

Fertilizers Recommendation System for Disease Prediction

PROJECT REPORT

Submitted by

Team ID: PNT2022TMID0341

AMILENENI DHANUSH

HEMANTH N

KANCHARLA LIKITH CHOWDARY

VENKATA ASI SUPREETH

In partial fulfilment for the award of the degree

Of

BACHELOR OF TECHNOLOGY

In

INFORMATION ENGINEERING



**SAVEETHA ENGINEERING COLLEGE ,
THANDALAM.**

1.INTRODUCTION

1.1 Overview Two datasets—the fruit dataset and the vegetable dataset—are gathered for this research. Convolutional Neural Networks, a deep learning neural network, is used to train and test the datasets that have been collected (CNN). The fruit dataset is first trained, and then CNN is tested. There are 6 courses total, and each class is trained and tested. The vegetable dataset is then tested and trained. Python is the programme used to train and test datasets. All of the Python code is initially created in the Jupyter notebook that comes with Anaconda Python, and it is then tested in the IBM cloud. Finally, Flask, a Python package, is used to construct a web-based framework. Along with their related files in the static folder, two html files are created and placed in the templates folder. The Spyder-Anaconda Python software "app.py" that interfaces with these two webpages was developed and tested.

1.2 Purpose This study is used to test samples of fruits and vegetables and find out which diseases they may have. Additionally, this project suggests fertilisers for certain ailments.

2.LITERATURE SURVEY

2.1 Existing issue Indumathi suggested a technique for spotting leaf illnesses and suggested fertilisers to treat them. However, the method's low number of train and test sets leads to subpar accuracy. A straightforward approach of prediction for a soil-based fertiliser prescription system for anticipated crop diseases was put out by Pandi Selvi. This approach offers less predictability and accuracy. Shiva Reddy proposed a machine learning-based IoT-based system for recommending fertiliser and detecting leaf disease that has less than 80% accuracy.

2.2 Proposed remedy A deep learning-based neural network is utilised in this project's effort to train and test the datasets that were gathered. CNN, a deep learning-based neural network, provides classification accuracy rates of greater than 90%. By boosting The accuracy rate can rise to 95% to 98% by adding more dense layers and changing hyperparameters like the number of epochs and batch size.

3.THEORITICAL ANALYSIS

3.1 Block diagram

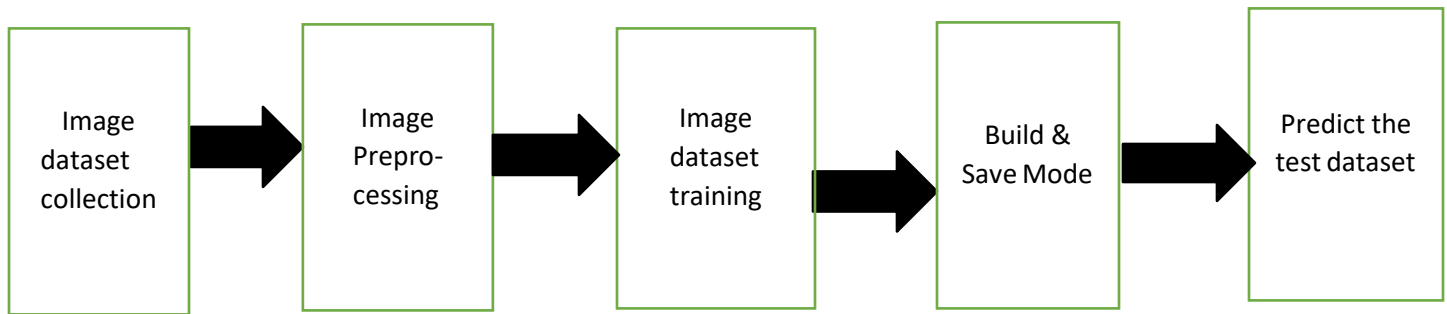


Figure.3.1. Block Diagram of the project

Fig.3.1 displays the project's overall block diagram. The collecting of picture datasets comes first, then comes image preparation. The training of picture datasets using various hyperparameter initializations is the third phase. After that, create the model and save it in.h5 format. The final step involves applying the trained model to test new or existing datasets.

3.2 Hardware/Software designing

Python is the programme used to train and test the dataset. Python programming is done in a notebook tool called Jupyter, which also works with the IBM cloud. Convolutional Neural Network is the neural network that was utilised to train and test the model (CNN).

The CNN has following layers:

- Convolutional layer (32x32 kernal (3x3))

- Max-pool layer (kernel(2x2))
- Flatten layer
- Dense layer (different layers with different size)
- Drop out layer (optional)
- Final output dense layer(size 6x1 for fruit dataset and 9x1 for Vegetable dataset)

Images are normalised to 1 and then downsized to 128x128 in the preparation stage. Different batch sizes are used to arrange the photos. Then, using the gathered datasets, train set and test set are created. The following Python libraries need to be imported before beginning the process in order to perform the aforementioned actions in Python:

- NumPy
- TensorFlow
- Keras
- Matplotlib (optional for data visualization)

The following activation functions used in the CNN training:

