

**SBM COLLEGE OF ENGINEERING
AND TECHNOLOGY**

**BACHELOR OF ENGINEERING
IN
ELECTRONICS AND
COMMUNICATION ENGINEERING.**

**FERTILIZER RECOMMENDATION
SYSTEM FOR DISEASE PREDICTION**

A PROJECT REPORT BY

P.NAGAJOTHI - 921619106041

M.BHARATHI PRIYA - 921619106009

T.DIVYADHARSHINI - 921619106018

S.YUVASHREE - 921619106067

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LITERATURE SURVEY

1) In Year 2022,"A Machine Learning Based New Recommendation System To The Farmer".

AUTHORS: D.N.V.S.L.S indra, M.Sobhana, A.H.L.Swaroop, V.PhaniKumar.

Totally 54% of India's land area is deemed arable, making it the world's largest agrarian economy. Soil infertility owing to over fertilization, as well as a lack of access and awareness of contemporary agricultural practices, are the different factors that contribute to low agricultural production. The main purpose of this research work is to develop a machine learning-based recommendation system to increase agricultural productivity. A variety of datasets were used in this study to design and develop advanced models to estimate the crop, recommend fertiliser, and identify plant disease. An algorithm called MobileNet uses an image of a leaf to identify the disease present in a plant. The XGBoost model predicts a suitable crop based on the local soil nutrients and rainfall. Random Forest [RF] model was used to propose fertilizer and develop ideas for improving soil fertility depending on nutrients present in the soil. When compared to other approaches, the proposed model delivers a high level of accuracy.

2) In Year 2022,"Crop Prediction And Disease Detection System".

AUTHORS: Sambhav Bhansali, Punit Shah, Jinay Shah, Priyal Vyas,Poonam Thakre.

Economy of India highly depends on agriculture. Still traditional ways of recommendations are used for agriculture. Currently, farmers use traditional ways of approximations for amount of fertilizer used and the type of crop to be sown. Agriculture extremely depends on the type of soil and climatic condition of the region. Therefore, it becomes vital to create advancement in this field. With the help of Machine Learning and Deep Learning Techniques we will create a Web-App which will be one-stop solutions for information regarding the agriculture. Crop and fertilizer recommendation system will help the farmers in increasing their yield production. We are going to take the soil parameters along with the weathers API to figure out the most suitable crop for that region. Using the decision tree and navies bayes algorithm we will make the recommendation model which will use the N-K-P, Ph. value.Basis on the crop and region of farming we will recommend the fertilizer and its uses to boost the yield productivity for farmers. Sometimes due to unwanted excess of rainfall or the pest attack can cause disease to crops. We will use the image classification technique where the user can upload the picture of the affected plant/crop and the system will figure out the type of disease which will be done using Support Vector Machine (SVM) or using the neural network techniques. And this disease detection will suggest that how that plant/crop can be cure or prevent. The aim is to make a common system for all the features and provide the results with the best accuracy for all the crops over most of the regions all over the India. Also, the price and news section will keep the farmers updated with daily market prices and government schemes and policies related to the agriculture and farming.

3) In Year 2022,"Agro-Farm-Crop,Fertilizer & Disease Prediction".

AUTHORS: SanidhyaPurohit, DeepSanghani, NamanSenjaliya, Prof.Anuradha Kapoor

Data mining is a rising studies area in crop yield analysis Yield prediciton is a complete essential problem in agriculture. Any farmer is interested in knowing how muchyield he is about to expect also, it will end-user-helpful to farmers for indicting which fertilizers to be used as well as knowing the crop diseases all at one place. The project comes with a model to be precise and accurate in predicting crop, fertilizers, Crop disease and deliver the end-user with proper recommendations about the required fertilizer ratio based on atmospheric and soil parameters of the land which enhance to increase the crop yield and increase farmer revenue.

4) In Year 2021,"Neural Network Based Fertilizers Recommendation System For Disorder Classification And Prediction In Petal Images".

AUTHORS: N.Valarmathi, M.Vengateshwaran, Kalaimani Shanmugam, R.Sudha.

The point of farming isn't just to take care of the ever-developing populace but at the same time is a basic wellspring of vitality and an answer for the emergency of an Earth-wide temperature boost. Determination of plant ailment is basic for early finding and control of it. The unaided eye method is generally utilized for the conclusion of ailments. This methodology requires experts who can recognize varieties in leaf shading. Ordinarily a similar malady is characterized by a few specialists as a different sickness. This arrangement is exorbitant, in light of the fact that it requires nonstop expert management. Makers need to follow their yields and perceive the primary signs at modest costs so as to abstain from spreading even a plant malady and spare a lot of income. Recruiting qualified ranchers can't be reasonable especially in far off geologically detached zones. AI calculations in an image can give a substitute strategy to following plants and an expert can deal with such a way to deal with offer their types of assistance at a lower cost. It incorporates picture division which incorporates the dynamic shape strategy and the picture arrangement approach which incorporates a neural system calculation for foreseeing various kinds of ailments. Or on the other hand grow the way to deal with suggest the composts dependent on the examination of power with estimations.

5) In Year 2020,"Design and Implementation of Fertilizer Recommendation System for Farmer".

AUTHORS: Dr.S.UshaKiruthikika, Dr.S.Kanaga Suba Raja,S.R. Ronak,S.Rengarajen, P.Ravindran.

India is an agrarian nation. But creating a profitable yield for the farmer in each crop cycle is becoming a major challenge on various factors. Picking the reasonable fertilizer for the land and yield is an important and basic part of agriculture. Deciding the supplement levels in soil utilizing lab hardware can be restrictively costly, particularly in developing nations. The current frameworks on deciding soil nutrient substance and proposal for fertilizer isn't sufficiently proficient efficient enough. This paper introduces a compelling technique for estimation of nutrient dimension in soil and suggestion for appropriate fertilizer. The proposed methodologies comprise of four stages: soil analysis, data pre-processing, data analysis and Recommendation. The soil sample is analyzed using an IoT based device utilizing NPK sensor with two electrodes are set to calculate collect the NPK ratio of the soil nutrient and for pre-processing, the data gathered from sensors are figured into correct dataset and machine learning algorithm is utilized to recognize the reasonable fertilizer. This venture is extremely valuable to farmer to pick the right fertilizer toward the start of product cycle and amplify the yield.

6) In Year 2019,"Fertilizers Recommendation System For Disease Prediction in Tree Leave".

AUTHORS: R.Neela , P.Nithya.

Agriculture is the main aspect of country development. Many people lead their life from agriculture field, which gives fully related to agricultural products. Plant disease, especially on leaves, is one of the major factors of reductions in both quality and quantity of the food crops. In agricultural aspects, if the plant is affected by leaf disease then it reduces the growth of the agricultural level. Finding the leaf disease is an important role of agriculture preservation. After pre-processing using a median filter, segmentation is done by Guided Active Contour method and finally, the leaf disease is identified by using Support Vector Machine. The disease-based similarity measure is used for fertilizer recommendation.

7) In Year 2018," Soil Toxicity Prediction and Recommendation System Using Data Mining in Precision Agriculture".

AUTHORS: Mayuri Pawar, Geetha Chillarage.

India is agricultural land. India ranks second worldwide in agriculture output, but GDP share is declining. There are many factors contribute for declining agriculture GDP which are inadequate irrigation, inadequate power supply, changing environmental conditions, conventional agricultural method etc. In this paper, the proposed system can help farmers by making them aware about their soil conditions. Farmers can maximize crops yield by knowing proportion of nutrients present in the soil. Soil toxicity

affects the soil nutrients which indirectly affects crops health. The proposed system predicts the level of toxicity present in the soil and makes farmer aware about it. Many farmers are depending on rainfall which is the one of the factor for poor growth and decreases crops yield. Thus the proposed system recommends the farmer about the crop, fertility of soil, level of toxicity and water supply. For this recommendation system, sensor's accuracy is very important as well as classification algorithm. For classification, decision tree J48 algorithm is used which is simple to implement and having more accuracy as compared with other classification algorithms. Issue of power supply can be overcome by using solar panel system.

8) In Year 2016,"Crop Recommendation System for Precision Agriculture".

AUTHORS:S.Pudumalar,E.Ramanujam,R.HarineRajashree,C.Kavya, T.Kiruthika,J.Nisha.

Data mining is the practice of examining and deriving purposeful information from the data. Data mining finds its application in various fields like finance, retail, medicine, agriculture etc. Data mining in agriculture is used for analyzing the various biotic and abiotic factors. Agriculture in India plays a predominant role in economy and employment. The common problem existing among the Indian farmers are they don't choose the right crop based on their soil requirements. Due to this they face a serious setback in productivity. This problem of the farmers has been addressed through precision agriculture. Precision agriculture is a modern farming technique that uses research data of soil characteristics, soil types, crop yield data collection and suggests the farmers the right crop based on their sitespecific parameters. This reduces the wrong choice on a crop and increase in productivity. In this paper, this problem is solved by proposing a recommendation system through an ensemble model with majority voiting technique using random tree.

Fertilizer Recomendation System

for Disease Prediction

PROBLEM STATEMENT:

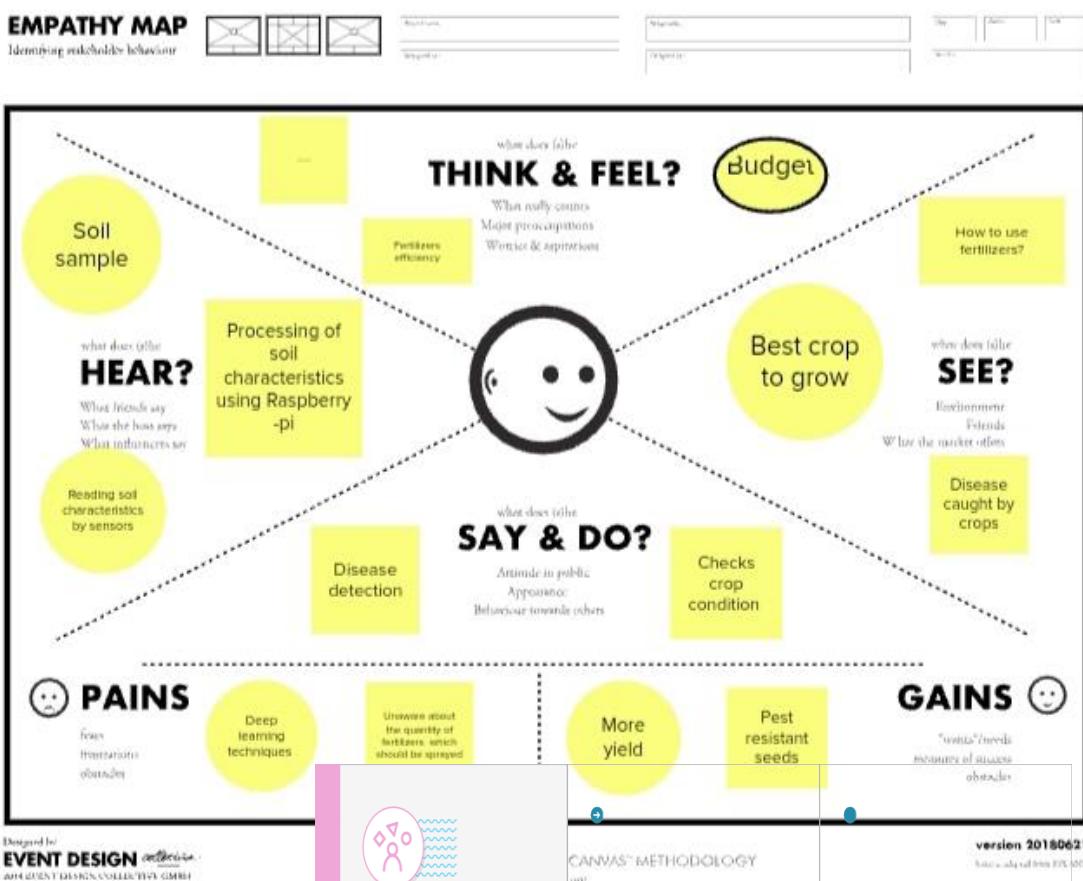
Mr.Narasimma Rao is a 65 years old man. He had a own farmingland and do Agriculture for past 30 Years , In this 30 Years he Faced a problem in Choosing Fertilizers and Controlling of Plant Disease.

