG",target_size=(128,128))
img
```

Out[41]:

In [42]:

```
x=tf.keras.utils.img_to_array(img)
x=np.expand_dims(x,axis=0)
y=np.argmax(model.predict(x),axis=1)
1/1 [=====] - 0s 47ms/step
```

In [43]:

```
index=['Apple___Black_rot', 'Apple___healthy',
'Corn_(maize)___Northern_Leaf_Blight',
'Corn_(maize)___healthy', 'Peach___Bacterial_spot',
'Peach___healthy']
```

In [44]:

```
index[y[0]]
```

Out[44]:

```
'Apple___Black_rot'
# Load the saved model (vegetable)
```

In [45]:

```
model=load_model('vegetable.h5')
```

In [51]:

```
img=keras.utils.load_img("Veg-
dataset/test_set/Tomato___Bacterial_spot/b5c376b6-3c0a-46d7-a053-
```

```
2b3a57168897___UF.GRC_BS_Lab_Leaf_9003.JPG",target_size=(128,128))
img
```

Out[51]:

In [52]:

```
x=tf.keras.utils.img_to_array(img)
x=np.expand_dims(x,axis=0)
y=np.argmax(model.predict(x),axis=1)
1/1 [=====] - 0s 26ms/step
```

In [53]:

```
index=['Pepper,_bell___Bacterial_spot','Pepper,_bell___healthy','Potato___Earl
y_blight','Potato___Late_blight','Potato___healthy','Tomato___Bacterial_spot',
'Tomato___Late_blight','Tomato___Leaf_Mold','Tomato___Septoria_leaf_spot']
```

In [54]:

```
index[y[0]]
```

Out[54]:

```
'Tomato___Septoria_leaf_spot'
```

In []:

Flask app creation

```
import requests
from tensorflow.keras.preprocessing import image
from tensorflow.keras.models import load_model
import numpy as np
import pandas as pd
import tensorflow as tf
from flask import Flask, request, render_template, redirect, url_for
import os
from werkzeug.utils import secure_filename
from tensorflow.python.keras.backend import set_session
```

```
app = Flask(__name__, template_folder="templates")
```

```
#load both the vegetable and fruit model
model=load_model("vegetable.h5")
modell=load_model("fruit.h5")
```



```

#home page
@app.route('/')
def home():
    return render_template("home.html")

@app.route('/prediction')
def prediction():
    return render_template("predict.html")

@app.route('/predict', methods=['POST'])
def predict():
    if request.method=='POST':
        #get the file from the post request
        f=request.files['image']
        #save the file to ./uploads
        basepath = os.path.dirname(__file__)
        file_path =os.path.join(basepath, 'uploads', secure_filename(f.filename))
        f.save(file_path)
        img=image.load_img(file_path,target_size=(128,128))
        x=image.img_to_array(img)
        x=np.expand_dims(x,axis=0)
        plant=request.form['plant']
        print(plant)
        if(plant=="vegetable"):
            #preds=model.predict(x)
            preds=np.argmax(model.predict(x),axis=1)
            print(preds[0])
            df=pd.read_excel('precautions-veg.xlsx')
            print(df.iloc[preds[0]]['caution'])
        else:
            #preds=model1.predict(x)
            preds=np.argmax(model1.predict(x),axis=1)
            df=pd.read_excel('precautions-fruits.xlsx')
            print(df.iloc[preds[0]]['caution'])
        return df.iloc[preds[0]]['caution']

if __name__ == '__main__':
    app.run(debug=False, port=8000)

```