FERTILIZERS RECOMMENDATION SYSTEM FOR DISEASE PREDICTION

USING ARTIFICIAL INTELLIGENCE

A PROJECT REPORT

Submitted by

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CONTENT

1. INTRODUCTION

- 1. Project Overview
- 2. Purpose

2. LITERATURE SURVEY

- 1. Existing problem
- 2. References
- 3. Problem Statement Definition

3. IDEATION & PROPOSED SOLUTION

- 1. Empathy Map Canvas
- 2. Ideation & Brainstorming
- 3. Proposed Solution
- 4. Problem Solution fit

4. REQUIREMENT ANALYSIS

- 1. Functional requirement
- 2. Non-Functional requirements

5. PROJECT DESIGN

- 1. Data Flow Diagrams
- 2. Solution & Technical Architecture
- 3. User Stories

6. PROJECT PLANNING & SCHEDULING

- 1. Sprint Planning & Estimation
- 2. Sprint Delivery Schedule
- 3. Reports from JIRA

7. CODING & SOLUTIONING (Explain the features added in the project along with code)

- 1. Feature 1
- 2. Feature 2
- **8.** Database Schema (if Applicable)

9. TESTING

- 1. Test Cases
- 2. User Acceptance Testing

10. RESULTS

1. Performance Metrics

11. ADVANTAGES & DISADVANTAGES

- 12. CONCLUSION
- 13. FUTURE SCOPE

14. APPENDIX

Source Code

GitHub & Project Demo Link

1. INTRODUCTION

PROJECT OVERVIEW

Agriculture is the most important sector in today's life. Most plants are affected by a wide variety of bacterial and fungal diseases. Diseases on plants placed a major constraint on the production and a major threat to food security. Hence, early and accurate identification of plant diseases is essential to ensure high quantity and best quality. In recent years, the number of diseases on plants and the degree of harm caused has increased due to the variation in pathogen varieties, changes in cultivation methods, and inadequate plant protection techniques.

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.

PURPOSE

This project is used to test the fruits and vegetables samples and identify the different diseases. Also ,this project recommends fertilizers for predicted diseases Agriculture serves as a means of supplying food to a population that is always expanding, as well as a significant source of energy and a means of combating globalwarming.

Plant diseases are very important because they can have a negative impact on the quality and quantity of crops produced in agriculture. Early detection of plant diseases is crucial for their treatment and management. Typically, illnesses are identified using the naked eye technique. Experts who can recognise variations in leaf colour are involved in this process. This method requires a lot of work, takes a while, and is not appropriate for fields with a lot of space. They forecast plant disease and recommend fertiliser for the damaged plants. This frequently involves a range of methods for assessing the qualities of the herbs that largely influence the plants. These complex systems that contain a large amount of datasets are forecasted using the neural network. By using artificial intelligence, complex manual systems' working models can be made simpler and more precise

2. LITERATURE SURVEY

EXISTING PROBLEM

Proposed a method for leaf disease detection and suggest fertilizers to cure leaf diseases. But the method involves less number of train and test sets which results in poor accuracy. Also proposed a simple prediction method for soil based fertilizer recommend ratio system for predicted crop diseases.

This method gives less accuracy and prediction. proposed an IoT based system for leaf disease detection and fertilizer recommendation which is based on Machine Learning techniques yields less 80 percentage accuracies.

REFERENCES

- [1]. R Indumathi.; N Saagari.; V Thejuswini.; R Swarnareka.," Leaf Disease Detectionand Fertilizer Suggestion", IEEE International Conference on System, Computation, Automation and Networking (ICSCAN), 29-30 March 2019, DOI: 10.1109/ICSCAN.2019.8878781.
- [2]. P. Pandi Selvi, P. Poornima, "Soil Based Fertilizer Recommendation System for Crop Disease Prediction System", International Journal of Engineering Trends and Applications (IJETA) Volume 8 Issue 2, Mar-Apr 2021.
- [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.

PROBLEM STATEMENT DEFINITION

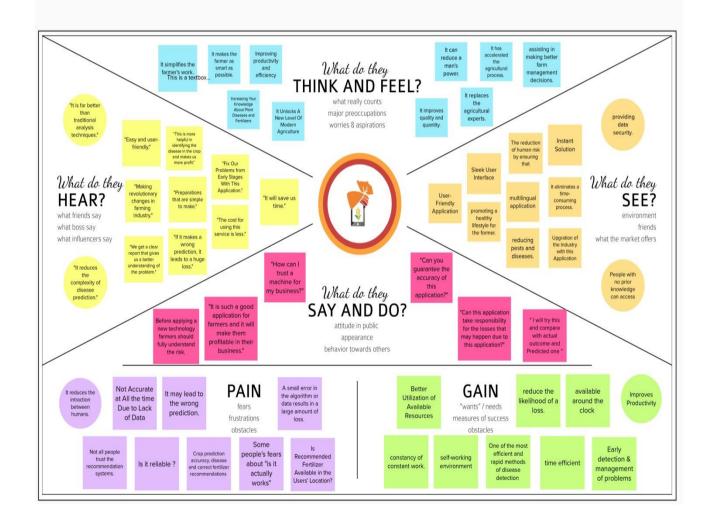
- Preprocess the images.
- Applying the CNN algorithm to the dataset.
- How deep neural networks detect the disease.
- You will be able to know how to find the accuracy of the model.
- You will be able to build web applications using the Flask framework.

3. IDEATION & PROPOSED SOLUTION

EMPATHY MAP CANVAS

Fertilizers Recommendation System For Disease Prediction

Agriculture is the main aspect of the economic development of a country. Agriculture is the heart and life of most Indians. By understanding their feelings and problems, we can create a better product and contribute to their lives. For our project, we are getting surveys from farmers to understand what they truly require and desire.



IDEATION & BRAINSTROMING



Brainstorm solo

Have each participant begin in the "solo brainstorm space" by sliently brainstorming ideas and placing them into the template. This "silent-storming" avoids group-think and creates an inclusive environment for introverts and extroverts alike. Set a time limit. Encourage people to go for quantity.

O 10 minutes

NANDHINI R

Can create a Identify User smart chat preferences bot to clear

Automation resolution of low quality

insuctive UI

SOWMIYA G

who use this

port for th

information about how

Securing

the data of

the user

SENTHIL KUMAR

Build kerss image classification model	Responsive and easy to interact UI	The application must be cros platform
	-	-

amount of

as input to predict the amount of

VELU N

images of the healthy trops after the use of fertilizer	several image processing algorithms	mode and light mode for the application UI
Based on the specifical suggest the diseases that	Based on the crop suggest the diseases that the crop	Suggest fertilizer to prevent the relapse of th

Cut review with Empire. Setting dark

Brainstorm as a group

Have everyone move their ideas into the "group sharing space" within the template and have the team silently read through them. As a team, sort and group them by thematic topics or similarities. Discuss and answer any questions that arise. Encourage "Nes. and.." and build on the ideas of other people along the way.