- Narasimma Rao wants to know the better recommendation for fertilizers for plants with the disease.
 - He has faced huge losses for a long time.
 - This problem is usually faced by most farmers.
- Mr. Narasimma Rao needs to know the result immediately.

Who does the problem affect?	Persons who do Agriculture
What are the boundaries of the problem?	People who Grow Crops and facing Issues of Plant Disease
What is the issue?	In agricultural aspects, if the plant is affected by leaf disease, then it reduces the growth and productiveness. Generally, the plant diseases are caused by the abnormal physiological functionalities of plants.
When does the issue occur?	During the development of the crops as they will be affected by various diseases.

Where does the issue occur?	The issue occurs in agriculture practicing areas, particularly in rural regions.
Why is it important that we fix the problem?	It is required for the growth of better quality food products. It is important to maximise the crop yield.
What solution to solve this issue?	An automated system is introduced to identify different diseases on plants by checking the symptoms shown on the leaves of the plant.
What methodology used to solve the issue?	Deep learning techniques are used to identify the diseases and suggest the precautions that can be taken for those diseases.

Empathy Map

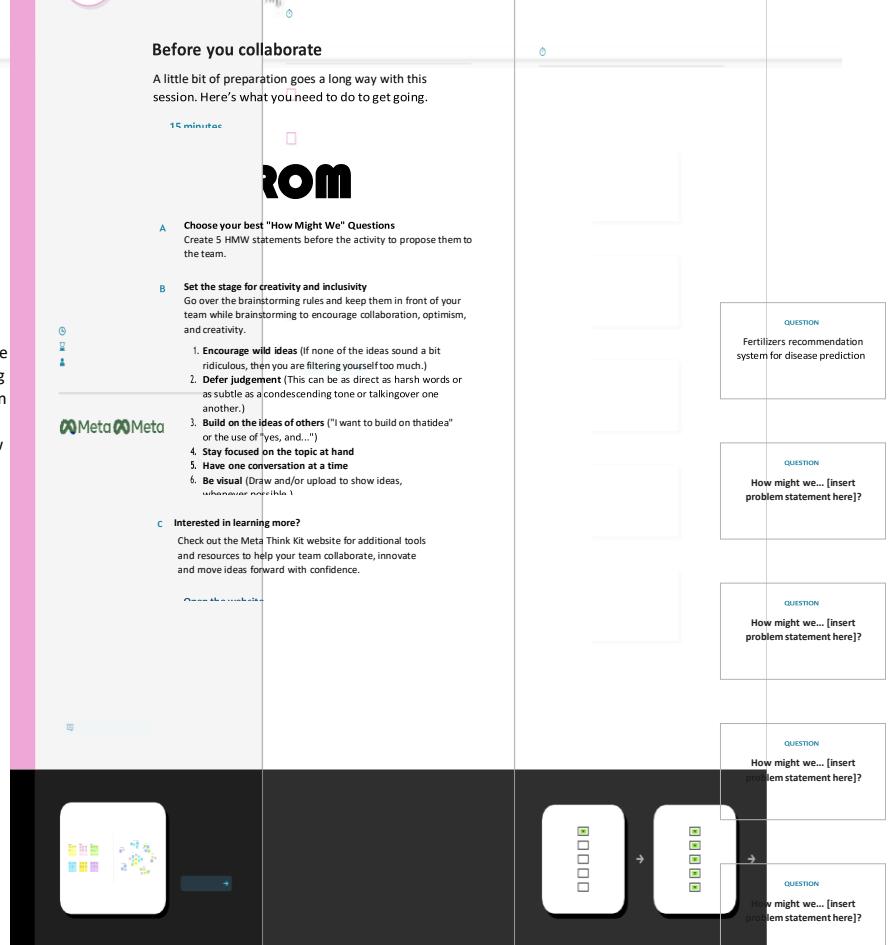


Conducting a brainstorm

Executing a brainstorm isn't unique; holding a productive brainstorm is. Great brainstorms are ones that settle the stage for fresh and generative thinking through simple guidelines and an open and collaborative environment. Use this when you're just kicking-off a new project and want to hit the ground running with big ideas that will move your business forward.

15 minutes to prepare
30-60 minutes to collaborate
3-8 people recommended

Created in partnership with



Project Design Phase-I

Proposed Solution

Template

Date	18 oct 2022
Team ID	PNT2022TMID49029
Project Name	Project – Fertilizer recommendation system for disease predication
Maximum Marks	2 Marks

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)	<p>Agriculture is having a great impact on the country's economy .</p> <p>Different diseases affect plant that reduces their production and is a major threat to food security.</p> <p>The major drawbacks that the farmers of our country are currently facing includes crop failure, lack of adequate knowledge, crop damage due to ignorance, lack of professional assistance, inaccessibility to agro-tech solutions. Most of the diseases are detected in later stages that to manually which is time consuming and results in heavy loss so it is important to build an automated system that detects disease at early stage and provides fertilizer recommendation accordingly.</p> <p>Farmer usually detect the crop diseases and plant disease with their naked eye which makes them take tough decisions on which fertilizer to use.</p> <p>It is necessary to develop crop yield prediction and fertilizer recommendation system which predicts crop yield based on soil nutrients crop yield data and recommend fertilizer for selected crop based on different dataset like fertilizer data, location data, and crop yield data.</p>

2.	Idea / Solution description	Implementation of artificial intelligence for identification of pests and recommendation of insecticides using TPF-CNN. The combination of two major things required in farming in one system is spraying proper insecticides and adding the needed fertilizer amount to the soil.
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		<p>Implementation of soil sensor for soil NPK nutrient analysis and recommendation of fertilizers accordingly.</p> <p>An automated system is built that takes the input as pictures of leaves which is uploaded by the user, identifies different disease on plants by checking the symptoms shown on the leaves of the plant.</p> <p>Deep learning techniques are used to identify the diseases and suggest the fertilizer needed for the plant.</p>
3.	Novelty / Uniqueness	<p>Efficient approach for controlling the overuse of insecticides and fertilizer in farming.</p> <p>Time efficient compared to KNN, SVM, and ANN.</p> <p>It can suggest and predict best and correct fertilizer for disease in the plant.</p> <p>It does not require user to consult any specialist for identification for disease that affected the leaves and the fertilizer that is required for the same.</p> <p>It detects plant disease at an early stage.</p>
4.	Social Impact / Customer Satisfaction	<p>Yield right crop at the right time, balancing the crop production, control plant disease, economic growth and planning to reduce the crop scarcity.</p> <p>Hence to detect and recognize the plant disease and to recommend fertilizer it is necessary to provide symptoms in identifying the disease at its earliest.</p> <p>Hence implemented new fertilizer recommendation system for crop disease prediction.</p> <p>The whole process of identifying disease and recommendation of fertilizer happens just by uploading image so it is user friendly.</p> <p>It helps farmers to get good yield out of the crop.</p> <p>People will get good quality food products.</p>
5.	Business Model (Revenue Model)	<p>Typically dedicate 10% of their AI investment to algorithms, 20% to technologies and 70% to embedding AI into business processes and agile ways of working.</p> <p>In other words, companies invest twice as much in people and processes as they do in technologies.</p> <p>Social media is the best way to spread the word about our application.</p> <p>And with influencers we can reach out to people.</p>

		Clustering and targeting the farmers for identifying disease on their plants and recommending them fertilizers for the same.
Cultivators Gardeners Plant	Scalability of the Solution	<p>This can be improved by introducing online purchases of crops, fertilizers, etc., easily.</p> <p>The farmers may not be aware of the infections or diseases that affected their plants.</p> <p>Even if they did, the nutrients required to cure may not be known. Identification of the right fertilizers and the quantity fertilizer that can be recommended for it among the list of fertilizer available.</p> <p>It can be used by anyone in the world.</p>

Project Title:

Project Design Phase-I - Solution Fit Template

Team ID:

PNT2022TMIDxxxxxx

Define CS, fit into CC

Explore AS, differentiate

1. CUSTOMER SEGMENT(S)	CS	6. CUSTOMER CONSTRAINTS	CC	5. AVAILABLE SOLUTIONS	AS
Who is your customer? i.e. working parents of 0-5 y.o. kids		What constraints prevent your customers from taking action or limit their choices of solutions? i.e. spending power, budget, no cash, network connection, available devices.		Which solutions are available to the customers when they face the problem or need to get the job done? What have they tried in the past? What pros & cons do these solutions have? i.e. pen and paper is an alternative to digital notetaking	

Fertilizers contain specific nutrients that are required for the proper development of the plant body. Some fertilizers soil fit plants indirectly by increasing water retention

Soil may not have adequate quantities of all nutrients. Rate of replenish of soil nutrients is much slower than the rate pf consumption. Hence fertilizers are required to balance these rates by providing enough nutrients to the soil and plants

An system that takes the image of leaves as input and identifies the different symptoms to decide on the disease that affects plant.

This will be done usinf Doop learning techniques.

Online portal is for accepting the input image and displaying the recommended fertilizers.

While offline, the image pre processing, segmentation, disease prediction etc are done

<p>Focus on J&P, tap into BE, understand RC</p>	<p>2. JOBS-TO-BE-DONE / PROBLEMS</p> <p>Which jobs-to-be-done (or problems) do you address for your customers? There could be more than one; explore different sides.</p>	J&P	<p>9. PROBLEM ROOT CAUSE</p> <p>What is the real reason that this problem exists? What is the back story behind the need to do this job? i.e. customers have to do it because of the change in regulations.</p>	RC	<p>7. BEHAVIOUR</p> <p>What does your customer do to address the problem and get the job done? i.e. directly related: find the right solar panel installer, calculate usage and benefits indirectly associated: customers spend free time on volunteering work (i.e. Greenpeace)</p>	BE
<p>Identify strong TR & EM</p>	<p>3. TRIGGERS</p> <p>What triggers customers to act? i.e. seeing their neighbour installing solar panels, reading about a more efficient solution in the news.</p>	TR	<p>10. YOUR SOLUTION</p> <p>If you are working on an existing business, write down your current solution first, fill in the canvas, and check how much it fits reality.</p> <p>If you are working on a new business proposition, then keep it blank until you fill in the canvas and come up with a solution that fits within customer limitations, solves a problem and matches customer behaviour.</p>	SL	<p>8. CHANNELS of BEHAVIOUR</p> <p>8.1 ONLINE What kind of actions do customers take online? Extract online channels from #7</p> <p>8.2 OFFLINE What kind of actions do customers take offline? Extract offline channels from #7 and use them for customer development.</p>	C
	<p>4. EMOTIONS: BEFORE / AFTER</p> <p>How do customers feel when they face a problem or a job and afterwards? i.e. lost, insecure > confident, in control - use it in your communication strategy & design.</p>	EM				