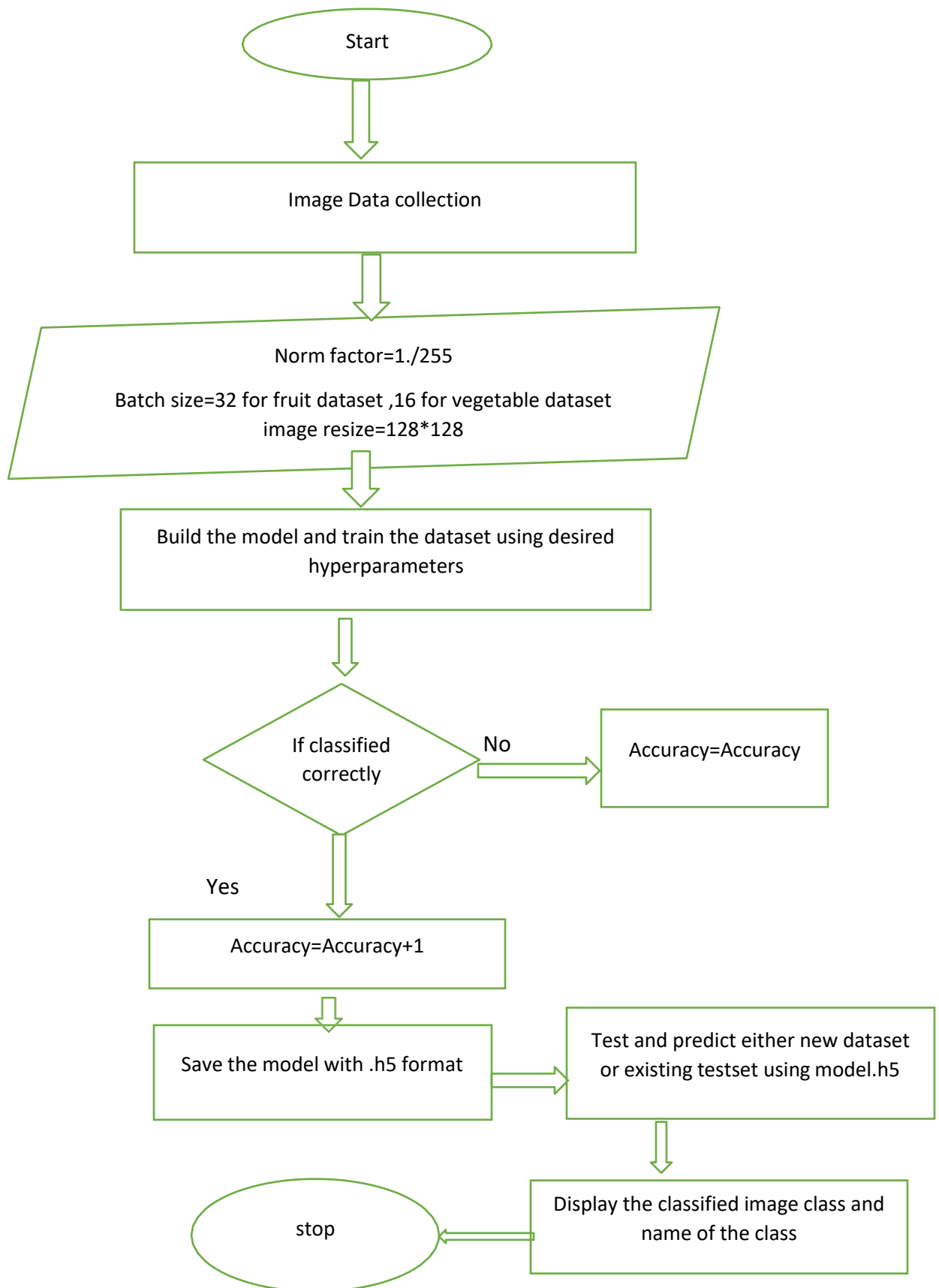
- Max Pool layer and RELU at the conclusion of the convolution layer
- End of output dense layer SoftMax
- The dataset argmax is utilised for testing, and it's optional

4.EXPERIMENTAL INVESTIGATIONS

Analysis performed when developing the solution There are various and tested batch sizes. The accuracy provided by CNN varies with batch size. Size of the batch

specifies how many iterations there are in each epoch. The quantity of epochs is a key hyper parameter. When compared to other hyper metrics, this affects accuracy and has a significant impact on accuracy. By increasing the number of epochs, the accuracy may be adjusted from 80% to 90% for the vegetable dataset and from 95% to 98% for the fruit dataset. The size of the training and test datasets also has a significant impact on accuracy. More photos can be added to the train dataset to boost accuracy. The size of the train dataset and the number of epochs both increase the computational time required to create a model. The train dataset and test dataset batch sizes are also very important in terms of processing time. When there are additional convolutional layers, the complexity of the neural network increases. A higher layer count will produce results with greater accuracy. The CNN algorithm requires more time to develop a model and spend more time training as the number of layers increases. The size of the train datasets affects the model.h5 size. However, the amount of the train dataset and the complexity of the CNN architecture determine the memory need.