O 15 Harusan

Provide location of the store where the fertilizer is available

Responsive and easy to interact UI

Automation resolution of low quality images

Generating report for the suggested fertilizer

Provide offers and discounts who use this application

Suggest fertilizer to prevent the relapse of the disease

PROBLEM STATEMENTS

Problem Statement (PS)	I am	I'm trying to	But	Because	Which makes mefeel
PS-1	a farmer	grow and harvest healthy crops	my crops are affected by some disease	8	sad and disappointed to seethe diseased crops
PS-2	a gardener	grow some vegetable plants in the home	there are spots appearing on the leaves	of not maintaining and providingthe needed nutrients	not productive and not useful
PS-3	a nursery manager	grow and sell the plant saplings	the baby plants are not growing after some stage	not doing the needed tasks in order to maintain the growth	unprofitable in the future



PROPOSED SOLUTION

Proposed Solution Template:

Project team shall fill the following information in proposed solution template.

S.No	Parameter	Description
1.	Problem Statement (Problem to be solved)	Agriculture is the most important sector in today's life. Most plants are affected by a wide variety of bacterial and fungal diseases. on plants placed a major constraint on the production and a major threatto food security. Hence, early and accurate identification of plant diseases is essential to ensure high quantity and best quality. In recent years, the number of diseases on plants and the degree of harm caused has increased due to the variation in pathogen varieties, changes in cultivation methods, and inadequate plant protection techniques. So, based on above situation, how to help the farmers to choose the right fertilizer based upon the data given by them to promote productivity as well as eradicate the crop disease? What kind of model is suitable for this task? These are the main problems which are needed to be considered Here in this scenario.
2.	Idea/Solution description	The proposed idea is based on the usage of CNN primarily. The images with respective class names are made to train using CNN to classify the disease present, then based upon The results, the required fertilizer will be suggested to the user via the web interface.

3.	Novelty/Uniqueness	The novelty of the project lies on the parameters used on the CNN model.
		Another aspect is the usage of both
		sample and
		Real time data during model training.
4.	Social Impact/ Customer	The proposed idea will impact the
	Satisfaction	farmers a lot such that the right kind
		of fertilizer will be suggested to
		eradicate the diseases present in crops
		as well as increase the production
		By promoting their growth. The
		farmer can have insights about the
		fertilizers as well.
5.	Business Model(Revenue	The proposed model can be deployed
	Model)	both in web as well as containerized
		model (docker,k8s)for enterprise
		usage. The Web app will be made for
		free with certain limitations to all
		users. The enterprise app could be
		provided to businesses as standalone
		app which they could run on their own
		servers and then provide
		services to consumers for specific
		amount.
6	Scalability of the Solution	At present, the model will be trained
		with certain crop diseases and
		provide suggestion son the fertilizer
		to use on the crops. By time, the
		model will be trained with more
		images
		To classify more crop diseases to
		predict and suggest fertilizers in
		efficient way.
		-

4. REQUIRMENT ANALYSIS

FUNCTIONAL REQUIREMENTS

Functional Requirements:

Following are the functional requirements of the proposed solution.

FR	Functional	Sub Requirement (Story /
No.	Requirement (Epic)	Sub-Task)
FR-1	User Registration	Registration through Form
		Registration through Gmail
		Registration through LinkedIN
FR-2	User Confirmation	Confirmation via Email
		Confirmation via OTP
FR-3	User Login	Login with user name Login
		with password
FR-4	Profile update	Update the user credentials
		Update the Contact details
FR-5	Uploading Images	Upload the image of the
		affected Crop
FR-6	Image processing	Upload image will be
		processed to predict the
		disease in leaf
FR-7	Recommendation	User will request the fertilizer
		Get the fertilizer
		recommendations as per the
		disease
FR-8	Ratings and Reviews	Share their Experiences Give
		the Feedback

NON-FUNCTIONAL REQUIREMENTS

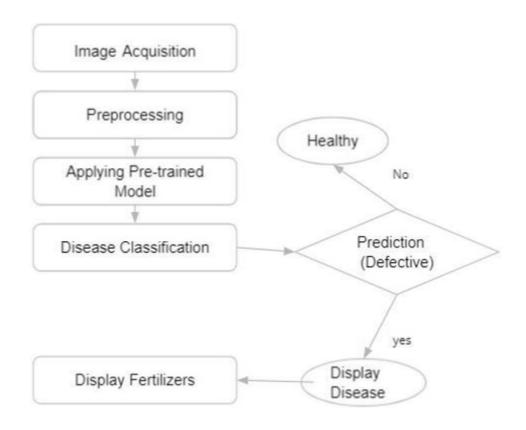
Non-functional Requirements:

Following are the non-functional requirements of the proposed solution.

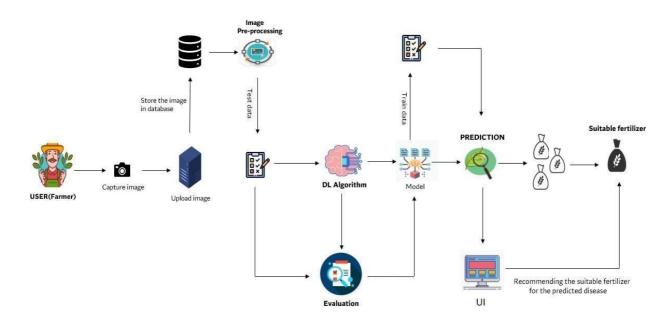
FR	Non-Functional	Description
No.	Requirement	
NFR-1	Usability	It can be used on both website and
		mobile browsers so that the
		usability of this application is very efficient.
NFR-2	Security	The information belongs to user and
		about leaf is highly secured.
NFR-3	Reliability	The leaf quality is necessary to
		predict the disease in the leaf.
NFR-4	Performance	The performance of application is
		very fast and efficient, based on the
		prediction of the disease.
NFR-5	Availability	It will predict any type of new
		disease by
		learning from the available data and
		predict the disease accurately.
NFR-6	Scalability	It can be accessed by a greater
		number of users at the same time
		without any
		performance issues.

5. PROJECT DESIGN

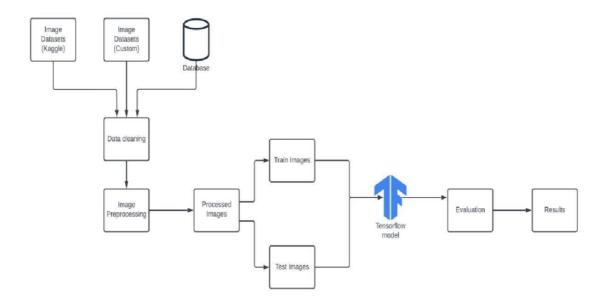
DATA FLOW DIAGRAMS



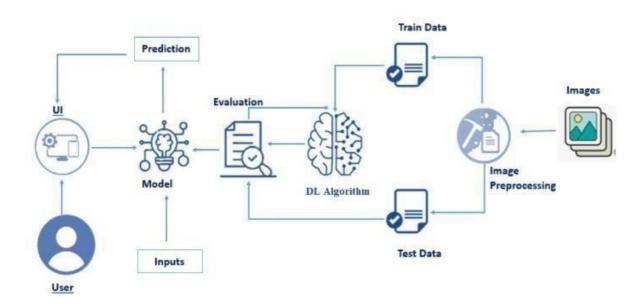
SOLUTION ARCHITECTURE



Plant crop disease is anticipated, and appropriate fertilizer is advised for a higher yield. The diseased plant photos are acquired and preprocessed in comparison to the dataset of diseased plants. The photos are processed using a Deep Learning algorithm, which is subsequently tested. A model is then created based on the evaluations, trained using a variety of inputs, and predictions are presented to the users, aiding in the fertilizer recommendation process. The inclusion of the Convolutional layers in the classification and processing of the images further aids in