Project Design Phase-II

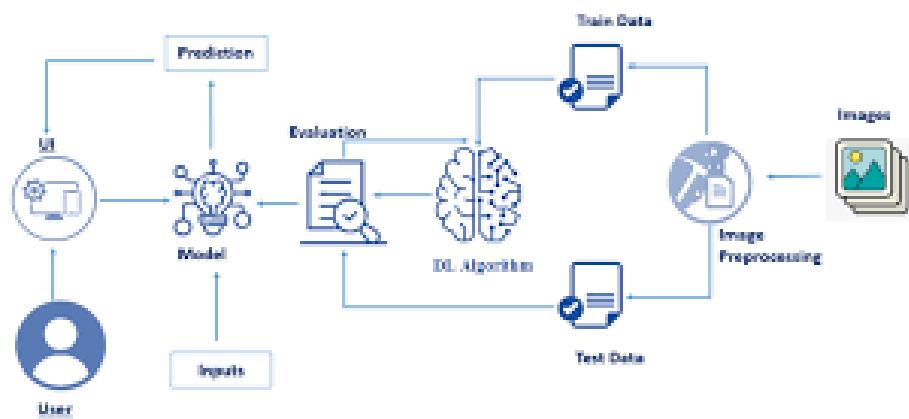
Data Flow Diagram & User Stories

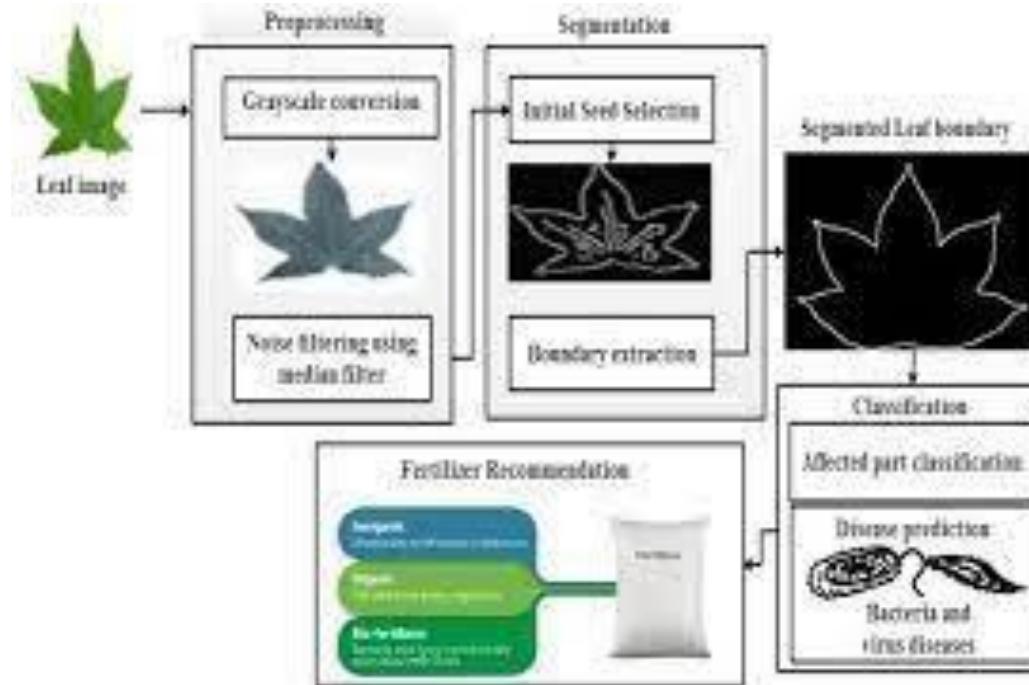
Date	20 October 2022
Team ID	PNT2022TMID49029
Project Name	Project – Fertilizer recommendation system for disease prediction
Maximum Marks	4 Marks

Data Flow Diagrams:

A Data Flow Diagram (DFD) is a traditional visual representation of the information flows within a system. A neat and clear DFD can depict the right amount of the system requirement graphically. It shows how data enters and leaves the system, what changes the information, and where data is stored.

Example:





User Stories

Use the below template to list all the user stories for the product.

Project Design Phase-II

Solution Requirements

(Functional & Non-functional)

Date	20 October 2022
Team ID	PNT2022TMID49029
Project Name	Project – Fertilizer recommendation system for disease prediction
Maximum Marks	4 Marks

Functional Requirements:

Following are the functional requirements of the proposed solution.

FR No.	Functional Requirement (Epic)	Sub Requirement (Story / Sub-Task)
FR-1	User Registration	<p>NAME: Enter Name</p> <p>EMAIL: Enter Mail</p> <p>PASSWORD: D: Enter Password</p> <p>PHONE: Enter Phone number (Registration through form Registration through gmail Registration through linked IN)</p>
FR-2	User Confirmation	<p>Thank you for your email</p> <p>We have received a request from your mail. Please confirm to proceed further.</p> <p>If any queries please contact our help centre to help get you an instant answer to your question. (Confirmation via Email Confirmation via OTP)</p>

FR-3	Product Features	<p>It provides data of the fertilizer to full fill the user's demands.</p> <p>Reading soil and plants characteristics by sensors.</p>
FR-4	Testing Features	<p>This estimation of nutrient in soil is done using an NPK monitoring unit with Arduino UNO as the microcontroller to read the values from it.</p> <p>Convolutional Neural Networks (CNN) algorithms recommend appropriate fertilizers that can be used to prevent damage to plants from pathogenic viruses.</p> <p>The fertilizer data is collected from various markets about the brand name and NPK ratio of the fertilizer is collected.</p>
FR-5	Objective	Smart farming and precision farming can be advanced by calculating NPK value for more accurate values.

		Analyzing the soil condition of any region and the requirements of the farmer to maximize the soil production.
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Non-functional Requirements:

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

FR No.	Non-Functional Requirement	Description
NFR-1	Usability	<p>It is very easily usable for the customer.</p> <p>The customer gets a notification whenever the insufficient nutrient or disease is detected in the plant.</p> <p>Help the farmer to identify the disease.</p>
NFR-2	Security	<p>Security is very much concerned regarding the data collected and customer details.</p> <p>These securities are mainly related to the cloud services, they have strict security across the network.</p> <p>The proposed method combines the two major aspects in farming, pest identification and insecticides recommendation.</p>
NFR-3	Reliability	<p>The use of artificial intelligence gives appropriate result.</p> <p>The CNN algorithm model has 95% accuracy.</p> <p>The reliability is more for the customers.</p> <p>It is easy to use so that health issues can be avoided.</p>
NFR-4	Performance	<p>The app runs on a mobile device under various loads and circumstances.</p> <p>Precision fertilizer and precision crops is mostly used.</p>

NFR-5	Availability	<p>There is a high availability for user's access. Anyone can make use of it.</p> <p>Reduces the losses as ammonia, nitrate leaching, apply the right rate and accurately.</p>
NFR-6	Scalability	<p>It is an effective way to minimize the damages for a plant by early detection of disease and recommending suitable fertilizers.</p> <p>If the soil is not replenished with nutrients through fertilizer, crop yield will deteriorate over time.</p>

Project design phase-II Technology stack(Architecture & stack)

Date	20 October 2022
Team id	PNT2022TMID49029
Project name	Fertilizer recommendation system for disease prediction
Maximum number	4 marks

Technical Architecture:

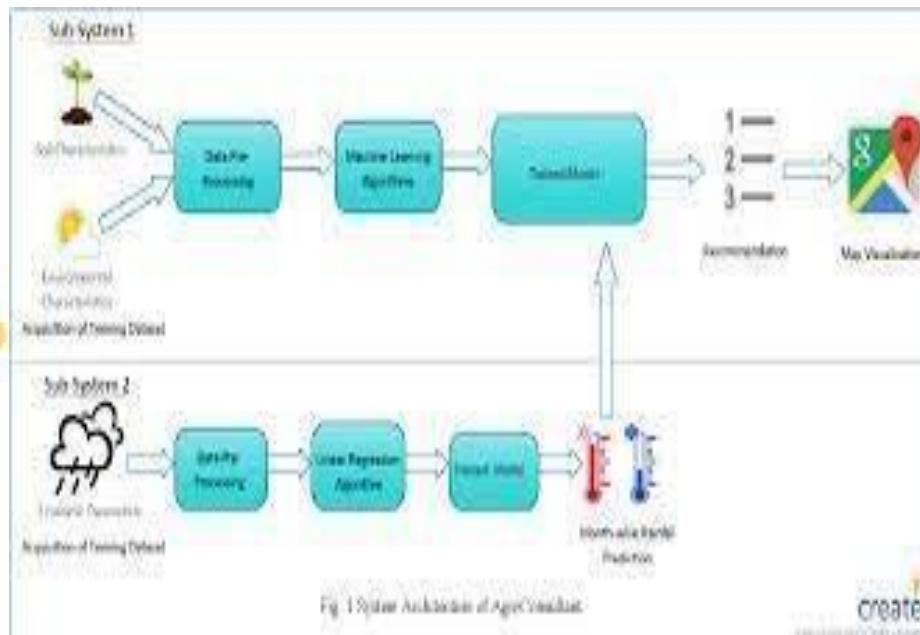


Table-1: Components & Technologies:

S.No	Component	Description	Technology
1.	User Interface	How user interacts with application e.g. Web UI, Mobile App, Chatbot etc.	HTML, CSS, JavaScript / Angular Js / React Js etc.
2.	Application Logic-1	Logic for a process in the application	Python
3.	Application Logic-2	Logic for a process in the application	IBM Watson STT service
4.	Application Logic-3	Logic for a process in the application	IBM Watson Assistant
5.	Database	Data Type, Configurations etc.	MySQL, NoSQL, etc.
6.	Cloud Database	Database Service on Cloud	IBM DB2, IBM Cloudant etc.
7.	File Storage	File storage requirements	IBM Block Storage or Other Storage Service or Local Filesystem
8.	External API-1	Purpose of External API used in the application	IBM Weather API, etc.
9.	External API-2	Purpose of External API used in the application	Aadhar API, etc.
10.	Machine Learning Model	Purpose of Machine Learning Model	Object Recognition Model, etc.
11.	Infrastructure (Server / Cloud)	Application Deployment on Local System / Cloud Local Server Configuration: Cloud Server Configuration :	Local, Cloud Foundry, Kubernetes, etc.

Table-2: Application Characteristics:

S.No	Characteristics	Description	Technology
1.	Open-Source Frameworks	List the open-source frameworks used	Anaconda Navigator, Tensor flow, Keras, Flask
2.	Security Implementations	List all the security / access controls implemented, use of firewalls etc.	e.g. SHA-256, Encryptions, IAM Controls, OWASP etc.
3.	Scalable Architecture	Justify the scalability of architecture (3 – tier, Micro-services)	Response time, Throughput, CPU and network usages, etc.
4.	Availability	Justify the availability of application (e.g. use of load balancers, distributed servers etc.)	All kind of users.
5.	Performance	Design consideration for the performance of the application (number of requests per sec, use of Cache, use of CDN's) etc.	Predicting disease, image processing, Visual similarity, rules, machine learning techniques, etc.