5.FLOWCHARTS



6.RESULTS

The following is a screenshot of the project's final results: Fruit dataset testing and training



```
In [40]: pred = model.predict_classes(x)

WARNING:tensorflow:From c:\python-input-40-f93bd030d5d\1: Sequential.predict_classes (from tensorflow.python.keras.engine.sequ
ential) is deprecated and will be removed after 2021-01-01.
Instructions for updating:
Please use `model.predict(x, axis=-1)`, if your model does multi-class classification (e.g. if it uses
a 'softmax' last-layer activation), or `(model.predict(x) > 0.5).astype('int32')`, if your model does binary classification
(e.g. if it uses a 'sigmoid' last-layer activation).

In [41]: pred
Out[41]: array([1], dtype=int64)

In [42]: index = ['Apple__Black_rot', 'Apple__healthy', 'Corn_(maize)__Northern_Leaf_Blight', 'Corn_(maize)__healthy', 'Peach__Bacterial_
spot', 'Peach__healthy']

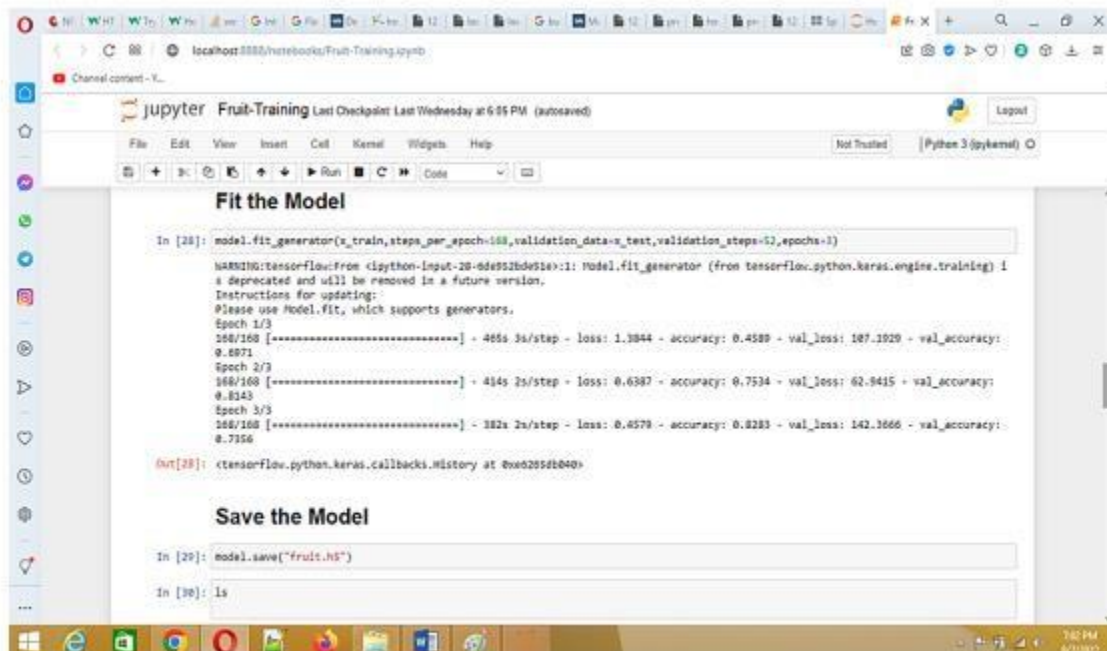
In [43]: print('the given image belongs to-', index[pred[0]])
the given image belongs to- Apple__healthy

Test Apple Black Rot class images

In [54]: img = image.load_img('E:\IBM_COURSE\Project\Dataset Plant Disease\fruit-dataset\fruit-dataset\test\Apple__Black_rot\0f3d46f6
a.jpg')

In [55]: x_test = image.img_to_array(img)
```

Figure.6.2 Test the Fruit dataset



```
Fit the Model

In [28]: model.fit_generator(x_train, steps_per_epoch=100, validation_data=x_test, validation_steps=10, epochs=1)

WARNING:tensorflow:From c:\python-input-28-d4d952bde51a\1: Model.fit_generator (from tensorflow.python.keras.engine.training) i
s deprecated and will be removed in a future version.
Instructions for updating:
Please use Model.fit, which supports generators.

Epoch 1/3
168/168 [=====] - 486s 3s/step - loss: 1.3844 - accuracy: 0.4589 - val_loss: 187.1929 - val_accuracy:
0.8971
Epoch 2/3
168/168 [=====] - 414s 2s/step - loss: 0.6387 - accuracy: 0.7534 - val_loss: 62.9415 - val_accuracy:
0.8543
Epoch 3/3
168/168 [=====] - 382s 2s/step - loss: 0.4579 - accuracy: 0.8283 - val_loss: 142.3666 - val_accuracy:
0.7356

Out[28]: <tensorflow.python.keras.callbacks.History at 0x0205d040>

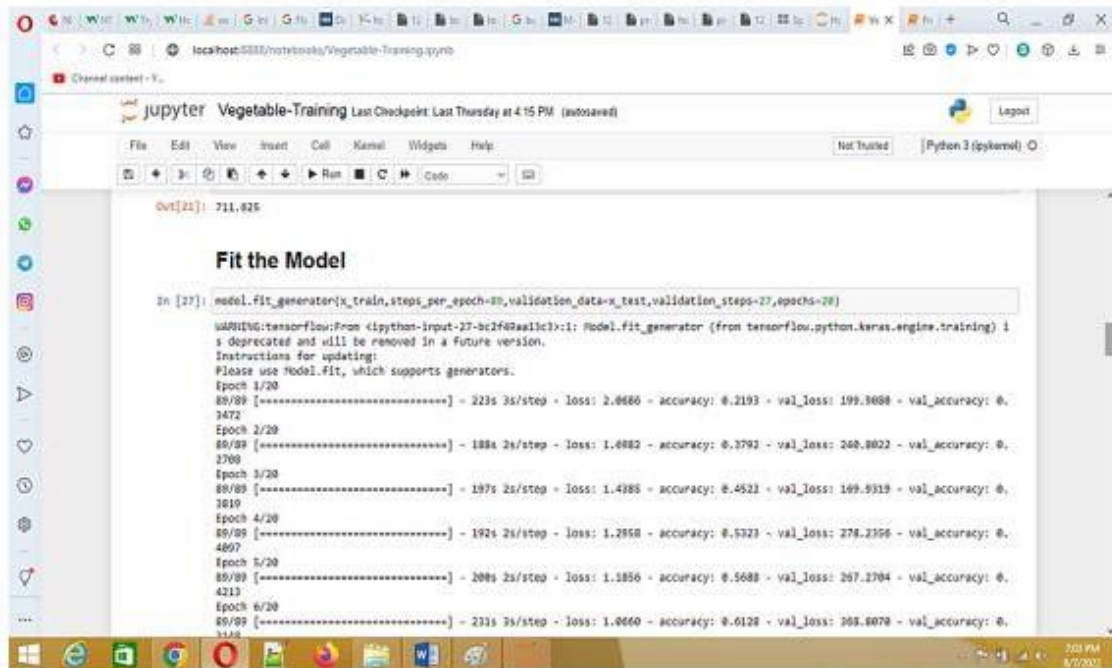
Save the Model

In [29]: model.save("fruit.h5")

In [30]: Is
```