TECHNICAL ARCHITECTURE



USER STORIES

A digital camera or similar devices are used to take images of different types, and then those are used to identify the affected area in leaves. Then different types of image-processing techniques are applied to them, the process those images, to get different and useful features needed for the purpose of analyzing later-Plant leaf disease identification is especially needed to predict both the quality and quantity of the First segmentation step primarily based on a mild polygonal leaf model is first achieved and later used toguide the evolution of an energetic contour. Combining global shape descriptors given by the polygonal model with local curvature

User Type	Functional Requirement (Epic)	User Story Number	User Story / Task	Acceptance criteria	Priority
Customer (Mobile user)	Download the database	USN-1	As a user, I can register for the application by entering my email,password, and confirming my password.	I can access my account/ dashboard	High
	Register	USN-2	As a user, I can register for the application by entering my email,password, and confirming my password.	I can receive confirmati on email & click confirm	High
	Login	USN-3	As a user, I will receive confirmation email once I have registered for the application	I can register & access the dashboard With Login	Low
	Upload the image	USN-4	As a user, I must upload the imageto identify the disease		Medium
Customer (Webuser)	The functional requirements Are same asmobile user	USN-5	Same as mobile user can fill thedetail asked here	Same as mobile user can register the details	High(when compare dto mobile users)
Administrat or		USN-6	As the administrator I predict theoutput	Show the result	High

6. PROJECT PLANNING & SCHEDULING

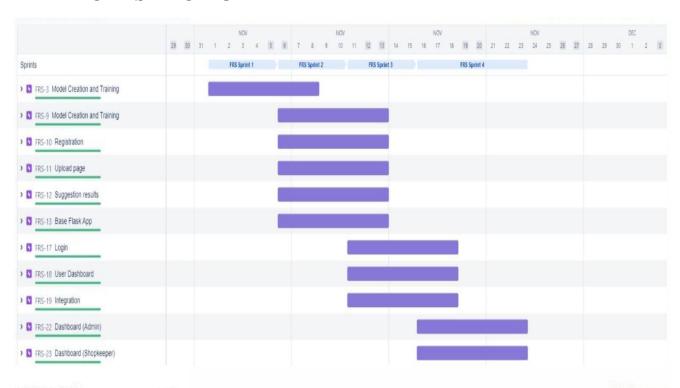
SPRINT PLANNING & ESTIMATION

Sprint	Functional Requireme nt (Epic)	User Story Numb er	User Story/Task	Story Points	Priority	Team Members
Sprint-1	Registration	USN-1	Collecting plant disease dataset	2	High	Nandhini R
Sprint-1		USN-2	Labeling the dataset according to class	1	High	Sowmiya G
Sprint-2	Testing training and creating a model	USN-3	Start initiating the model	2	Low	Senthilkumar M
Sprint-1		USN-4	Adding different layers of CNN(convolution, pooling dense, flat ten)	2	Medium	Velu N
Sprint-1		USN-5	Training the data	1	High	Sowmiya G, Nandhini R
Sprint-3			Testing the data	1	Medium	Senthilkumar M, Velu N
Sprint-3	Flask and framework design		Creating backend framework with flask	2	High	Nandhini R, SowmiyaG, Senthilkumar M, Velu N
Sprint-4			Predicting disease and recommend fertilizer	2		Nandhini R, Sowmiya G, Senthilkumar M, Velu N

SPRINT DELIVERY SCHEDULE

Sprint	Total Story Points	Duration	Sprint Start Date	Sprint End Date (Planned)	Story Points Completed (as on Planned End Date)	Sprint Release Date (Actual)
Sprint-1	20	6Days	24Oct2022	29Oct2022	10	05 NOV 2022
Sprint-2	20	6Days	31Oct2022	05Nov2022	15	10 Nov 2022
Sprint-3	20	6Days	07Nov2022	12Nov2022	15	15 Nov 2022
Sprint-4	20	6Days	14Nov2022	19Nov2022	10	20 Nov 2022

REPORTS FROM JIRA



6. CODING & SOLUTIONING

FEATURE 1

The application's registration page is created. User registration is carried out if the user hasn't already done so. Enough work was put into making this process seamless. If the user has registered, he can now log in directly. Email address, name, and password were required for registration.

MODEL BUILDING

1. Import The Libraries

Import the libraries that are required to initialize the neural network layer, and create and add different layers to the neural network model.

```
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
```

2. Initializing The Model

Keras has 2 ways to define a neural network:

- Sequential
- Function API

The Sequential class is used to define linear initializations of network layers which then, collectively, constitute a model. In our example below, we will use the Sequential constructor to create a model, which will then have layers added to it using the add () method. Now, will initialize our model. Initialize the neural network layer by creating a reference/object to the Sequential class.

```
model=Sequential()
```

3. ADD CNN Layers

We will be adding three layers for CNN

- Convolution layer
- Pooling layer
- Flattening layer

Add Convolution Layer

The first layer of the neural network model, the convolution layer will be added. To create a convolution layer, Convolution2D class is used. It takes a number of feature detectors, feature detector size, expected input shape of the image, and activation function as arguments. This **19** layer applies feature detectors on the input image and returns a feature map (features from the image).

Activation Function: These are the functions that help us to decide if we need to activate the node or not. These functions introduce non-linearity in the networks.

```
model.add(Convolution2D(32,(3,3),input_shape = (128,128,3),activation = 'relu'))
```

Add the 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. After the convolution layer, a pooling layer is added. Max pooling layer can be added using MaxPooling2D class. It takes the pool size as a parameter. Efficient size of the pooling matrix is (2,2). It returns the pooled feature maps. (Note: Any number of convolution layers, pooling and dropout layers can be added)

```
model.add(MaxPooling2D(pool_size = (2,2)))
```

Add the 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 layers.

```
model.add(Flatten())
```

4. Add Dense Layers

Now, let's add Dense Layers to know more about dense layers click below

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.

Adding Hidden layers

This step is to add a dense layer (hidden layer). We flatten the feature map and convert it into a vector or single dimensional array in the Flatten layer. This vector array is fed it as an input to the neural network and applies an activation function, such as sigmoid or other, and returns the output.

- init is the weight initialization; function which sets all the weights and biases of a network to values suitable as a starting point for training.
- units/ output_dim, which denote is the number of neurons in the hidden layer.
- The activation function basically decides to deactivate neurons or activate them to get the desired output. It also performs a nonlinear transformation on the input to get better results on a complex neural network.
- You can add many hidden layers, in our project we are added two hidden layers. The 1st hidden layer with 40 neurons and 2nd hidden layer with 20neurons.

Adding the output layer

This step is to add a dense layer (output layer) where you will be specifying the number of classes your dependent variable has, activation function, and weight initializer as the arguments. We use the add () method to add dense layers. the output dimensions here is 6

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

5. Train And Save The Model

Compile the model

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

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

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

```
model.fit_generator(x_train, steps_per_epoch = 168,epochs = 3,validation_data = x_test,validation_steps = 52)
```

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 file using save().