Project Planning Phase

Project Planning Template (Product Backlog, Sprint Planning, Stories, Story points)

Date	18 October 2022
Team ID	PNT2022TMID49029
Project Name	Project - Fertilizers Recommendation system for disease prediction
Maximum Marks	8 Marks

Product Backlog, Sprint Schedule, and Estimation (4 Marks)

Use the below template to create product backlog and sprint schedule

Sprint	Functional Requirement (Epic)	User Story Number	User Story / Task	Story Points	Priority	Team Members
Sprint-1	Registration	USN-1	As a user, I can sign up and register respective sites to access the required details and data. And import the required libraries for the processes.	2	High	P.Nagajothi M.Bharathi priya T.Divyadharshini S.Yuvashree
Sprint-2	Login	USN-2	As a user, I will access the page and test and train the CNN model to predict or detect the plant disease.	2	High	P.Nagajothi M.Bharathi priya T.Divyadharshini S.Yuvashree
Sprint-3	Customer Service	USN-3	As a customer care executive , I am available to the customers . so if the customers have any issues or in need of any assistance they will get help and solve them.	1	Medium	P.Nagajothi M.Bharathi priya T.Divyadharshini S.Yuvashree
Sprint-4	Dashboard	USN-4	As a user, I will have the access to know about the activities in the plant.	2	High	P.Nagajothi M.Bharathi priya T.Divyadharshini S.Yuvashree

Project Tracker, Velocity & Burndown Chart: (4 Marks)

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	6 Days	24 Oct 2022	29 Oct 2022	20	04 Nov 2022
Sprint-2	20	6 Days	31 Oct 2022	05 Nov 2022		06 Nov 2022
Sprint-3	20	6 Days	07 Nov 2022	12 Nov 2022		09 Nov 2022
Sprint-4	20	6 Days	14 Nov 2022	19 Nov 2022		12 Nov 2022

Velocity:

Imagine we have a 10-day sprint duration, and the velocity of the team is 20 (points per sprint). Let's calculate the team's average velocity (AV) per iteration unit (story points per day)

$$AV = \frac{\text{sprint duration}}{\text{velocity}} = \frac{20}{10} = 2$$

AV:

Sprint 1 = $20/6 = 3.33$,

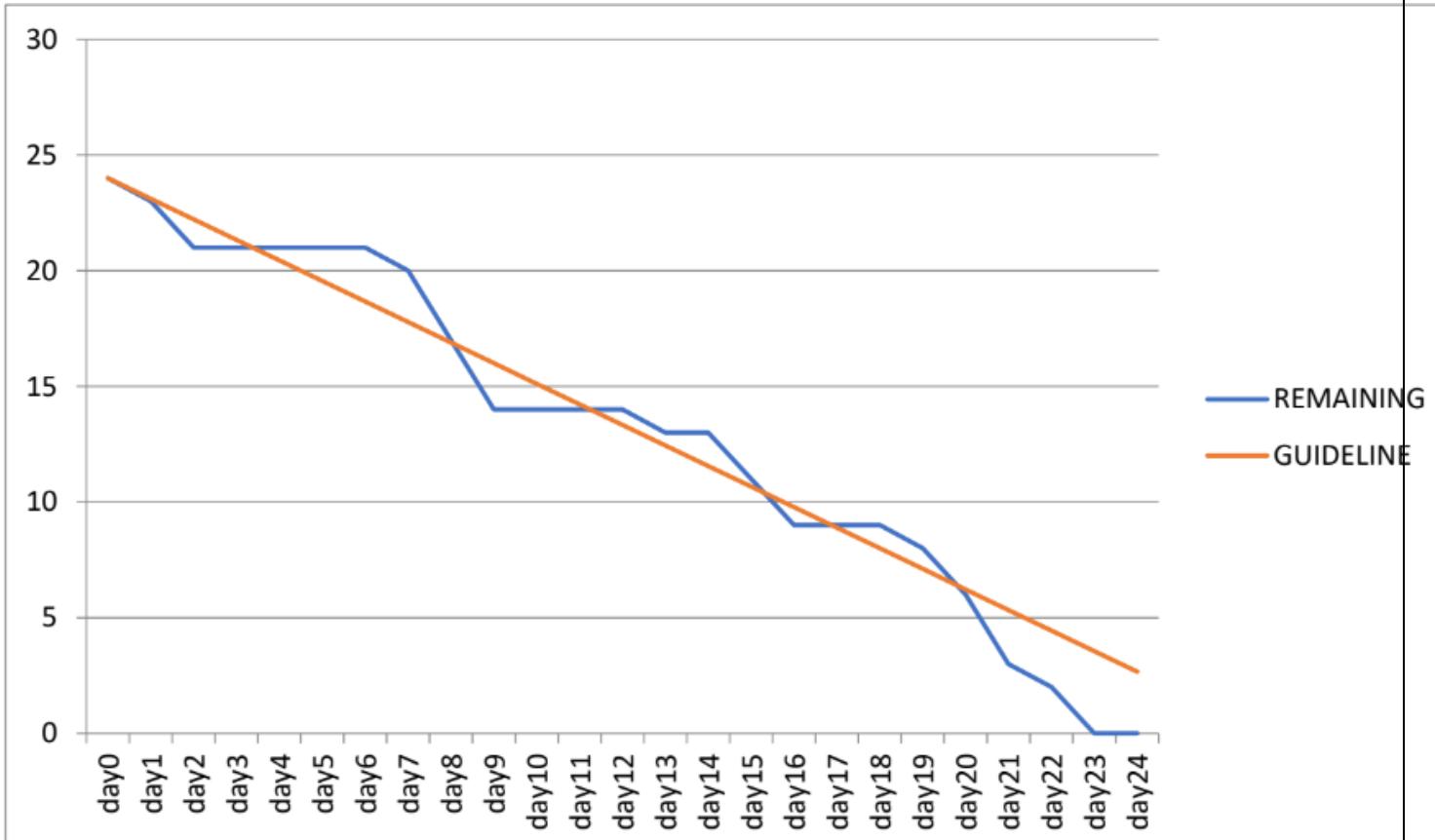
Sprint 2 = $20/6 = 3.33$,

Sprint 3 = $20/6 = 3.33$,

Sprint 4 = 20/6= 3.33.

Burndown Chart:

A burndown chart is a graphical representation of work left to do versus time. It is often used in agile software development methodologies such as Scrum. However, burn down charts can be applied to any project containing measurable progress over time.



Fertilizers Recommendation System for Disease Prediction

PROJECT REPORT

Submitted by

Team ID: PNT2022TMID49029

P.NAGAJOTHI

M.BHARATHIPRIYA

T.DIVYADHARSHINI

S.YUVASHREE

In partial fulfilment for the award of the

degree Of

BACHELOR OF ENGINEERING

In

ELECTRONICS AND COMMUNICATION ENGINEERING



SBM COLLEGE OF ENGINEERING AND TECHNOLOGY,DINDIGUL.

1. INTRODUCTION

1.1 Overview In this project, two datasets name fruit dataset and vegetable dataset are collected. The collected datasets are trained and tested with deep learning neural network named Convolutional Neural Networks (CNN). First, the fruit dataset is trained and then tested with CNN. It has 6 classes and all the classes are trained and tested. Second, the vegetable dataset is trained and tested. The software used for training and testing of datasets is Python. All the Python codes are first written in Jupyter notebook supplied along with Anaconda Python and then the codes are tested in IBM cloud. Finally, a web-based framework is designed with help of Flask a Python library. There are 2 html files are created in templates folder along with their associated files in static folder. The Python program 'app.py' used to interface with these two webpages is written in Spyder-Anaconda python and tested.

1.2 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.

2. LITERATURE SURVEY

2.1 Existing problem Indumathi 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. Pandi selvi proposed a simple prediction method for soil-based fertilizer recommendation system for predicted crop diseases. This method gives less accuracy and prediction. Shiva reddy proposed an IoT based system for leaf disease detection and fertilizer recommendation which is based on Machine Learning techniques yields less 80 percentage accuracies.

2.2 Proposed solution In this project work, a deep learning based neural network is used to train the collected datasets and test the same. The deep learning based

neuralnetwork is CNN which gives more than 90% classification accuracies. By
increasing

the more number of dense layers and by modifying hyperparameters such as number of epochs, batch size, the accuracy rate can be increased to 95% to 98%.

3. THEORITICAL ANALYSIS

3.1 Block diagram

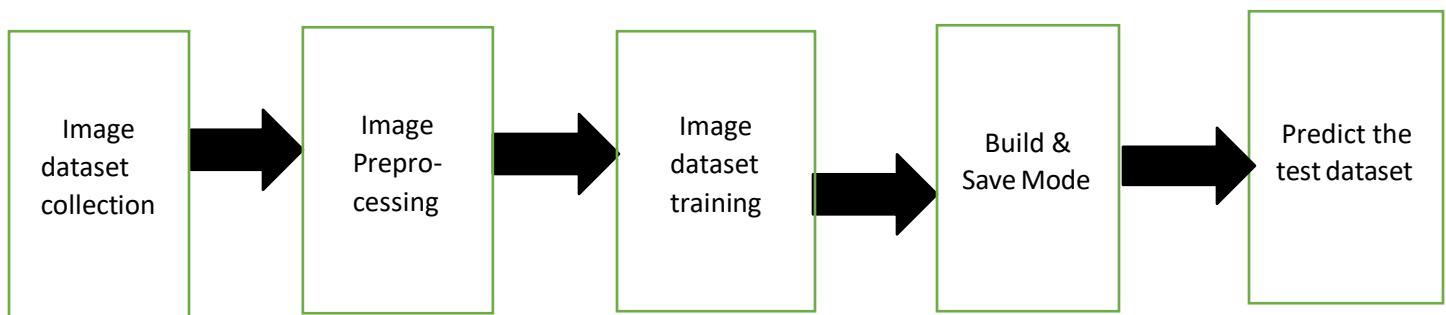


Figure 3.1. Block Diagram of the project

The block diagram of the entire project is shown in Fig.3.1. First step is the image dataset collection followed by image preprocessing. The third step is the training of image datasets with initializing different hyper parameters. Then build the model and save the model file with .h5 format. The final stage is the testing of existing or new datasets using the trained model.

3.2 Hardware/Software designing

The software used for training and testing the dataset is Python. The Jupyter notebook (Notebook of IBM cloud also) is used for python programming. The neural network used for training and testing the model is Convolutional Neural Network (CNN).

The CNN has following layers:

- Convolutional layer (32x32 kernel (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)

In the preprocessing step, images are normalized to 1 and then resized to 128x128. The images are arranged in different batch sizes. Then train set and test set are formed from the collected datasets. In order to do the above steps in Python, the following Python libraries must be imported before starting the process:

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

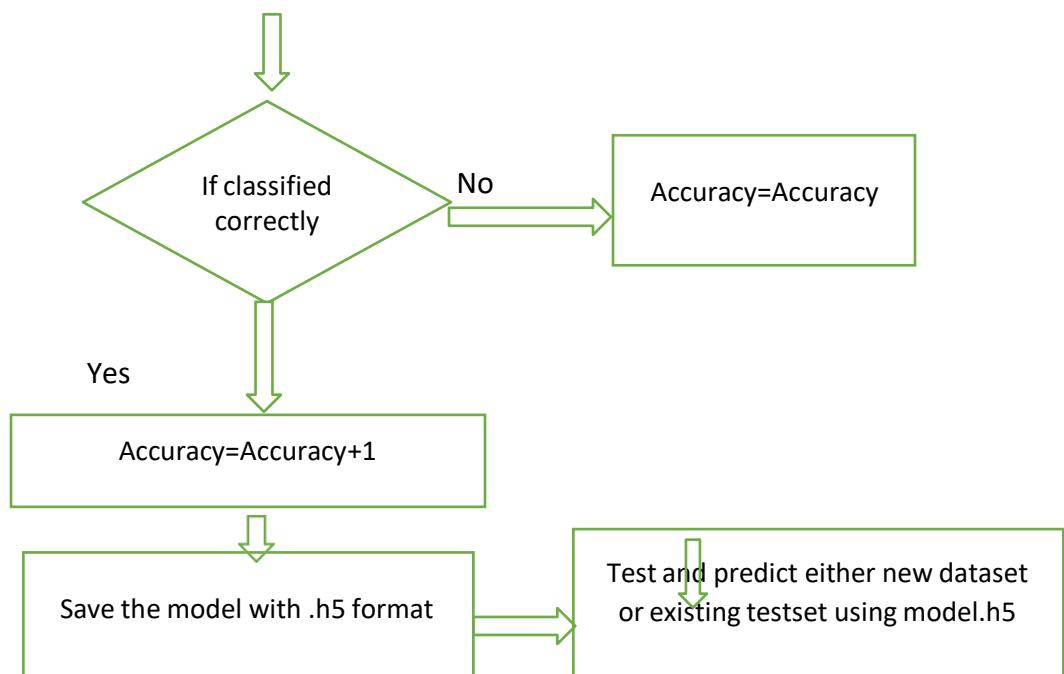
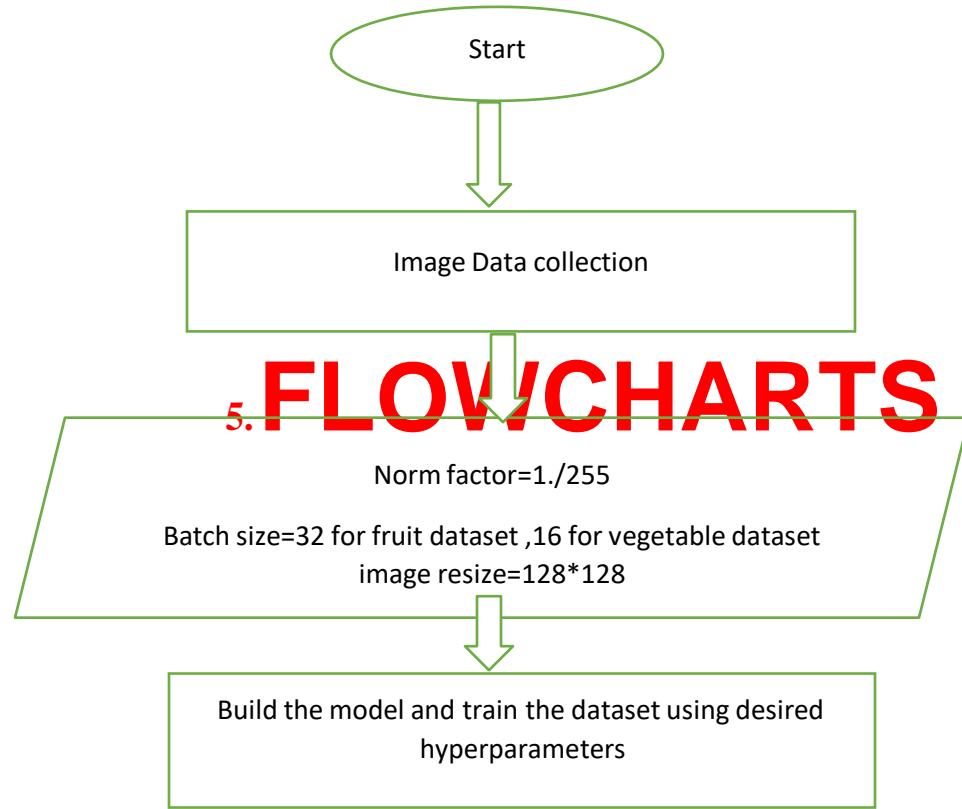
The following activation functions used in the CNN training:

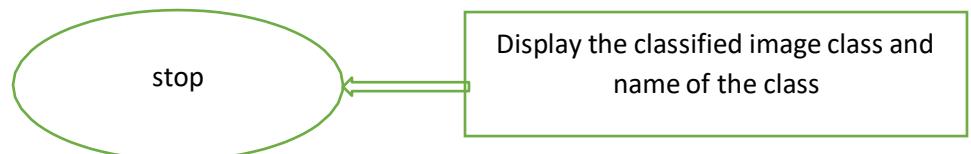
- RELU at the end of convolution layer and Max Pool layer
- SoftMax at the end of output dense layer
- For testing the dataset argmax is used, its an optional

4. EXPERIMENTAL INVESTIGATIONS

Analysis made while working on the solution The batch sizes are varied and tested. For different batch sizes, the CNN gives different accuracies. The batch size

determines the number of iterations per epoch. Another important hyper parameter is the number of epochs. This determines accuracy and it has high influence on accuracy compared to other hyper parameters. The accuracy can be varied from 80% to 90% in vegetable dataset and 95% to 98% in the case of fruit dataset by increasing the number of epochs. The size of test dataset and train dataset also has very high influence on accuracies. The accuracy can be increased by using more number of images in train dataset. The computational time for model building is increased when the size of the train dataset increased and also number of epochs increased. The batch size of train dataset and test dataset also play a vital role in computational time. The Neural Network complexity is increased when more number of convolutional layers increased. If the number of layers increased, better accuracy result will obtain. At the same time, increasing the number of layers in CNN leads to more training time and also requires more time to build a model. The model .h5 size depends on the size of train datasets. But the memory requirement depends on the size of train dataset and CNN architecture complexity.





6. RESULTS

Final findings(output) of the project given below in the form of screenshot:Training and Testing of Fruit dataset

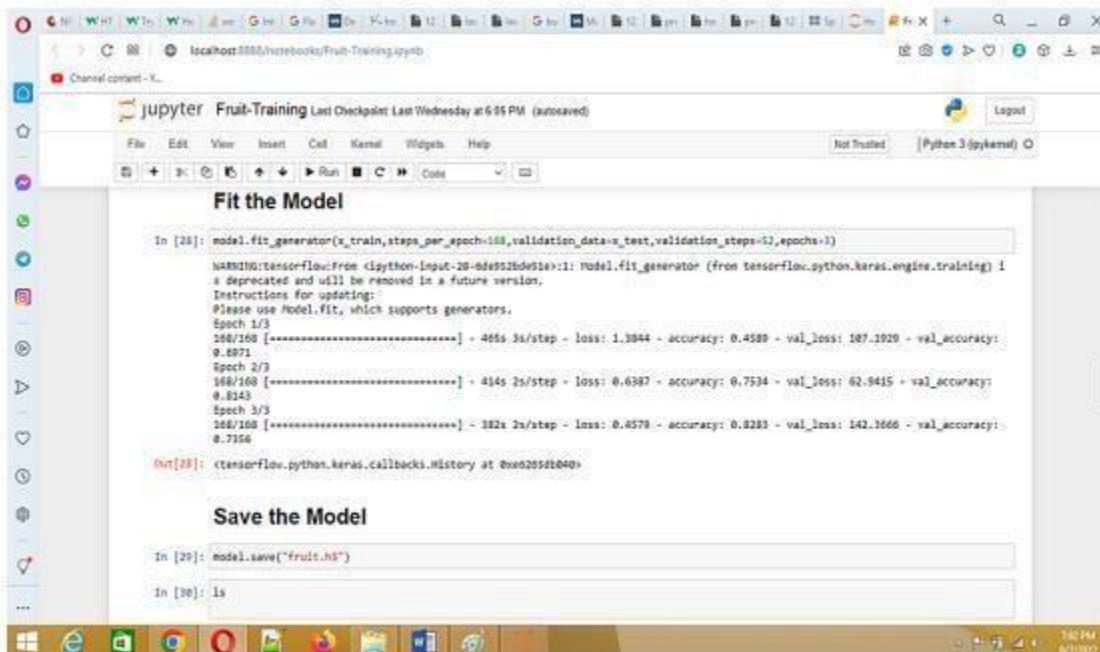


```
In [48]: pred = model.predict_classes(x)
WARNING:tensorflow:From <ipython-input-48-f93dd63985d>:1: Sequential.predict_classes (from tensorflow.python.keras.engine.sequential) is deprecated and will be removed after 2021-01-01.
Instructions for updating:
Please use instead: `np.argmax(model.predict(x), axis=-1)` , if your model does multi-class classification (e.g. if it uses a 'softmax' last-layer activation). `(model.predict(x) > 0.5).astype('int32')` , if your model does binary classification (e.g. if it uses a 'sigmoid' last-layer activation).
In [49]: pred
Out[49]: array([1], dtype=int64)
In [50]: index =['Apple__Black_rot','Apple__healthy','Corn_(maize)_Northern_Leaf_Blight','Corn_(maize)_healthy','Pear__Bacterial_Rust']
In [51]: print('the given image belongs to:',index[pred[0]])
the given image belongs to: Apple__Black_rot
```

Test Apple Black Rot class Images

```
In [52]: img = image.load_img("E:/IBM_HY_COURSE/Project/Dataset/Plant Disease/fruit-dataset/fruit-dataset/test/Apple__Black_rot/Bf3dd6d4.jpg")
In [53]: x=image.img_to_array(img)
```

Figure.6.2 Test the Fruit dataset



```
In [28]: model.fit_generator(x_train,steps_per_epoch=100,validation_data=x_test,validation_steps=50,epochs=1)
WARNING:tensorflow:From <ipython-input-28-0de952bd691a>: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 1/3
100/100 [=====] - 465s 3s/step - loss: 1.3844 - accuracy: 0.4589 - val_loss: 107.1929 - val_accuracy: 0.8971
Epoch 2/3
100/100 [=====] - 414s 2s/step - loss: 0.6387 - accuracy: 0.7534 - val_loss: 62.9435 - val_accuracy: 0.8143
Epoch 3/3
100/100 [=====] - 382s 2s/step - loss: 0.6579 - accuracy: 0.8283 - val_loss: 142.3666 - val_accuracy: 0.7156
Out[28]: <tensorflow.python.keras.callbacks.History at 0xe02054bb040>
```

Fit the Model

```
In [29]: model.save("fruit.h5")
In [30]: ls
```

Figure.6.1. Fit a model for Fruit dataset

Train and Test Vegetable dataset

```
In [27]: model.fit_generator(x_train,steps_per_epoch=80,validation_data=x_test,validation_steps=27,epochs=27)
WARNING:tensorflow:From tensorflow.python.keras.layers.core.py:110: 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
00:09 [=====] - 223s 3s/step - loss: 2.0686 - accuracy: 0.2193 - val_loss: 199.3698 - val_accuracy: 0.
1472
Epoch 2/20
00:09 [=====] - 1884.2s/step - loss: 1.6882 - accuracy: 0.3792 - val_loss: 248.8022 - val_accuracy: 0.
2708
Epoch 3/20
00:09 [=====] - 197s 2s/step - loss: 1.4388 - accuracy: 0.4521 - val_loss: 169.9319 - val_accuracy: 0.
3819
Epoch 4/20
00:09 [=====] - 192s 2s/step - loss: 1.2858 - accuracy: 0.5323 - val_loss: 278.2396 - val_accuracy: 0.
4867
Epoch 5/20
00:09 [=====] - 208s 2s/step - loss: 1.1856 - accuracy: 0.5688 - val_loss: 267.2704 - val_accuracy: 0.
4213
Epoch 6/20
00:09 [=====] - 233s 3s/step - loss: 1.0660 - accuracy: 0.6128 - val_loss: 308.8078 - val_accuracy: 0.
5148
```