Figure.6.1. Fit a model for Fruit dataset

Train and Test Vegetable dataset



The screenshot shows a Jupyter Notebook titled "Vegetable-Training" with a Python 3 (ipykernel) environment. The notebook is running a code cell that trains a model. The output shows the model's performance over 20 epochs. The training loss decreases from 2.0666 to 1.0660, and the validation accuracy increases from 0.2193 to 0.6128.

```
Out[21]: 711.825
```

Fit the Model

```
In [27]: model.fit_generator(x_train, steps_per_epoch=80, validation_data=x_test, validation_steps=27, epochs=20)
```

WARNING:tensorflow:From <ipython-input-27-bc2f42aa1c1>:1: Model.fit_generator (from tensorflow.python.keras.engine.training) is deprecated and will be removed in a future version. Instructions for updating: Please use Model.fit, which supports generators.

Epoch 1/20
80/80 [=====] - 223s 3s/step - loss: 2.0666 - accuracy: 0.2193 - val_loss: 199.8668 - val_accuracy: 0.3472

Epoch 2/20
80/80 [=====] - 188s 2s/step - loss: 1.6882 - accuracy: 0.3792 - val_loss: 240.8022 - val_accuracy: 0.2708

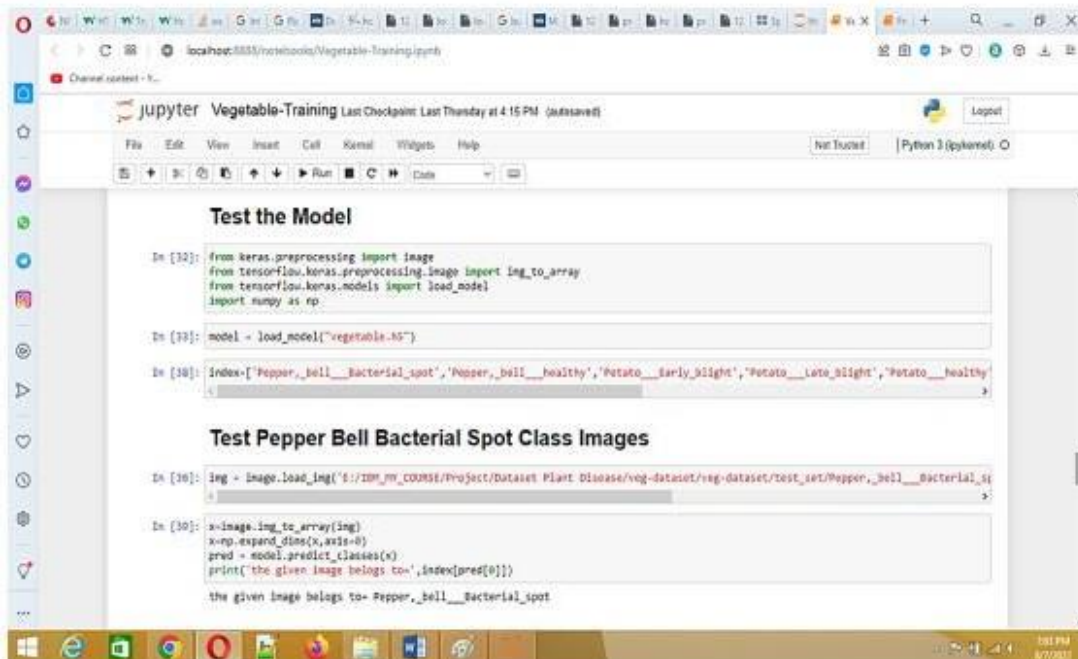
Epoch 3/20
80/80 [=====] - 197s 2s/step - loss: 1.4085 - accuracy: 0.4521 - val_loss: 169.9319 - val_accuracy: 0.3610

Epoch 4/20
80/80 [=====] - 192s 2s/step - loss: 1.2858 - accuracy: 0.5329 - val_loss: 278.2356 - val_accuracy: 0.4807

Epoch 5/20
80/80 [=====] - 208s 2s/step - loss: 1.1856 - accuracy: 0.5688 - val_loss: 267.2784 - val_accuracy: 0.4213

Epoch 6/20
80/80 [=====] - 231s 3s/step - loss: 1.0660 - accuracy: 0.6128 - val_loss: 265.6078 - val_accuracy: 0.3318

Figure.6.3. Train the Vegetable dataset



The screenshot shows a Jupyter Notebook titled "Vegetable-Training" with a Python 3 (ipykernel) environment. The notebook is running a code cell that tests the model. The output shows the model's performance on a test set of images. The model correctly identifies the class of the images.

Test the Model

```
In [32]: from keras.preprocessing import image  
from tensorflow.keras.preprocessing.image import img_to_array  
from tensorflow.keras.models import load_model  
import numpy as np
```

```
In [33]: model = load_model("vegetable.h5")
```

```
In [38]: index=['Pepper_bell__Bacterial_spot', 'Pepper_bell__healthy', 'Potato__Early_blight', 'Potato__Late_blight', 'Potato__healthy']
```