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

model.summary() can be used to see all parameters and shapes in each layer in our models.

6. Test The Model

The model is to be tested with different images to know if it is working correctly.

Import the packages and load the saved model

Import the required libraries

```
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
```

Initially, we will be loading the fruit model. You can test it with the vegetable model in a similar way.

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

Load the test image, pre-process it and predict Pre-processing the image includes converting the image to array and resizing according to the model. Give the pre-processed image to the model to know to which class your model belongs to.

```
img = image.load_img('apple_healthy.JPG',target_size = (128,128))
```

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

pred = model.predict_classes(x)

pred
[1]
```

The predicted class is 1.

FEATURE 2

Python Code

After the model is built, we will be integrating it into a web application so that normal users can also use it. The user needs to browse the images to detect the disease.

To Build a flask application do the following:

Step 1: Load the required packages

Step 2: Initialize the flask app and load the model

Step 3: Configure the home page

Step 4: Pre-process the frame and run

The trained machine learning model can predict the output from an image that is uploaded, and the nutrition facts are also displayed on the same page. The model's accuracy was determined to be 95%, and when it was trained on the IBM cloud, it reached 100%.

7. TESTING

TEST CASES

The test cases include invalid email and unrecognizable images. For the image part, a text fileor other format files were uploaded as a corner case.

USER ACCEPTANCE TESTING

Acceptance testing is performed on a collection of business functions in a Production environment and after the completion of functional testing. This is the final Stage in the testing process before the system is accepted for operational use. This testing should be done with original data and with the presence of the users. This test confirms the systemready for production.

Purpose of Document

The purpose of this document is to briefly explain the test coverage and open issues of the [Fertilizer Recommendation system for plant disease prediction] project at the time of the release to User Acceptance Testing(UAT).

Defect Analysis

This report shows the number of resolved or closed bugs at each severity level, and how they were resolved.

Resolution	Severity 1	Severity 2	Severity 3	Severity 4	Subtot al
Leaf spots	10	4	2	3	19
Mosaic leaf pattern	9	6	3	6	24
Misshapen leaves	2	7	0	1	10
Yellow leaves	11	4	3	20	38
Fruit rots	3	2	1	0	6
Fruit spots	5	3	1	1	10
Blights	4	5	2	1	12
Totals	44	31	13	32	11 9

8. RESULTS

PERFORMANCE METRICS

Model Performance Testing:

Project team shall fill the following information in model performance testing template.

S.No.	Parameter	Values	Screenshot
1.		Totalparams:896 Trainableparams:896 Non- trainableparams:0	model.summary() Model: "sequential" Laver (type) Output Shape Param #
			conv2d (Conv2D) (None, 126, 126, 32) 896 max_pooling2d (MaxPooling2D (None, 63, 63, 32) 0) flatten (Flatten) (None, 127008) 0
	Accuracy	Training Accuracy– 96.55	Total params: 896 Trainable params: 896 Non-trainable params: 0 (spoth 1780 25/225 [
	recuruey	ValidationAccuracy-97.45	S88 9986/step - 10810 - 20225 - accuracy: 8-9042 - val_loss: 0-3015 - val_accuracy: 8-070
			655 3768/1409 - 855 37686/1409 - 10555 0.1280 - accuracy; 0.9580 - val_loss; 0.1380 - val_accuracy; 0.9573 185 37786/1409 - 10555 0.1285 - accuracy; 0.9585 - val_loss; 0.1580 - val_accuracy; 0.9573 185 37786/1409 - 10555 0.1295 - accuracy; 0.9585 - val_loss; 0.1680 - val_accuracy; 0.9586 -

Model Summary

Accuracy

```
l.fit_generator(x_train,steps_per_epoch=len(x_train),validation_data=x_test,validation_steps=len(x_test),epochs=10)
sers\Sree Ram\AppData\Local\Temp\ipykernel 13228\1582812018.py:1: UserWarning: `Model.fit generator` is deprecated and will
emoved in a future version. Please use `Model.fit`, which supports generators.
lel.fit_generator(x_train,steps_per_epoch=len(x_train),validation_data=x_test,validation_steps=len(x_test),epochs=10)
1/10
1 2/10
225 [================ ] - 88s 393ms/step - loss: 0.2825 - accuracy: 0.9042 - val loss: 0.3015 - val accuracy:
75
1 3/10
1 4/10
225 [========== ] - 84s 374ms/step - loss: 0.1576 - accuracy: 0.9463 - val loss: 0.2424 - val accuracy:
1 5/10
225 [=========================== - 84s 372ms/step - loss: 0.1719 - accuracy: 0.9389 - val loss: 0.1330 - val accuracy:
32
1 6/10
225 [=========: 0.1240 - accuracy: 0.9580 - val loss: 0.1340 - val accuracy:
1 7/10
78
1 8/10
225 [================= - 83s 371ms/step - loss: 0.1012 - accuracy: 0.9643 - val loss: 0.1468 - val accuracy:
51
1 9/10
1 10/10
```

9. ADVANTAGES & DISADVANTAGES

ADVANTAGES:

- The suggested model yields extremely high classification accuracy
- It can train and test on very large datasets.
- It can resize very high-quality images within itself.

DISADVANTAGES:

- Prediction is limited to few plants as we havent trained all the plants.
- The proposed model is computationally expensive to train and test.
- The neural network architecture used in this project work is highly complex.

10. CONCLUSION

The model here involves classifying images from datasets of fruits and vegetables. The number of epochs was increased to boost categorization accuracy. Different classification accuracies are obtained for different batch sizes. The accuracies are increased by adding more convolution layers. The accuracy of classification is also increased by adjusting the number of dense layers. The accuracies are different while varying the size of the train and test datasets.

11. 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.
- As of now we have just built the web application which apparently takes the input as an image and then predict the out in the near future we can develop an application which computer vision and AI techniques to predict the infection once you keep the camera near the plant or leaf this could make our project even more usable
- This further research is implementing the proposed algorithm with the existing public datasets. Also, various segmentation algorithms can be implemented to improve accuracy. The proposed algorithm can be modified further to identify the disease that affects the various plant organs such as vegetables and fruits.

12. APPENDIX

SAMPLE SOURCE CODE:-

```
import tensorflow as tf
import time import numpy as npimport os
start = time.time()
#try:
# Total iterations
final iter = 1000
# Assign the batch value
batch size = 20
# 20% of the data will automatically be used for validation
validation size = 0.2
img_size = 128
num channels = 3 train path = r'data\Train'
# Prepare input data
if not os.path.exists(train path):
print("No such directory")raise Exception
classes = os.listdir(train path)num classes = len(classes)
# We shall load all the training and validation images and labels into
memory using openCV and use that during training
data = dataset.read train sets(train path, img size, classes,
validation size=validation size)
# Display the stats
print("Complete reading input data. Will Now print a snippet of it")
print("Number of files in Training-
set:\t\t{}".format(len(data.train.labels)))print("Number of files in
Validation-set:\t{}".format(len(data.valid.labels))) session =
tf.compat.v1.Session()
x = tf.compat.v1.placeholder(tf.float32, shape=[None, img size, img size,
num channels], name='x')
## LabeLs
y true = tf.compat.v1.placeholder(tf.float32, shape=[None, num classes],
name='v true')
y true cls = tf.argmax(y true, dimension=1)
##Network graph params
    filter size conv1
    num filters conv1
    filter size conv2
    num filters conv2
```