Figure.6.3. Train the Vegetable dataset

```
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 [39]: img = Image.open('E:/IBM_DM_COURSE/Project/Dataset/Plant Disease/veg-dataset/veg-dataset/test_set/Pepper_bell__Bacterial_spot/1.jpg')
          x=np.expand_dims(x,axis=0)
          pred = model.predict_classes(x)
          print('the given image belongs to ',index[pred[0]])

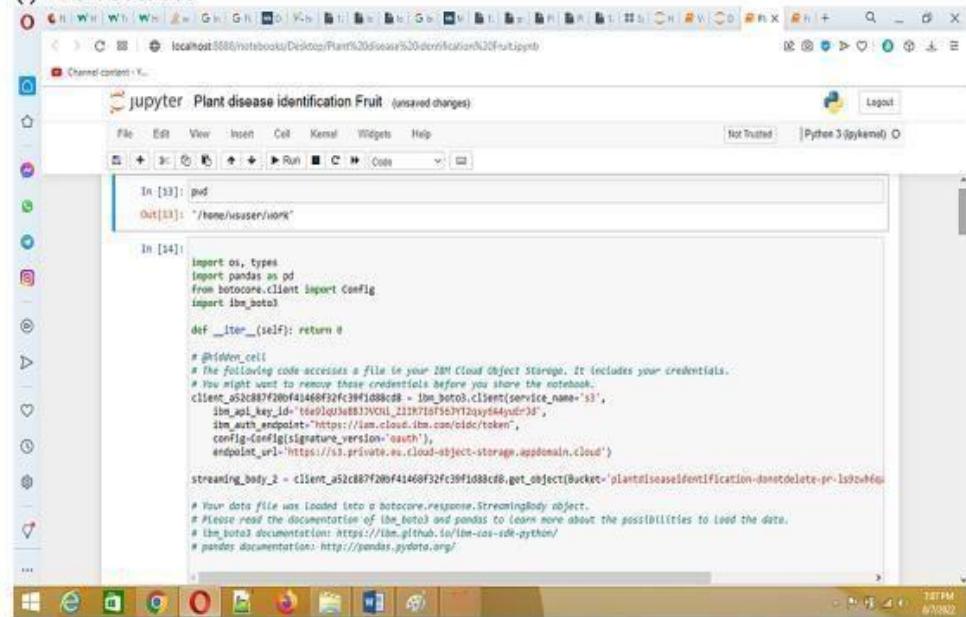
the given image belongs to Pepper_bell__Bacterial_spot
```

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

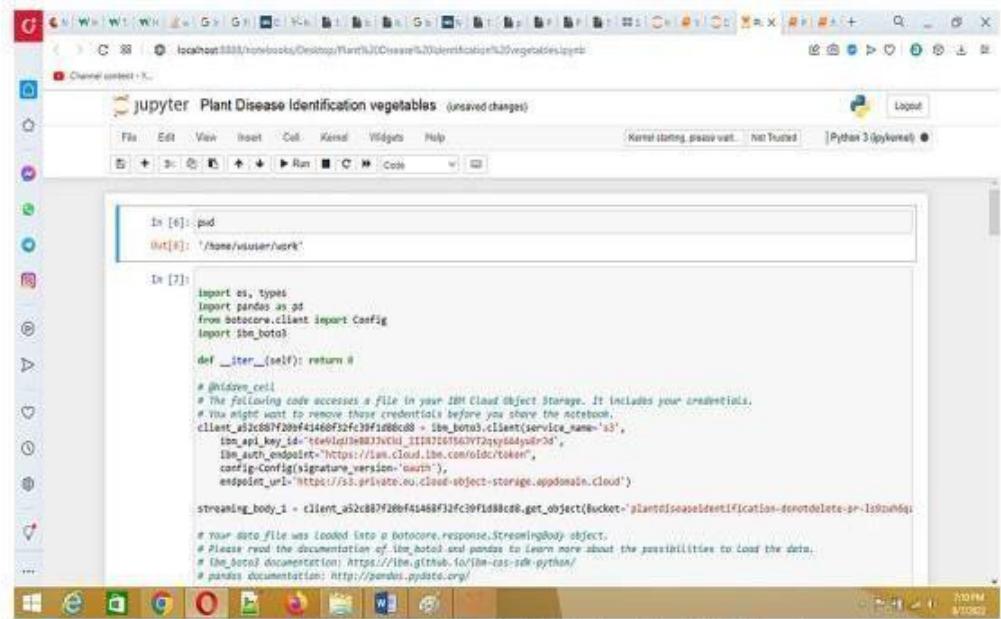
(i). Fruit dataset



The screenshot shows a Jupyter Notebook interface with the title "jupyter Plant disease identification Fruit (untrusted)". The notebook has two cells:

- In [13]:** `pd.read_csv('plant/disease/fruit.csv')`
- In [14]:** A longer block of Python code for interacting with IBM Cloud Object Storage to download a file named "fruit.csv". The code includes imports for os, types, pandas, boto3, Config, and ibm_boto3. It defines a generator function `_iter_(self)` and uses `StreamingBody` to handle the download. A note at the top of the code block says: "# 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." The code also specifies service name 's3', endpoint 'http://objectstorage.us-south.cloud.bluemix.net:19998', auth endpoint 'https://iam.cloud.ibm.com/oidc/token', config signature version 'v4', and endpoint url 'https://s3.private.us.cloud-object-storage.appdomain.cloud'.