Test Pepper Bell Bacterial Spot Class Images

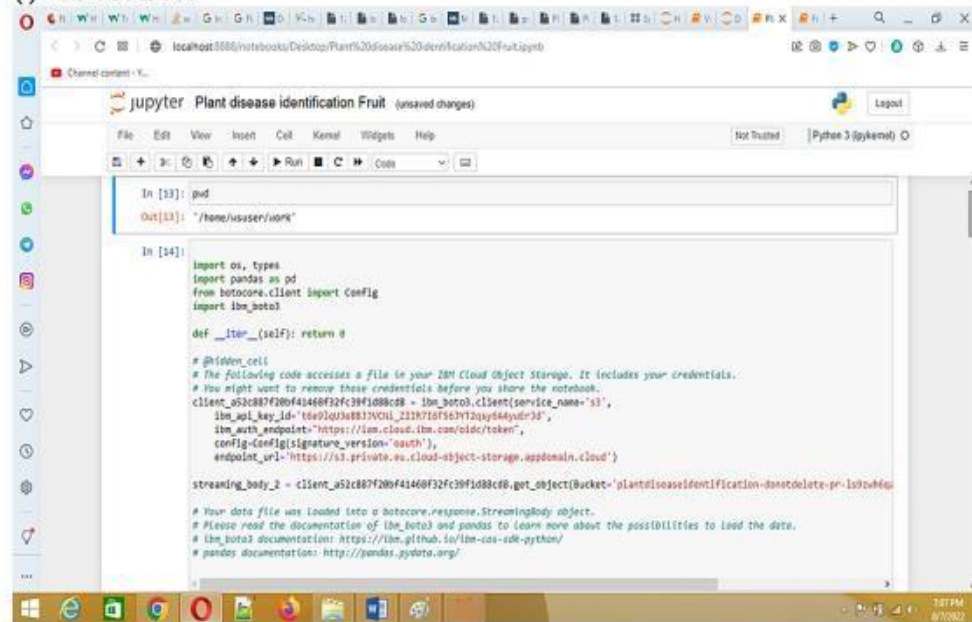
```
In [36]: img = image.load_img('F:/DITM_MY_COURSE/Project/Dataset Plant Disease/veg-dataset/veg-dataset/test_set/Pepper_bell__Bacterial_spot')  
x = image.img_to_array(img)  
x = np.expand_dims(x, axis=0)  
pred = model.predict_classes(x)  
print('the given image belongs to-', index[pred[0]])  
the given image belongs to- Pepper_bell__Bacterial_spot
```

Figure.6.4. Test the Vegetable dataset

Train and Test Vegetable dataset IBM Cloud

Due to CUH limit exceeds, I have downloaded the notebooks and opened in Jupyter notebook

(i). Fruit dataset



```
In [13]: pd
Out[13]: "/home/user/work"

In [14]:
import os, types
import pandas as pd
from botocore.client import Config
import ibm_botocore

def __iter__(self): return 0

# @iiden_cell
# The following code accesses a file in your IBM Cloud Object Storage. It includes your credentials.
# You might want to remove these credentials before you share the notebook.
client = boto3.client('s3',
    aws_access_key_id='t6a9303e8832vchi_233k7167563712ay644udr3d',
    aws_secret_access_key='t6a9303e8832vchi_233k7167563712ay644udr3d',
    config=Config(signature_version='s3v4'),
    endpoint_url='https://s3.private.eu.cloud-object-storage.appdomain.cloud')

streaming_body_2 = client.get_object(Bucket='plantdiseaseidentification-donotdelete-pr-1s9zwh6u',
    Key='fruit_dataset.csv')

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

Figure.6.5. Training Fruit Dataset in IBM Cloud

```
In [6]: pid
Out[6]: '/home/vsuser/work'

In [7]:
import os, types
import pandas as pd
from botocore.client import Config
import boto3

def __iter__(self): return 0

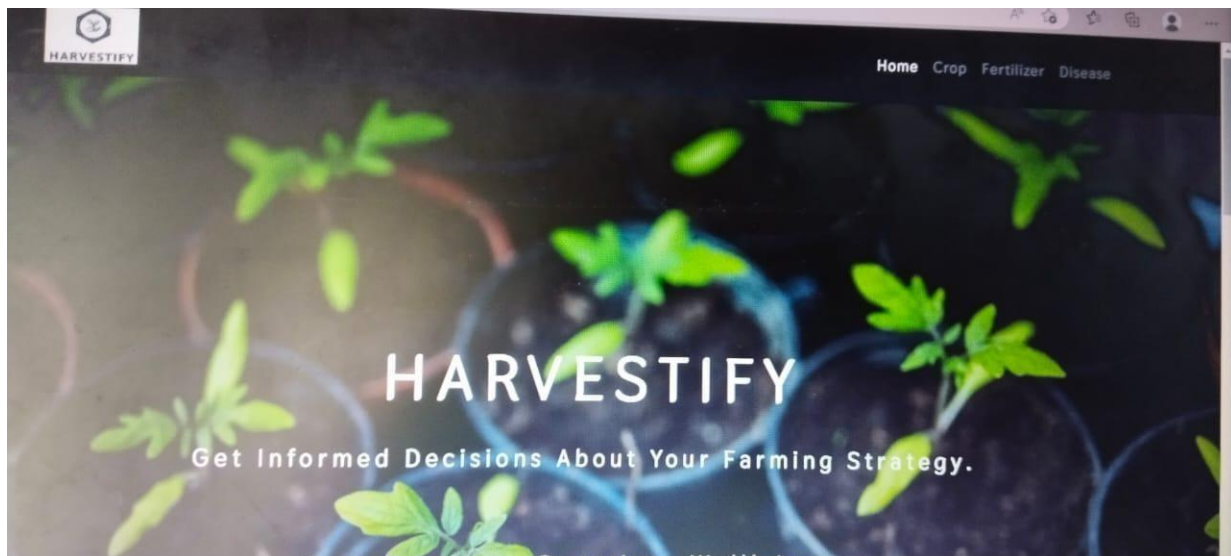
# @hidden_cell
# The following code accesses a file in your IBM Cloud Object Storage. It includes your credentials.
# You might want to remove these credentials before you share the notebook.
client = boto3.client(service_name='s3',
    iam_api_key_id='idw4idsh837XUJ_12IDIG056V72pydddydRd',
    iam_auth_endpoint='https://iam.cloud.ibm.com/oidc/token',
    config=Config(signature_version='auth'),
    endpoint_url='https://s3.private.eu.cloud-object-storage.appdomain.cloud')

streaming_body_1 = client.get_object(Bucket='plantdiseaseidentification-donotdelete-pr-15huhdq',
    Key='data/vegetables/train.csv')