```
filter_size_conv3
    num_filters_conv3
fc layer size = 128
def create weights(shape):
return tf. Variable(tf.random.truncated normal(shape, stddev=0.05))
def create biases(size):
return tf.Variable(tf.constant(0.05, shape=[size]))
def make generator model(input, num input channels, conv filter size,
num filters):
## We shall define the weights that will be trained using create weights
function. weights = create weights(shape=[conv filter size,
conv filter size, num input channels, num filters])
## We create biases using the create_biases function. These are also
trained.
biases = create_biases(num_filters)
## Creating the convolutional layerlayer = tf.nn.conv2d(input=input,
filter=weights,
strides=[1, 1, 1, 1], padding='SAME')
layer += biases
## We shall be using max-pooling. layer = tf.nn.max pool(value=layer,
ksize=[1, 2, 2, 1],
strides=[1, 2, 2, 1], padding='SAME')
## Output of pooling is fed to Relu which is the activation function for
us.
layer = tf.nn.relu(layer)
return layer
# Function to create a Flatten Layer
def create flatten layer(layer):
# We know that the shape of the layer will be [batch size img size
img sizenum channels]
# But let's get it from the previous layer.
layer_shape = layer.get_shape()
## Number of features will be img_height * imq width* num channels. But
we shall calculate it in place of hard-coding it.
num_features = layer_shape[1:4].num_elements()
## Now, we Flatten the layer so we shall have to reshape to num features
layer = tf.reshape(layer, [-1, num_features])
```

```
# Function to create a Fully - Connected Layer
def create fc layer(input, num inputs, num outputs, use relu=True):
# Let's define trainable weights and biases.
weights = create_weights(shape=[num_inputs, num outputs])biases =
create biases(num outputs)
# Fully connected layer takes input x and produces wx+b.Since, these are
matrices, we use matmul function in Tensorflow
layer = tf.matmul(input, weights) + biases
if use relu:
layer = tf.nn.relu(layer)
return layer
# Create all the layers
layer conv1 = make generator model(input=x,
num input channels=num channels, conv filter size=filter size conv1,
num filters=num filters conv1)
layer conv2 = make generator model(input=layer conv1,
num input channels=num filters conv1, conv filter size=filter size conv2,
num filters=num filters conv2)
layer conv3 = make generator model(input=layer conv2,
num input channels=num filters conv2, conv filter size=filter size conv3,
num filters=num filters conv3)
layer flat = create flatten layer(layer conv3)
layer fc1 = create fc layer(input=layer flat,
num inputs=layer flat.get shape()[1:4].num elements(),
num outputs=fc layer_size,
use relu=True)
layer fc2 = create fc layer(input=layer fc1, num inputs=fc layer size,
num outputs=num classes,
use relu=False)
y pred = tf.nn.softmax(layer fc2, name='y pred')
y pred cls = tf.argmax(y pred, dimension=1)
session.run(tf.compat.v1.global variables initializer())
cross entropy =
tf.nn.softmax_cross_entropy_with_logits_v2(logits=layer_fc2,
labels=y true)
cost = tf.reduce mean(cross entropy)
```

```
optimizer = tf.compat.v1.train.AdamOptimizer(learning rate=1e-
4).minimize(cost)correct prediction = tf.equal(y pred cls, y true cls)
accuracy = tf.reduce mean(tf.cast(correct prediction, tf.float32))
session.run(tf.compat.v1.global variables initializer())
# Display all stats for every epoch
def show progress(epoch, feed dict train, feed dict validate,
val losstotal epochs):
acc = session.run(accuracy, feed_dict=feed_dict_train) val_acc =
session.run(accuracy, feed_dict=feed_dict validate)
msg = "Training Epoch {0}/{4} --- Training Accuracy: {1:>6.1%},
ValidationAccuracy: {2:>6.1%}, Validation Loss: {3:.3f}"
print(msg.format(epoch + 1, acc, val acc, val loss, total epochs))
total iterations = 0
saver = tf.compat.v1.train.Saver()print("")
# Training Function
def train(num iteration):
global total_iterations
for i in range(total_iterations,
total iterations + num iteration):
x_batch, y_true_batch, _, cls_batch = data.train.next_batch(batch_size)
x_valid_batch, y_valid_batch, _, valid_cls_batch =
data.valid.next_batch(batch_size)
feed dict tr = \{x: x \text{ batch},
y true: y true batch}feed dict val = {x: x valid batch,
y true: y valid batch} session.run(optimizer, feed dict=feed dict tr)
if i % int(data.train.num_examples / batch_size) == 0:
val loss = session.run(cost, feed dict=feed dict val) epoch = int(i /
int(data.train.num examples / batch size))
# print(data.train.num examples)
# print(batch size)
# print(int(data.train.num_examples/batch_size))# print(i)
total epochs = int(num iteration / int(data.train.num examples /
batch_size)) + 1show_progress(epoch, feed_dict_tr, feed_dict_val,
val loss, total epochs)
saver.save(session, 'trained_model') total iterations += num iteration
```

```
train(num iteration=final iter)
#except Exception as e: #print("Exception:",e)
# Calculate execution time
end = time.time()dur = end-start print("")
if dur<60:
print("Execution Time:",dur,"seconds")elif dur>60 and dur<3600:</pre>
dur=dur/60
print("Execution Time:",dur,"minutes")else:
dur=dur/(60*60)
print("Execution Time:",dur,"hours")
from flask import Flask, render template, flash, request,
session, send file
from flask import render_template, redirect, url for, request
import warningsimport datetimeimport cv2
import tensorflow as tf
import numpy as np
from tkinter import *
import os
app = Flask( name )app.config['DEBUG']
app.config['SECRET KEY'] = '7d441f27d441f27567d441f2b6176a'
@app.route("/")
def homepage():
return render template('index.html')
@app.route("/Test")
def Test():
return render template('Test.html')
@app.route("/train", methods=['GET', 'POST'])def train():
if request.method == 'POST':
import model as model
return render_template('Tranning.html')
@app.route("/testimage", methods=['GET', 'POST'])def testimage():
if request.method == 'POST':
file = request.files['fileupload'] file.save('data/alien_test/Test.jpg')
```

```
img = cv2.imread('data/alien_test/Test.jpg')
train path = r'data\train'if not os.path.exists(train path): print("No
such directorv")
raise Exception
# Path of testing images dir_path = r'data\alien_test' if not
os.path.exists(dir_path):print("No such directory") raise Exception
# Walk though all testing images one by onefor root, dirs, files in
os.walk(dir path):for name in files:
print("")
print(filename)
num channels = 3images = []
image path = name
filename = dir_path + '\\' + image_pathimage_size =128
if os.path.exists(filename):
# Reading the image using OpenCV
image1 = cv2.imread(filename)
import_file_path = filename
image = cv2.imread(import file path)
fnm = os.path.basename(import file path)filename = 'Test.jpg'
cv2.imwrite(filename, image)
# print("After saving image:")
print("\n*************\nImage : " + fnm +
"\n****************")img = cv2.imread(import_file_path)
if img is None: print('no data')
print(img.shape)
img1 = cv2.imread(import file path)
img = cv2.resize(img, ((int)(img.shape[1] / 5),
(int)(img.shape[0] / 5)))
original = img.copy() neworiginal = img.copy() cv2.imshow('original',
img1)
gray = cv2.cvtColor(img1, cv2.COLOR BGR2GRAY)
cv2.imshow('Original image', img1)orimage = 'static/Out/Test.jpg'
cv2.imwrite(orimage, img1)
```