Figure 6.5. Training Fruit Dataset in IBM Cloud

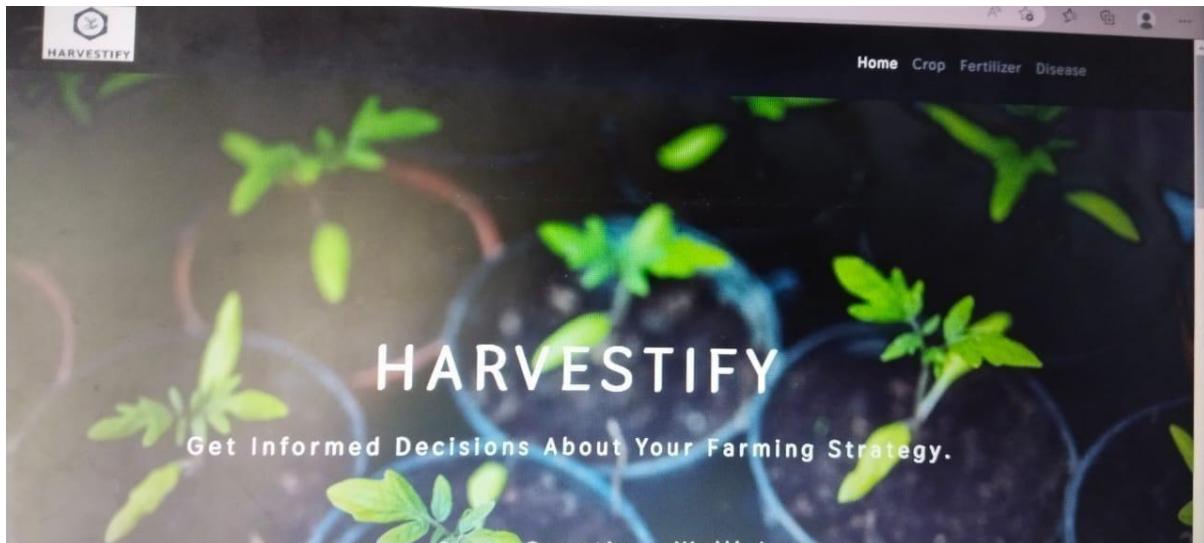


The screenshot shows a Jupyter Notebook interface in a web browser. The title bar says "Jupyter Plant Disease Identification vegetables (unsaved changes)". The notebook has two cells:

```
In [6]: pid  
Out[8]: '/home/visioner/work'  
  
In [7]:  
import es_types  
import pandas as pd  
from botocore.client import Config  
import ibm_boto3  
  
def __iter__(self): return it  
  
# This 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 = ibm_boto3.client(service_name='s3',  
    ibm_api_key_id='8be1d13e837cfd_212875054072a55a49f3d',  
    ibm_auth_endpoint='https://iam.cloud.ibm.com/oidc/token',  
    config=Config(signature_version='oauth'),  
    endpoint_url='https://s3.private.eu.cloud-object-storage.appdomain.cloud')  
  
streaming_body_1 = client._make_request('GET', 's3://200ff414408f32fc39f1a080d8 + ibm_boto3.client(service_name='s3',  
    ibm_api_key_id='8be1d13e837cfd_212875054072a55a49f3d',  
    ibm_auth_endpoint='https://iam.cloud.ibm.com/oidc/token',  
    config=Config(signature_version='oauth'),  
    endpoint_url='https://s3.private.eu.cloud-object-storage.appdomain.cloud')  
  
# 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://github.com/IBM/ibm-boto3-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)

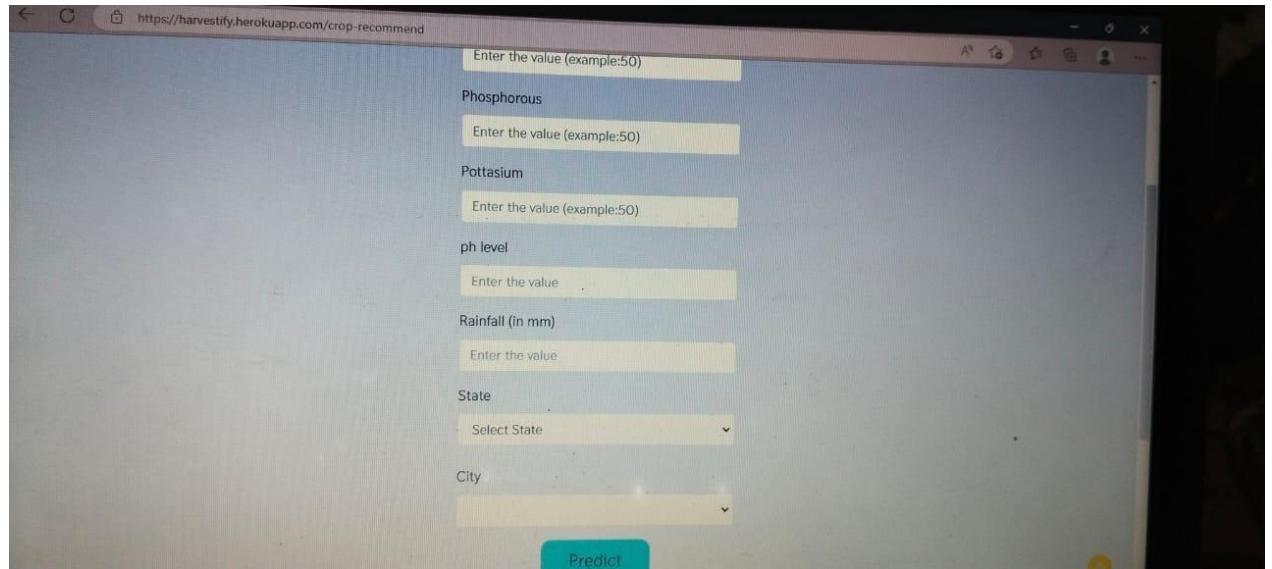
Enter the value

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)

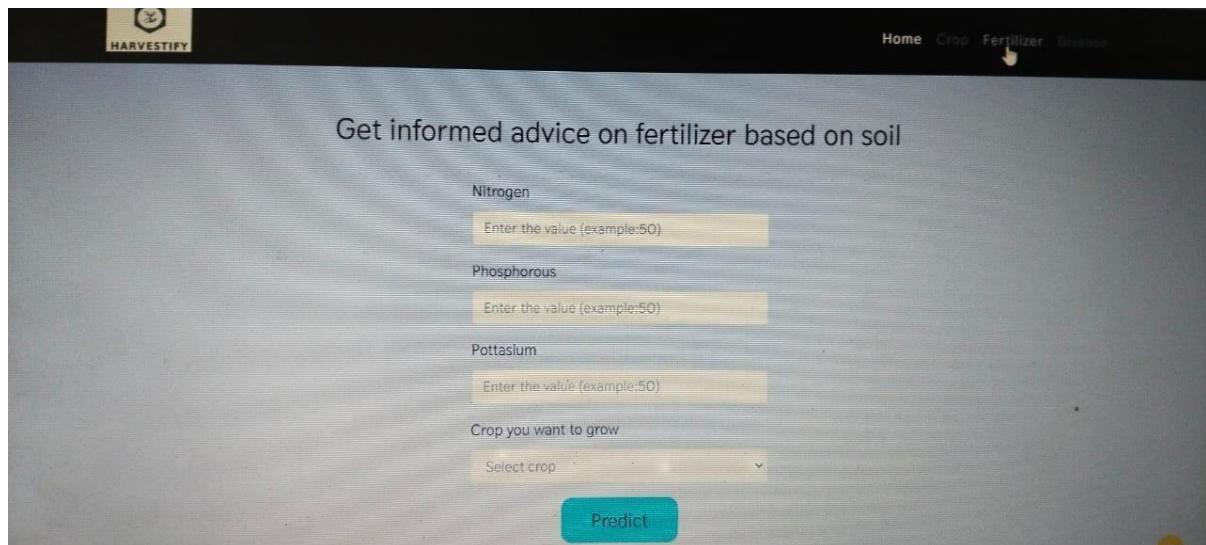
Pottasium

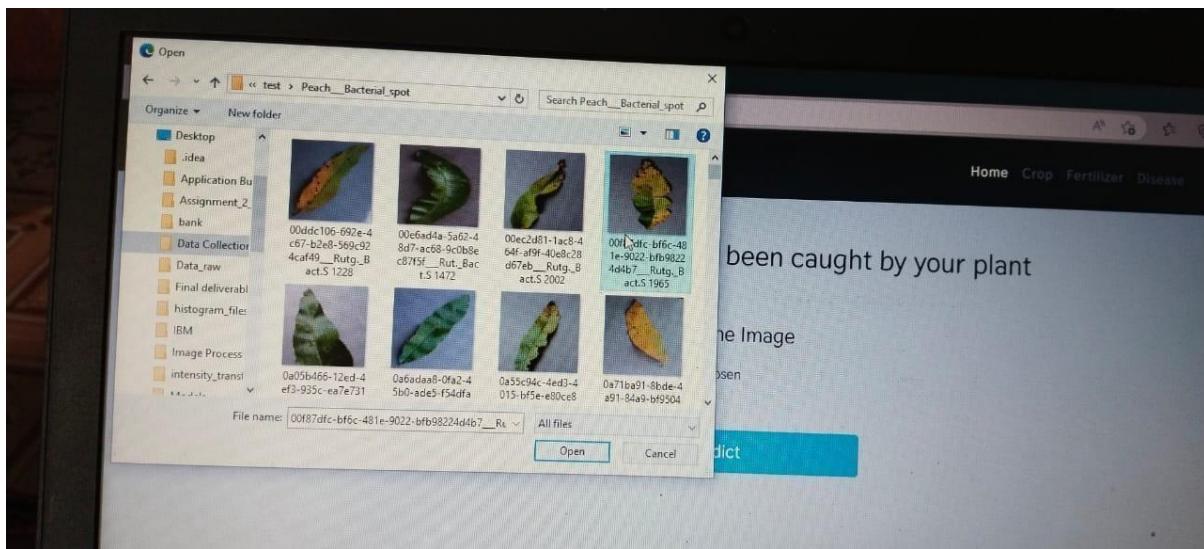
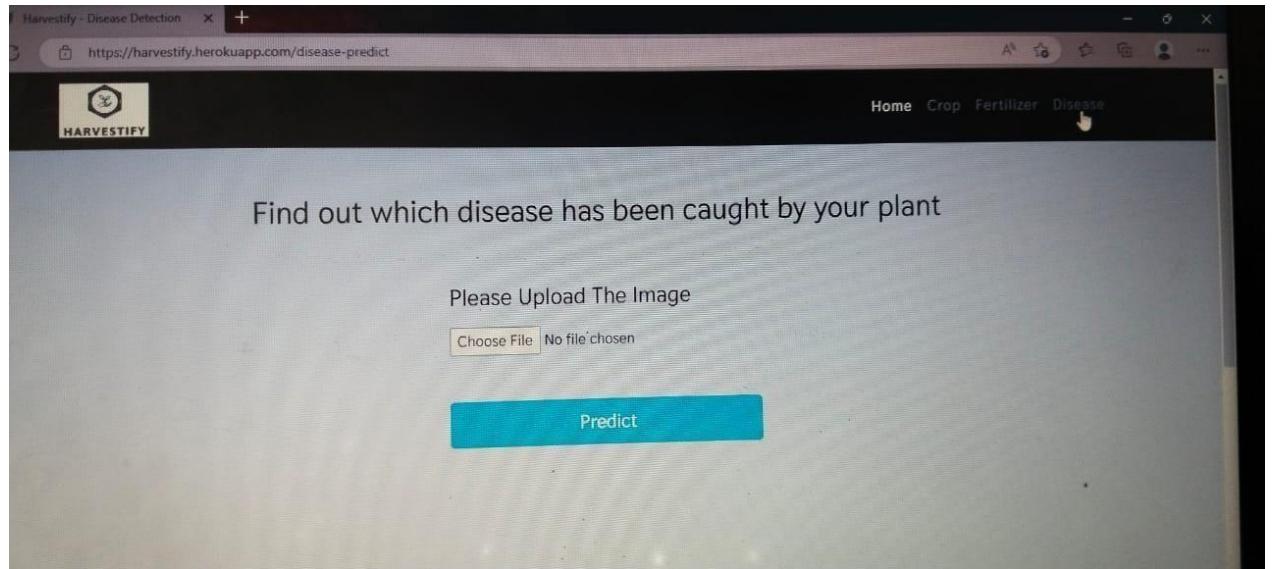
Enter the value (example:50)

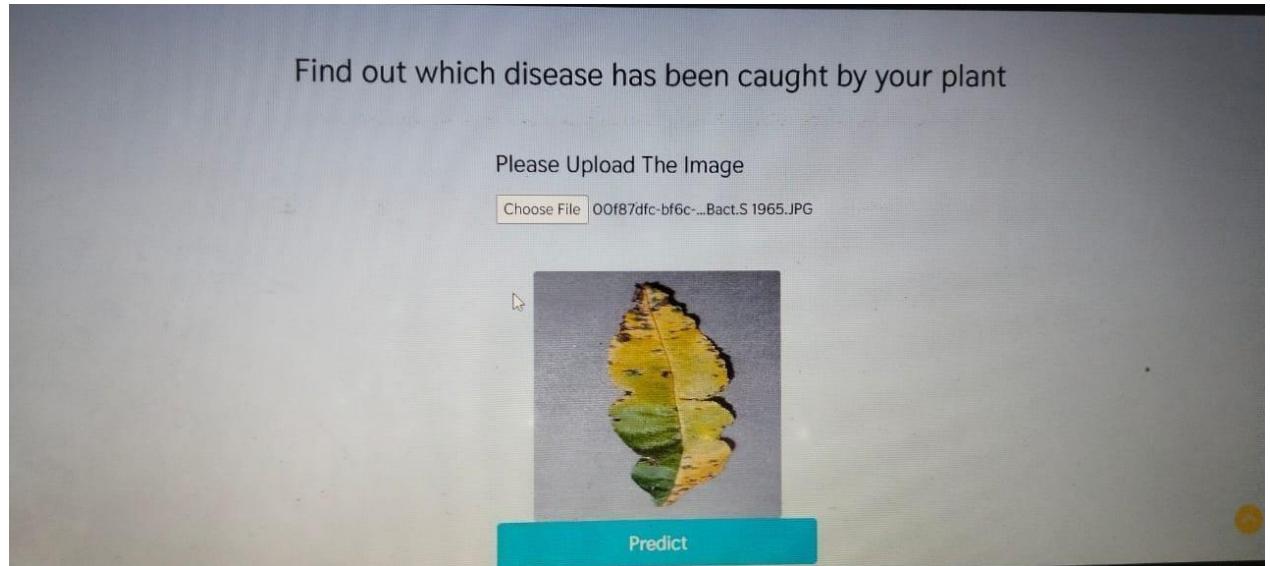
Crop you want to grow

Select crop

Predict







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.

How to prevent/cure the disease

1. The most effective management strategy is the use of pathogen-free certified seeds and disease-free transplants to prevent the introduction of the pathogen into greenhouses and field production areas. Inspect plants very carefully and reject infected transplants- including your own!
2. In transplant production greenhouses, minimize overwatering and handling of seedlings when they are wet.
3. Trays, benches, tools, and greenhouse structures should be washed and sanitized between seedlings crops.
4. Do not spray, tie, harvest, or handle wet plants as that can spread the disease.

7. ADVANTAGES & DISADVANTAGES

List of advantages

- The proposed model here produces very high accuracy of classification.
 - Very large datasets can also be trained and tested.
 - Images of very high can be resized within the proposed itself.

List of disadvantages

- For training and testing, the proposed model requires very high computational time.
- The neural network architecture used in this project work has high complexity.

8. APPLICATIONS

1. The trained network model used to classify the image patterns with high accuracy.
2. The proposed model not only used for plant disease classification but also for other image pattern classification such as animal classification.

3. This project work application involves not only image classification but also for pattern recognition.

9. CONCLUSIONS

The model proposed here involves image classification of fruit datasets and vegetable datasets. The following points are observed during model testing and training:

- The accuracy of classification increased by increasing the number of epochs.
- For different batch sizes, different classification accuracies are obtained.
 - The accuracies are increased by increasing more convolution layers.
 - The accuracy of classification also increased by varying dense layers.
- Different accuracies are obtained by varying the size of kernel used in the convolution layer output.
- Accuracies are different while varying the size of the train and test datasets.

10. FUTURE SCOPE

The proposed model in this project work can be extended to image recognition. The entire model can be converted to application software using python to exe software. The real time image classification, image recognition and video processing are possible with help OpenCV python library. This project work can be extended for security applications such as figure print recognition, iris recognition and face recognition.

11. BIBILOGRAPHY

- [1]. R Indumathi Leaf Disease Detection and 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.

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='categorical')  
  
# 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=['accuracy' ]) # 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,validation_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 PlantDisease/fruitdataset/fruit-dataset/test/Apple
                               _____healthy/00fc0da-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_Blight','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
                           PlantDisease/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 PlantDisease/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]])

```

FINAL CODE:

Index.html:

```

<!DOCTYPE html>
<html lang="en">
  <head>
    <!-- basic -->
    <meta charset="utf-8">

    <meta http-equiv="X-UA-Compatible" content="IE=edge">
    <!-- mobile metas -->

    <meta name="viewport" content="width=device-width, initial-scale=1">

    <meta name="viewport" content="initial-scale=1, maximum-scale=1">
    <!-- site metas -->

    <title>IBM</title>
    <meta name="keywords" content="">
    <meta name="description" content="">
    <meta name="author" content="">

    <!-- favicon -->

<link rel="icon" href="C:\Users\uma25\project\flask\static\images\favicon.png" type="image/gif" />

    <!-- bootstrap css -->
<link rel="stylesheet" href="C:\Users\uma25\project\flask\static\css\bootstrap.min.css">
    <!-- style css -->

<link rel="stylesheet" href="C:\Users\uma25\project\flask\static\css\style.css">

    <!-- Responsive-->
<link rel="stylesheet" href="C:\Users\uma25\project\flask\static\css\responsive.css">