# Your data file was loaded into a botocore.response.StreamingBody object.
# Please read the documentation of boto3 and pandas to learn more about the possibilities to load the data.
# boto3 documentation: https://boto3.amazonaws.com/v1/documentation/api/latest/guide/quickstart.html#python
# pandas documentation: http://pandas.pydata.org/
```

Figure.6.6. Training Vegetable Dataset in IBM Cloud

Out put



← ↻ https://harvestify.herokuapp.com/crop-recommend

Enter the value (example:50)

Phosphorous

Enter the value (example:50)

Pottasium

Enter the value (example:50)

ph level

Enter the value

Rainfall (in mm)

Enter the value


State

Select State ▾

City

▾

Predict

 HARVESTIFY

Home Crop Fertilizer Disease

Get informed advice on fertilizer based on soil

Nitrogen

Enter the value (example:50)

Phosphorous

Enter the value (example:50)

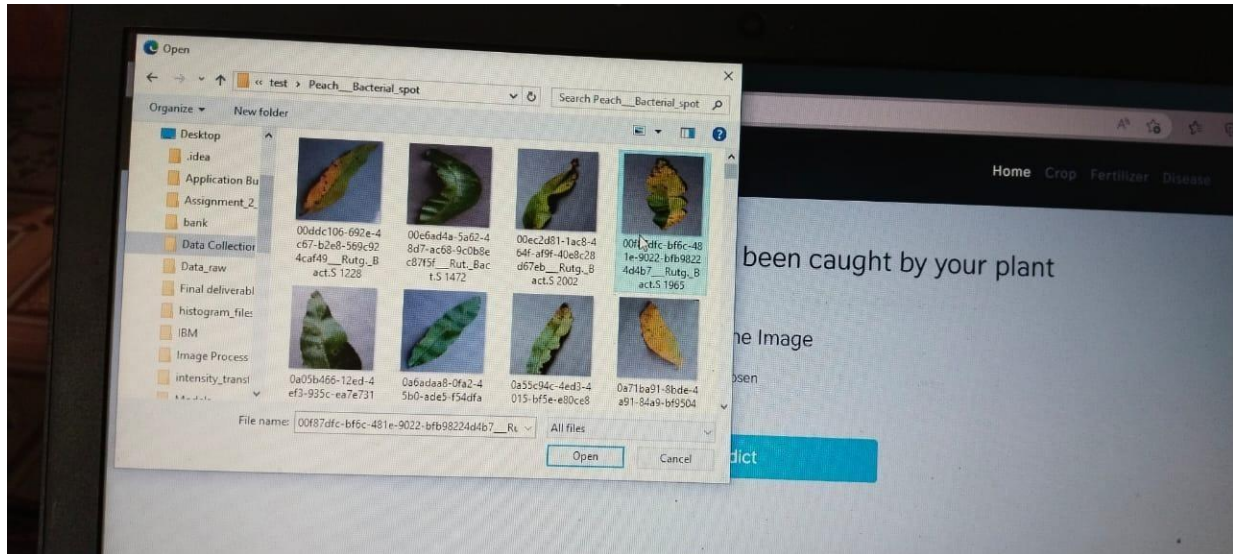
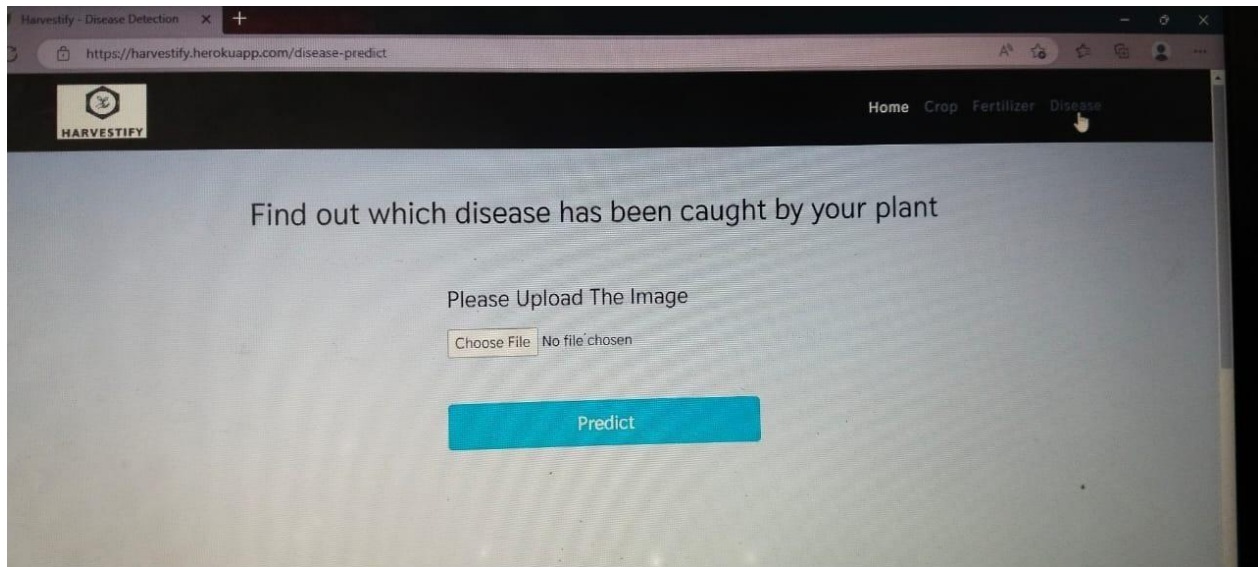
Pottasium

Enter the value (example:50)

Crop you want to grow

Select crop ▾

Predict



Find out which disease has been caught by your plant

Please Upload The Image

Choose File 00f87dfc-bf6c-...Bact.S 1965.JPG



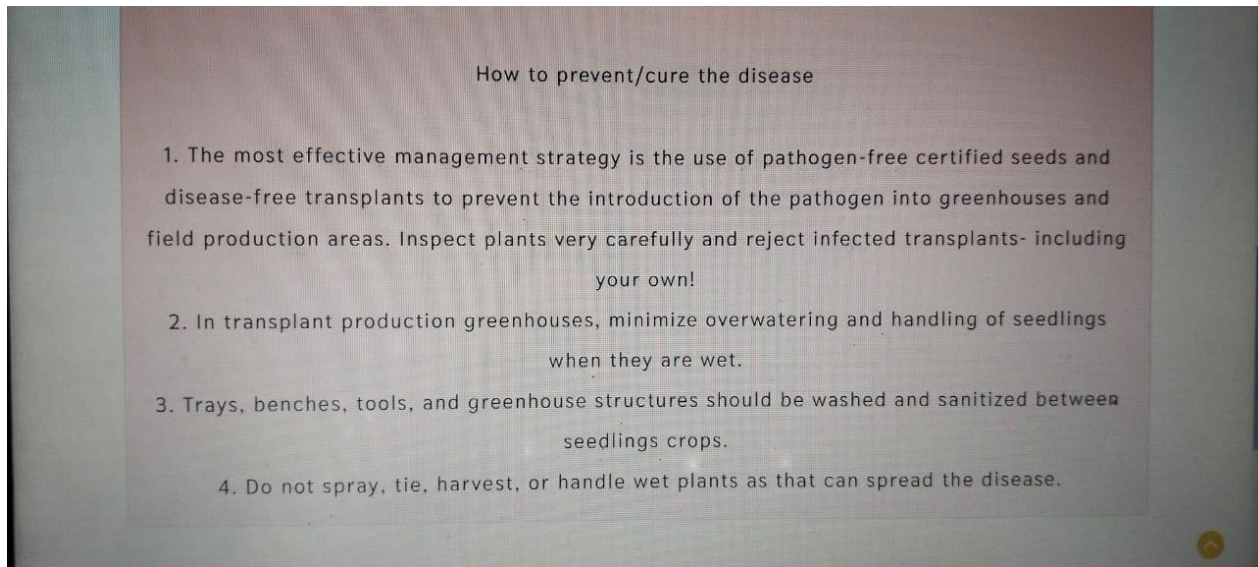
Predict

Crop: Peach

Disease: Bacterial Spot

Cause of disease:

1. The disease is caused by four species of *Xanthomonas* (*X. euvesicatoria*, *X. gardneri*, *X. perforans*, and *X. vesicatoria*). In North Carolina, *X. perforans* is the predominant species associated with bacterial spot on tomato and *X. euvesicatoria* is the predominant species associated with the disease on pepper.
2. All four bacteria are strictly aerobic, gram-negative rods with a long whip-like flagellum (tail) that allows them to move in water, which allows them to invade wet plant tissue and cause infection.



7.ADVANTAGES & DISADVANTAGES

List of advantages

- The model that is being suggested here achieves extremely high categorization accuracy.
- It is also possible to train on and test very big datasets.\
- Very high resolution images can be scaled inside the proposal itself.

List of disadvantages

- The suggested model has a very high computational time requirement for both training and testing.
- This project's utilisation of a neural network design involves great levels of complexity.