```
cv2.imshow('Gray image', gray)gry = 'static/Out/gry.jpg'
cv2.imwrite(gry, gray)
p = 0
for i in range(img.shape[0]):
for j in range(img.shape[1]):
B = img[i][j][0]
G = img[i][i][1]
R = img[i][j][2]
if (B >110 and G >110 and R >110):
p += 1
totalpixels = img.shape[0] * img.shape[1]
per white = 100 \cdot p / totalpixels
if per white >10:
# Guassian blur
img[i][j] = [500, 300, 200]
cv2.imshow('color change', img)
blur1 = cv2.GaussianBlur(img, (3, 3), 1)
# mean-shift algo
newimg = np.zeros((img.shape[0], img.shape[1], 3), np.uint8)
criteria = (cv2.TERM CRITERIA EPS +cv2.TERM CRITERIA MAX ITER, 10, 1.0)
img = cv2.pyrMeanShiftFiltering(blur1, 20, 30, newimg, 0,criteria)
cv2.imshow('means shift image', img)noise = 'static/Out/noise.jpg'
cv2.imwrite(noise, img)
# Guassian blur
blur = cv2.GaussianBlur(img, (11, 11), 1)
blur = cv2.GaussianBlur(img, (11, 11), 1)
# Canny-edge detection
canny = cv2.Canny(blur, 160, 290)
canny = cv2.cvtColor(canny, cv2.COLOR_GRAY2BGR)
# contour to find leafs
bordered = cv2.cvtColor(canny, cv2.COLOR_BGR2GRAY)
contours, hierarchy = cv2.findContours(bordered, cv2.RETR TREE,
cv2.CHAIN APPROX NONE)
maxC = 0 for x in range(len(contours)):if len(contours[x]) > maxC:
maxC = len(contours[x]) maxid = x
perimeter = cv2.arcLength(contours[maxid], True)
```

```
# print perimeter
Tarea = cv2.contourArea(contours[maxid])
cv2.drawContours(neworiginal, contours[maxid], -1, (0, 0,
255))
cv2.imshow('Contour', neworiginal)
# cv2.imwrite('Contour complete leaf.jpg',neworiginal)# Creating
rectangular roi around contour
height, width, _ = canny.shape
min x, min y = width, heightmax x = max y = 0
frame = canny.copy()
# computes the bounding box for the contour, and draws it on the frame,
for contour, hier in zip(contours, hierarchy):
(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, w, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, max x = min(x, y, h) = cv2.boundingRect(contours[maxid])min x, 
min_x), max(x + w, max_x)min_y, max_y = min(y, min_y), max(y + h, max_y)
if w >80 and h >80:
# cv2.rectangle(frame, (x,y), (x+w,y+h), (255, 0, 0), 2) #we do not draw
therectangle as it interferes with contour later on
roi = img[y:y + h, x:x + w]
originalroi = original[y:y + h, x:x + w]
if (max x - min x > 0 and max y - min y > 0):
roi = img[min_y:max_y, min_x:max_x]
originalroi = original[min_y:max_y, min_x:max_x] cv2.rectangle(frame,
(\min x, \min y), (\max x, \max y), (255,
0, 0),
2) # we do not draw the rectangle as it interferes with contour
cv2.imshow('ROI', frame)
roi12 = 'static/Out/roi.jpg'
cv2.imwrite(roi12, frame)
cv2.imshow('rectangle ROI', roi)
img = roi
# Changing colour-space
# imghsv = cv2.cvtColor(img, cv2.COLOR BGR2HSV)
imghls = cv2.cvtColor(roi, cv2.COLOR_BGR2HLS)
cv2.imshow('HLS', imghls)
imghls[np.where((imghls == [30, 200, 2]).all(axis=2))] = [0,
200, 01
# Only hue channel
cv2.imshow('new HLS', imghls)
huehls = imghls[:, :, 0]
cv2.imshow('img_hue hls', huehls)
# ret, huehls = cv2.threshold(huehls,2,255,cv2.THRESH BINARY)
huehls[np.where(huehls == [0])] = [35]
cv2.imshow('img_hue with my mask', huehls)
# Thresholding on hue image
ret, thresh = cv2.threshold(huehls, 28, 255, cv2.THRESH BINARY INV)
cv2.imshow('thresh', thresh)
```

```
# Masking thresholded image from original image
mask = cv2.bitwise_and(originalroi, originalroi, mask=thresh)
cv2.imshow('masked out img', mask)
# Resizing the image to our desired size and preprocessing will be done
exactly asdone during training
image = cv2.resize(image1, (image size, image size), 0, 0,
cv2.INTER LINEAR) images.append(image)
images = np.array(images, dtype=np.uint8)images =
images.astype('float32')
images = np.multiply(images, 1.0 / 255.0)
# The input to the network is of shape [None image size image size
num channels].Hence we reshape.
x batch = images.reshape(1, image size, image size, num channels)
# Let us restore the saved model
sess = tf.compat.v1.Session()
# Step-1: Recreate the network graph. At this step only graph is created.
saver = tf.compat.v1.train.import_meta_graph('models/trained_model.meta')
# Step-2: Now let's load the weights saved using the restore method.
saver.restore(sess, tf.train.latest checkpoint('./models/'))
# Accessing the default graph which we have restored
graph = tf.compat.v1.get_default_graph()
# Now, let's get hold of the op that we can be processed to get the
output.
# In the original network y_pred is the tensor that is the prediction of
the network
y pred = graph.get tensor by name("y_pred:0")
## Let's feed the images to the input placeholders
x = graph.get tensor by name("x:0")
y true = graph.get tensor by name("y true:0") y test images =
np.zeros((1, len(os.listdir(train path))))
# Creating the feed dict that is required to be fed to calculate y pred
feed_dict_testing = {x: x_batch, y_true: y_test_images}
result = sess.run(y_pred, feed_dict=feed_dict_testing)
# Result is of this format [[probabiliy_of_classA probability_of_classB
            print(result)
# Convert np.array to list
a = result[0].tolist()
r = 0
# Finding the maximum of all outputs
max1 = max(a)
index1 = a.index(max1)predicted class = None
```

```
# Walk through directory to find the label of the predicted output
count = 0
for root, dirs, files in os.walk(train path):
for name in dirs:
if count == index1:
predicted class = namecount += 1
# If the maximum confidence output is largest of all by a big margin then
# print the class or else print a warning
for i in a:
if i != max1:
if max1 - i < i:r = 1
out = ''
pre = "" if r == 0:
print(predicted class)
if (predicted_class == "Black spot"):
out = predicted class
pre = 'Griffin Fertilizer reducing the fungus'elif (predicted_class ==
"canker"):
out = predicted_class
pre = 'sprayed with Bordeaux mixture 1.0 per cent.'
elif (predicted class == "greening"):
out = predicted_class
pre = 'Mn-Zn-Fe-B micronutrient fertilizer'
elif (predicted class == "healthy"):
out = predicted class
#_messagebox.showinfo("Uses", '')
elif (predicted_class == "Melanose"):
out = predicted_class
pre = 'strobilurin fungicide'
else:
out = 'Could not classify with definite confidence'
else:
print("File does not exist")
org = 'static/Out/Test.jpg'
gry ='static/Out/gry.jpg' noise = 'static/Out/noise.jpg'roi12 =
```

```
'static/Out/roi.jpg'
return
render template('Test.html', result=out, org=org, gry=gry, inv=noise, noi=roi1
2, fer=pre)
def sendmsg(targetno,message):
import requests
requests.post("http://smsserver9.creativepoint.in/api.php?username=fantasy
&password=596692&to=" + targetno + "&from=FSSMSS&message=Dear user your
msg is " +
message + " Sent By FSMSG
FSSMSS&PEID=1501563800000030506&templateid=1507162882948811640")
if__name__== '__main__': app.run(debug=True, use reloader=True)
import cv2 import os import glob
from sklearn.utils import shuffle
import numpy as np
def load train(train path, image size, classes):images = []
labels = [] img names = []cls = []
print('Going to read training images')for fields in classes:
index = classes.index(fields)
print('Now going to read {} files (Index: {})'.format(fields, index))path
= os.path.join(train path, fields, '*g')
files = glob.glob(path)
for fl in files:
image = cv2.imread(f1)
image = cv2.resize(image, (image size, image size),0,0,cv2.INTER LINEAR)
image = image.astype(np.float32)
image = np.multiply(image, 1.0 / 255.0)images.append(image)
label = np.zeros(len(classes))label[index] = 1.0
labels.append(label)
flbase = os.path.basename(fl)img_names.append(flbase) cls.append(fields)
images = np.array(images) labels = np.array(labels) img names =
np.array(img names) cls = np.array(cls)
return images, labels, img names, cls
class DataSet(object):
def_init (self, images, labels, img names, cls):self. num examples =
images.shape[0]
self._images = images self._labels = labels self._img_names = img_names
```

```
self._cls = cls self._epochs done = 0
self. index in epoch = 0
@property
def images(self):
return self._images
@property
def labels(self):
return self. labels
@property
def img_names(self):
return self. img names
@property
def cls(self):
return self._cls
@property
def num_examples(self):
return self. num examples
@property
def epochs_done(self):
return self. epochs done
def next_batch(self, batch_size):
"""Return the next `batch size` examples from this data set."""
start = self._index_in_epoch self._index_in_epoch += batch_size
if self. index in epoch >self. num examples:
# After each epoch we update this
self._epochs_done += 1
start = 0
self._index_in_epoch = batch_size
assert batch size <= self. num examplesend = self. index in epoch</pre>
return self._images[start:end], self._labels[start:end],
self. img names[start:end], self. cls[start:end]
def read_train_sets(train_path, image_size, classes, validation_size):
class DataSets(object):
pass
data sets = DataSets()
images, labels, img_names, cls = load_train(train_path, image_size,
classes)images, labels, img names, cls = shuffle(images, labels,
img names, cls)
if isinstance(validation size, float):
```

```
validation size = int(validation size * images.shape[0])
validation images = images[:validation size] validation labels =
labels[:validation_size] validation_img_names =
img names[:validation size]validation cls = cls[:validation size]
train images = images[validation size:] train labels =
labels[validation_size:] train_img_names = img_names[validation_size:]
train cls = cls[validation size:]
data sets.train = DataSet(train images, train labels, train img names,
train_cls)
data_sets.valid = DataSet(validation_images, validation_labels,
validation img names, validation cls)
return data_sets import tensorflow as tfimport numpy as np
from tkinter import *
import os
from tkinter import filedialog
import cv2
import time
from matplotlib import pyplot as plt
from tkinter import messagebox
def endprogram():
print ("\nProgram terminated!")sys.exit()
def training():
import Training as tr
def imgtraining():
import file path = filedialog.askopenfilename()
image = cv2.imread(import file path)filename = 'Test.jpg'
cv2.imwrite(filename, image)print("After saving image:")
gray = cv2.cvtColor(image, cv2.COLOR BGR2GRAY)
cv2.imshow('Original image', image)cv2.imshow('Gray image', gray)
# import_file_path = filedialog.askopenfilename()
```

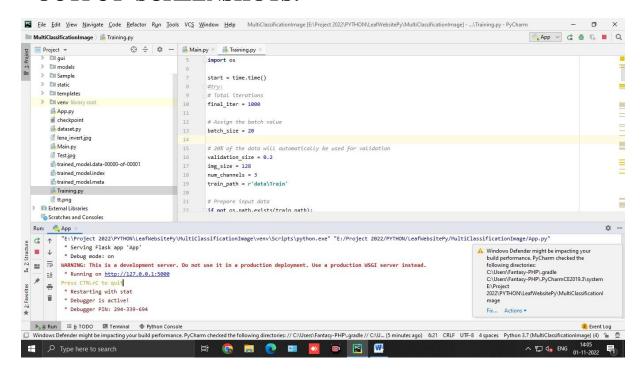
```
print(import file path)
fnm = os.path.basename(import file path)
print(os.path.basename(import file path))
from PIL import Image, ImageOps
im = Image.open(import_file_path)im_invert = ImageOps.invert(im)
im invert.save('lena_invert.jpg', quality=95)
im = Image.open(import_file_path).convert('RGB')im_invert =
ImageOps.invert(im) im invert.save('tt.png')
image2 = cv2.imread('tt.png')cv2.imshow("Invert", image2)
""""
img = image
gray = cv2.cvtColor(img, cv2.COLOR BGR2GRAY) cv2.imshow('Original image',
img)
#cv2.imshow('Gray image', gray)
dst = cv2.fastNlMeansDenoisingColored(img, None, 10, 10, 7, 21)
cv2.imshow("Nosie Removal", dst)
gray = cv2.cvtColor(img, cv2.COLOR BGR2GRAY)
print("\n**************\nImage : " + fnm +
"\n****************")img = cv2.imread(import_file_path)
if img is None:
print('no data')
img1 = cv2.imread(import file path)print(img.shape)
img = cv2.resize(img, ((int)(img.shape[1] / 5), (int)(img.shape[0] / 5)))
original = img.copy()
neworiginal = img.copy() cv2.imshow('original', img1)
gray = cv2.cvtColor(img1, cv2.COLOR_BGR2GRAY)
cv2.imshow('Original image', img1)
# cv2.imshow('Gray image', gray)
p = 0
for i in range(img.shape[0]):
for j in range(img.shape[1]):
B = img[i][j][0]
G = img[i][j][1]
R = img[i][j][2]
if (B >110 and G >110 and R >110):
totalpixels = img.shape[0] * img.shape[1]per white = 100 * p /
totalpixels
```