8.APPLICATIONS

1. The image patterns were accurately classified using the trained network model.
2. The suggested model is employed not only for the classification of plant diseases but also for the classification of other image patterns, such as animal classification.

3. This project work application uses pattern recognition in addition to image classification.

9.CONCLUSIONS

The model here involves classifying images from datasets of fruits and vegetables. Observations made during model testing and training include the following:

- The number of epochs was increased to boost categorization accuracy.
- Different classification accuracies are found for various batch sizes.
- By adding more convolution layers, the accuracy is increased.
- The use of several dense layers significantly improved categorization accuracy.
- By adjusting the size of the kernel utilised in the convolution layer output, different accuracies can be achieved.
- The accuracy varies with the size of the train and test datasets.

10.FUTURE SCOPE

The model that is being provided in this project work can be expanded to recognise images. Using python to exe software, the complete model may be turned into application software. With the aid of the OpenCV Python library, real-time image classification, image recognition, and video processing are all made possible. This project's work can be expanded to include security applications like face, iris, and figure print recognition.

11.BIBILOGRAPHY

[1]. R Indumathi Leaf Disease Detection and Fertilizer Suggestion, IEEE Conference on System, Computation, Automation and Networking (ICSCAN), March 29–30, 2019, DOI: 10.1109/ICSCAN.2019.8878781.

[2]. International Journal of Engineering Trends and Applications (IJETA) - Volume 8 Issue 2, Mar-Apr 2021, P. Pandi Selvi and P. Poornima, "Soil Based Fertilizer Recommendation System for Crop Disease Prediction System."

[3]. H Shiva reddy, Ganesh hedge, Prof. DR Chinnaya3, "IoT based Leaf Disease Detection and Fertilizer Recommendation", International Research Journal of Engineering and Technology (IRJET), Volume: 06 Issue: 11, Nov 2019, e-ISSN: 2395- 0056.

APPENDIX

A. Source Code (Jupyter notebook python code)

fruit.ipynb (due to limited page size the code vegetable.ipynb uploaded in github)