```
if per white >10:
img[i][j] = [500, 300, 200]
cv2.imshow('color change', img)
# Guassian blur
blur1 = cv2.GaussianBlur(img, (3, 3), 1)
# mean-shift algo
newimg = np.zeros((img.shape[0], img.shape[1], 3), np.uint8)
criteria = (cv2.TERM_CRITERIA_EPS + cv2.TERM_CRITERIA_MAX_ITER, 10, 1.0)
img = cv2.pyrMeanShiftFiltering(blur1, 20, 30, newimg, 0, criteria)
cv2.imshow('means shift image', img)
# Guassian blur
blur = cv2.GaussianBlur(img, (11, 11), 1)cv2.imshow('Noise Remove', blur)
corners = cv2.goodFeaturesToTrack(gray, 27, 0.01, 10)corners =
np.int0(corners)
# we iterate through each corner,
# making a circle at each point that we think is a corner.
for i in corners:
x, y = i.ravel()
cv2.circle(image, (x, y), 3, 255, -1)plt.imshow(image), plt.show()
def testing():
global testing_screen
testing_screen = Toplevel(main_screen)testing screen.title("Testing")
# login_screen.geometry("400x300")
testing screen.geometry("600x450+650+150") testing screen.minsize(120, 1)
testing_screen.maxsize(1604, 881)
testing_screen.resizable(1, 1)
# login screen.title("New Toplevel")
Label(testing_screen, text='''Upload Image''', background="#d9d9d9".
disabledforeground="#a3a3a3",
foreground="#000000", bg="turquoise", width="300", height="2",
font=("Calibri",16)).pack()
Label(testing_screen, text="").pack() Label(testing_screen,
text="").pack() Label(testing_screen, text="").pack()
Button(testing_screen, text='''Upload Image''', font=(
'Verdana', 15), height="2", width="30", command=imgtest).pack()
def imgtest():
import file path = filedialog.askopenfilename()
image = cv2.imread(import file path)print(import file path)
filename = 'data/alien test/Test.jpg'cv2.imwrite(filename, image)
print("After saving image:")
```

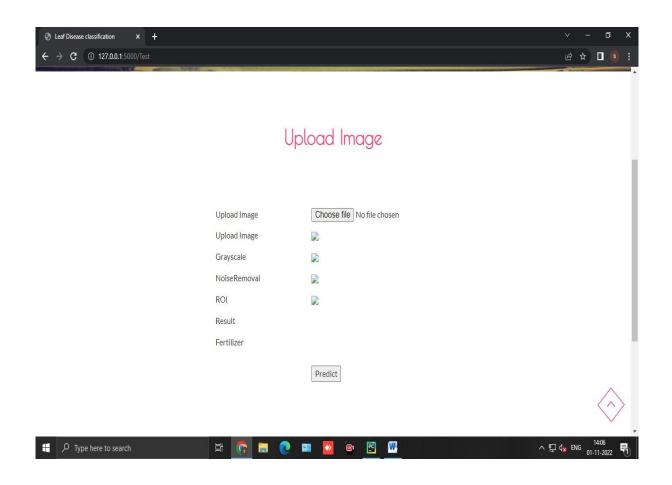
```
def main account screen():
from PIL import Image, ImageTk
global main_screen main_screen = Tk()width = 600
height = 600
screen_width = main_screen.winfo_screenwidth() screen_height =
main_screen.winfo_screenheight()x = (screen_width / 2) - (width / 2)
y = (screen_height / 2) - (height / 2) main_screen.geometry("%dx%d+%d+%d"
% (width, height, x, y)) main screen.resizable(0, 0)
# main_screen.geometry("300x250")
main screen.title("Leaf Disease classification")
Label(text="Leaf Disease classification", bg="turquoise", width="300",
height="5", font=("Calibri", 16)).pack()
Label(text="").pack()
Label(text="").pack()
image = ImageTk.PhotoImage(Image.open('gui/12344.jpg'))
Label(main screen, text='Hello', image=image, compound='left',
height="100", width="200",).pack()
Button(text="Training", font=(
'Verdana', 15), height="2", width="30", command=training,
highlightcolor="black").pack(side=TOP)
Label(text="").pack() Button(text="Testing", font=(
'Verdana', 15), height="2", width="30", command=testing).pack(side=TOP)
Label(text="").pack()
main screen.mainloop()
main account screen()
```

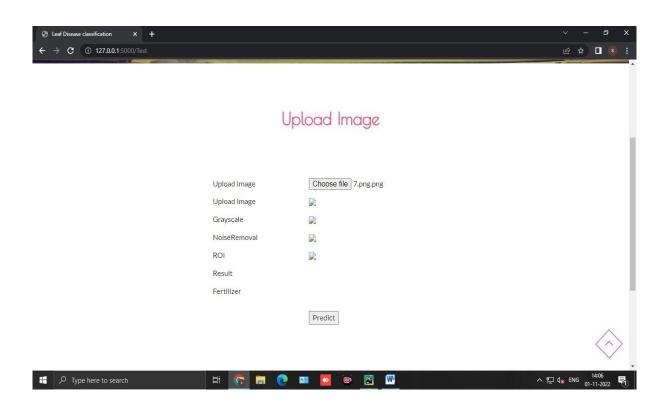
APPENDIX -2

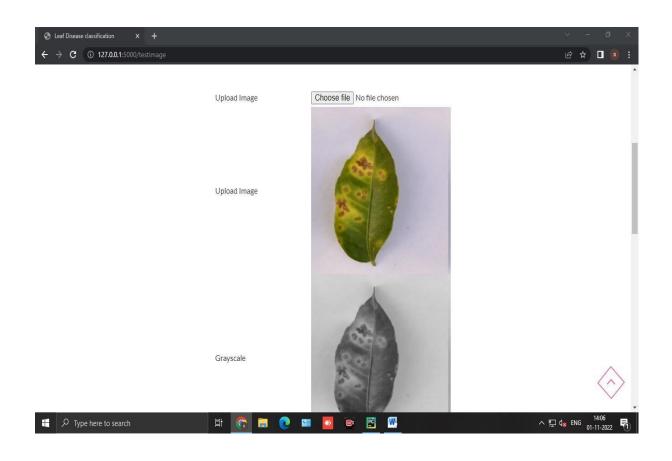
OUTPUT SCREENSHOTS:

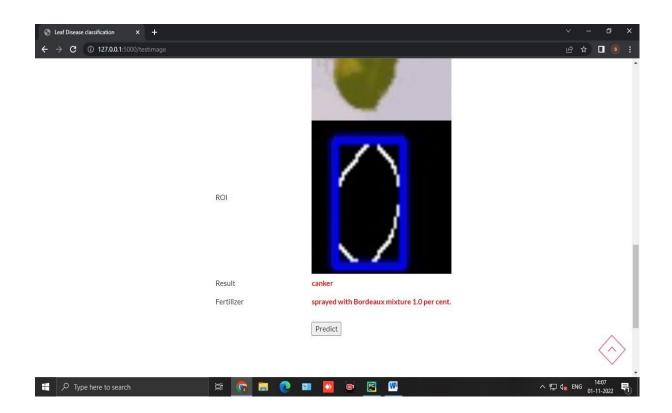












GITHUB:

https://github.com/IBM-EPBL/IBM-Project-28925-1660119014

DEMO VIDEO LINK:

https://drive.google.com/file/d/15g577gIuyUb0dPvop-337Y7pB64MEMUC/view?usp=share_link