```
#!/usr/bin/env python
```

```
# coding: utf-8
```

```
# In[1]: pwd
```

```
# In[2]: cd E:/IBM_MY_COURSE/Project/Dataset Plant Disease/fruit-  
dataset/fruit-dataset
```

```
## Apply ImageDataGenerator functionality to Train and Test set
```

```
## Preprocessing # In[3]: from keras.preprocessing.image import  
ImageDataGenerator train_datagen =
```



```

ImageDataGenerator(rescale=1./255,shear_range=0.2,zoom_range=0.2,horizontal_
flip=True) test_datagen = ImageDataGenerator(rescale=1) # In[4]: pwd

#                               In[5]:                               x_train                               =
train_datagen.flow_from_directory('E:/IBM_MY_COURSE/Project/Dataset
Plant                               Disease/fruit-
dataset/fruitdataset/train',target_size=(128,128),batch_size=32,class_mode='cate
gorical')

#                               In[6]:
x_test=test_datagen.flow_from_directory('E:/IBM_MY_COURSE/Project/Datas
et    Plant    Disease/fruit-dataset/fruit-dataset/test',target_size=(128,128),
batch_size=32,class_mode='categorical') # # Import the models

#   In[7]:   from   tensorflow.keras.models   import   Sequential   from
tensorflow.keras.layers import Dense,Convolution2D,MaxPool2D,Flatten

# # Initializing the models 10

# In[8]: model=Sequential()

# # Add CNN Layers

#                               In[9]:
model.add(Convolution2D(32,(3,3),input_shape=(128,128,3),activation='relu'))

# In[10]: x_train.class_indices

# # Add Pooling layer

# In[11]: model.add(MaxPool2D(pool_size=(2,2)))

# # Add Flatten layer # In[12]: model.add(Flatten())

# # Add Dense Layer

```

```

# In[21]: model.add(Dense(40, kernel_initializer='uniform',activation='relu'))
model.add(Dense(20, kernel_initializer='random_uniform',activation='relu'))

# # Add Output Layer # In[24]: model.add(Dense(6,activation='softmax',
kernel_initializer='random_uniform'))

# # Compile the model # In[25]:
model.compile(loss='categorical_crossentropy',optimizer='adam',metrics=['accur
acy' ]) # In[26]: len(x_train)

# In[27]: 5384/32

# # Fit the Model

# In[28]:
model.fit_generator(x_train,steps_per_epoch=168,validation_data=x_test,validat
ion_steps=52,epochs=3)

# # Save the Model

# In[29]: model.save("fruit.h5")

# In[30]: ls

# # Test the Model

# In[32]: from keras.preprocessing import image from
tensorflow.keras.preprocessing.image import img_to_array from
tensorflow.keras.models import load_model import numpy as np

# In[33]: model = load_model("fruit.h5")

# # Test Apple_Healthy Class images

```

```

# In[37]: img = image.load_img('E:/IBM_MY_COURSE/Project/Dataset Plant
Disease/fruitdataset/fruit-dataset/test/Apple___healthy/00fca0da-2db3-481b-
b98a9b67bb7b105c___RS_HL_7708.JPG',target_size=(128,128)) 11

# In[39]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0)

# In[40]: pred = model.predict_classes(x)

# In[41]: pred

#                                     In[45]:                               index
=['Apple___Black_rot','Apple___healthy','Corn_(maize)___Northern_Leaf_Blig
ht','Corn_(maize)___healthy','Peach___Bacterial_spot','Peach___healthy']

# In[46]: print('the given image belongs to=',index[pred[0]])

# # Test Apple Black Rot class images # In[54]: img =
image.load_img('E:/IBM_MY_COURSE/Project/Dataset Plant
Disease/fruitdataset/fruit-dataset/test/Apple___Black_rot/0f3d45f4-e121-42cd-
a5b6-be2f866a0574___JR_FrgE.S_2870.JPG',target_size=(128,128))

# In[55]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0) pred =
model.predict_classes(x) print('the given image belongs to=',index[pred[0]])

# # Test Corn Northern leaf Blight class images

# In[56]: img = image.load_img('E:/IBM_MY_COURSE/Project/Dataset Plant
Disease/fruitdataset/fruit-
dataset/test/Corn_(maize)___Northern_Leaf_Blight/00a14441-7a62-4034-bc40-
b196aeab2785___RS_NLB_3932.JPG',target_size=(128,128))

# In[57]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0) pred =
model.predict_classes(x) print('the given image belongs to=',index[pred[0]])

```

```

# # Test Corn Healthy class images # In[58]: img =
image.load_img('E:/IBM_MY_COURSE/Project/Dataset Plant
Disease/fruitdataset/fruit-dataset/test/Corn_(maize)___healthy/0a68ef5a-027c-
41ae-b227-159dae77d3dd___R.S_HL 7969 copy.jpg',target_size=(128,128))

# In[59]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0) pred =
model.predict_classes(x) print('the given image belongs to=',index[pred[0]]) # #
Test Peach Bacterial spot class images # In[60]: img =
image.load_img('E:/IBM_MY_COURSE/Project/Dataset Plant
Disease/fruitdataset/fruit-dataset/test/Peach___Bacterial_spot/00ddc106-692e-
4c67-b2e8-569c924caf49___Rutg._Bact.S 1228.JPG',target_size=(128,128)) 12
# In[61]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0) pred =
model.predict_classes(x) print('the given image belongs to=',index[pred[0]])

# # Test Peach Healthy class images

# In[62]: img = image.load_img('E:/IBM_MY_COURSE/Project/Dataset Plant
Disease/fruitdataset/fruit-dataset/test/Peach___healthy/1a07ce54-f4fd-41cf-
b088-144f6bf71859___Rutg._HL 3543.JPG',target_size=(128,128))

# In[63]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0) pred =
model.predict_classes(x) print('the given image belongs to=',index[pred[0]])

```