```

```
<!-- Scrollbar Custom CSS -->
<link rel="stylesheet" href="C:\Users\uma25\project\flask\static\css\jquery.mCustomScrollbar.min.css ">

<!-- Tweaks for older IEs-->

<link rel="stylesheet" href="https://netdna.bootstrapcdn.com/font-awesome/4.0.3/css/font-
awesome.css">

<link rel="stylesheet" href="https://cdnjs.cloudflare.com/ajax/libs/fancybox/2.1.5/jquery.fancybox.min.css"
media="screen">

<!--[if lt IE 9]>
<script src="https://oss.maxcdn.com/html5shiv/3.7.3/html5shiv.min.js"></script>
<script src="https://oss.maxcdn.com/respond/1.4.2/respond.min.js"></script><![endif]-
->
</head>

<!-- body -->
<body class="main-layout">

<!-- loader -->

<div class="loader_bg">
<div class="loader"></div>
</div>

<!-- end loader -->

<!-- header -->

<header>

<!-- header inner -->

<div class="header-top">
<div class="header">
```

```
<div class="container-fluid">

    <div class="row">
        <div class="col-xl-2 col-lg-4 col-md-4 col-sm-3 col logo_section">

            <div class="full">
                <div class="center-desk">

                    <div class="logo">

<a href="index.html"></a>

                    </div>
                </div>
            </div>
            </div>
        </div>
    <div class="col-xl-10 col-lg-8 col-md-8 col-sm-9">

        <div class="menu-area">
            <div class="limit-box">

                <nav class="main-menu ">

                    <ul class="menu-area-main">
                        <li class="active"><a href="index.html">Home</a> </li>
                        <li><a href="#about">About</a> </li>
                        <li><a href="predict.html">Predict</a> </li>
                    </ul>
                </nav>
            </div>
        </div>
        </div>
        </div>
    </div>
</div>
```

```
</div>

<!-- end header inner -->
<!-- end header -->

<section class="slider_section">
<div id="myCarousel" class="carousel slide" data-ride="carousel">

    <ol class="carousel-indicators">
        <li data-target="#myCarousel" data-slide-to="0" class="active"></li>
        <li data-target="#myCarousel" data-slide-to="1"></li>
    </ol>

    <div class="carousel-inner">

        <div class="carousel-item active">
            <div class="container-fluid padding_dd">

                <div class="carousel-caption">
                    <div class="row">

                        <div class="col-xl-6 col-lg-6 col-md-6 col-sm-12">
                            <div class="text-bg">


# Fertilizers Recommendation System For Disease Prediction In Plants



<form class="Vegetable">

</div>

</div>
<div class="col-xl-6 col-lg-6 col-md-6 col-sm-12">

<div class="images_box">

<figure></figure>

</div>

</div>
</div>
```

```
</div>

</div>
</div>

<div class="carousel-item">
<div class="container-fluid padding_dd">

<div class="carousel-caption">
<div class="row">

<div class="col-xl-6 col-lg-6 col-md-6 col-sm-12">
<div class="text-bg">

<h2 style="font-size: 35px;">It is used to predict diseases in plants and to suggest fertilizers which would be required to eradicate the disease and to fight off the disease in the future too.</h2>

</div>

</div>
<div class="col-xl-6 col-lg-6 col-md-6 col-sm-12">

<div class="images_box">

<figure></figure>

</div>

</div>
</div>

</div>
</div>

</div>

</div>

</div>

</div>

</div>

<a class="carousel-control-prev" href="#myCarousel" role="button" data-slide="prev">
```

```
<span class="carousel-control-prev-icon" aria-hidden="true"></span>
<span class="sr-only">Previous</span>
</a>
<a class="carousel-control-next" href="#myCarousel" role="button" data-slide="next">
<span class="carousel-control-next-icon" aria-hidden="true"></span>
<span class="sr-only">Next</span>
</a>
</div>
</section>
</div>
</header>
<!-- about -->
<div id="about" class="about">
<div class="container-fluid">
<div class="row">
<div class="col-xl-6 col-lg-6 col-md-6 col-sm-12">
<div class="about-box">
<h2>About us</h2>
```

<p>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. </p>

```
        </div>

        </div>
<div class="col-xl-6 col-lg-6 col-md-6 col-sm-12 padding_rl">

    <div class="about-box_img">

<figure></figure>

    </div>

    </div>
</div>

    </div>

    </div>

    </div>

    </div>

    <!-- end abouts -->
    <!-- end vegetable -->

    <!-- Javascript files-->

<script src="C:\Users\uma25\project\flask\static\js\jquery.min.js"></script>

<script src="C:\Users\uma25\project\flask\static\js\popper.min.js"></script>

<script src="C:\Users\uma25\project\flask\static\js\bootstrap.bundle.min.js"></script>

<script src="C:\Users\uma25\project\flask\static\js\jquery-3.0.0.min.js"></script>

    <script src="C:\Users\uma25\project\flask\static\js\plugin.js"></script>
        <!-- sidebar -->
<script src="C:\Users\uma25\project\flask\static\js\jquery.mCustomScrollbar.concat.min.js"></script>

    <script src="C:\Users\uma25\project\flask\static\js\custom.js"></script>
```

```
<script src="https:cdnjs.cloudflare.com/ajax/libs/fancybox/2.1.5/jquery.fancybox.min.js "></script>
</body>
</html>
```

Predict.html:

```
<!DOCTYPE html>
<html lang="en">

    <head>
        <!-- basic -->
        <meta charset="utf-8">
        <meta http-equiv="X-UA-Compatible" content="IE=edge">
        <!-- mobile metas -->
        <meta name="viewport" content="width=device-width, initial-scale=1">
        <meta name="viewport" content="initial-scale=1, maximum-scale=1">
        <!-- site metas -->
        <title>IBM</title>
        <meta name="keywords" content="">
        <meta name="description" content="">
        <meta name="author" content="">
        <!-- fevicon -->
        <link rel="icon" href="C:\Users\uma25\project\flask\static\images\fevicon.png" type="image/gif" />
        <!-- bootstrap css -->
        <link rel="stylesheet" href="C:\Users\uma25\project\flask\static\css\bootstrap.min.css">
```

```
<!-- style css -->
<link rel="stylesheet" href="C:\Users\uma25\project\flask\static\css\style.css">

<!-- Responsive-->
<link rel="stylesheet" href="C:\Users\uma25\project\flask\static\css\responsive.css">

<!-- Scrollbar Custom CSS -->

<link rel="stylesheet" href="C:\Users\uma25\project\flask\static\css\jquery.mCustomScrollbar.min.css ">

<!-- Tweaks for older IEs-->
<link rel="stylesheet" href="https://netdna.bootstrapcdn.com/font-awesome/4.0.3/css/font-awesome.css">

<link href="{{ url_for('static', filename='css/final.css') }}" rel="stylesheet">

<link rel="stylesheet" href="https://cdnjs.cloudflare.com/ajax/libs/fancybox/2.1.5/jquery.fancybox.min.css"
      media="screen">

<!--[if lt IE 9]>
<script src="https://oss.maxcdn.com/html5shiv/3.7.3/html5shiv.min.js"></script>

<script src="https://oss.maxcdn.com/respond/1.4.2/respond.min.js"></script><![endif]-
      ->
</head>

<body>

<!-- predict -->
<div id="predict" class="vegetable">

<div class="container">
  <div class="row">

    <div class="col-md-12">
      <div class="titlepage">
```

```

<h2> Plant Disease <strong class="llow">Prediction</strong> </h2>

</div>
</div>

</div>
<div class="row">

<div class="col-xl-5 col-lg-5 col-md-5 col-sm-12 ">

<h4 style="color: green;"><b>Drop the image to get theprediction!!!...</b></h4>

<div class="vegetable_shop">
<form action="" id="upload-file" method="post" enctype="multipart/form-data">

<h5 style="color: red;"><b>Choose Category</b></h5>

<select name="plant">

<option value="fruit" name="fruit">Fruits</option>
<option value="vegetable" name="vegetable">Vegetables</option>

</select>
<br><br>

<input id="imageUpload" type="file" name="image" value="CHOOSE" accept="image/jpeg, image/jpg, image/png" onchange="document.getElementById('output').src=window.URL.createObjectURL(this.files[0])">

<br><br>

<img src="" id="output">

<br><br>

<button type="button" class="btn btn-info btn-lg " id="btn-predict" style="background: #28272c;">Predict</button>

</form>
<!--drop down,file upload,button-->

</div>

```

```
</div>

<div class="loader" style="display:none;"></div>
<h3>

<span id="result" style="font-size:17px; "> </span>
</h3>

<div class="col-xl-7 col-lg-7 col-md-7 col-sm-12 ">
<div class="vegetable_img">

<right>

<figure></figure>
</right>

</div>
</div>

</div>
</div>

<!-- Javascript files-->

<script src="C:\Users\uma25\project\flask\static\js\jquery.min.js"></script>
<script src="C:\Users\uma25\project\flask\static\js\popper.min.js"></script>
<script src="C:\Users\uma25\project\flask\static\js\bootstrap.bundle.min.js"></script>

<script src="C:\Users\uma25\project\flask\static\js\jquery-3.0.0.min.js"></script>
<script src="C:\Users\uma25\project\flask\static\js\plugin.js"></script>
<script src="C:\Users\uma25\project\flask\static\js\main.js"></script>
<!-- sidebar -->

<script src="C:\Users\uma25\project\flask\static\js\jquery.mCustomScrollbar.concat.min.js"></script>
<script src="C:\Users\uma25\project\flask\static\js\custom.js"></script>
```

```
<script src="https:cdnjs.cloudflare.com/ajax/libs/fancybox/2.1.5/jquery.fancybox.min.js "></script>
</body>
</html>
```

App.py :

```
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__)
```

```
model=load_model('vegetable.h5')
model1=load_model('fruit.h5')
```

```
#homepage
@app.route('/')
def home():
    return render_template('index.html')
```

```
#prediction page
@app.route('/prediction')def
    prediction():

        return render_template('predict.html')

@app.route('/predict',methods=['POST'])def predict():

    if request.method=='POST': f=request.files['image']
    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(preds)print(preds)
        df=pd.read_excel('precautions_veg.xlsx')
        print(df.iloc[preds]['caution'])
    else:
        preds = model1.predict(x)
        preds=np.argmax(preds)
```

```
df=pd.read_excel('precautions_fruits.xlsx')print(df.iloc[preds]['caution'])

return df.iloc[preds]['caution']if __name__ == "__main__":
    app.run(debug=False)
```

Fertilizers

Reco

mendation

Syste

mFor Disease Prediction

Team ID :
PNT2022TMID49029

Project Objectives:

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

Project Flow :

- A web Application is built where;
- Farmers can 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
 - To accomplish the above task you must complete the below activities and tasks;

- Download the dataset.
- Classify the dataset into train and test sets.
 - Add the neural network layers.
- Load the trained images and fit the model.
 - Test the model.
- Save the model and its dependencies.
- Build a Web application using a flask that integrates with the model built.

Prior Knowledge :

- **Supervised and Unsupervised learning:**

- Supervised learning, as the name indicates, has the presence of a supervisor as a teacher. Basically supervised learning is when we teach or train the machine using data that is welllabelled. Which means some data is already tagged with the correct answer. After that, the machine is provided with a new set of data.so that the supervised learning algorithm analyses the training data and produces a correct outcome from labelled data.
- Unsupervised learning is the training of a machine using information that is neither classified nor labeled and allowing the algorithm to act on that information without guidance. Here the task of the machine is to group unsorted information according to similarities, patterns, and differences without any prior training of data.

- **Regression Classification and Clustering:**

- Clustering is an unsupervised technique. With clustering, the algorithm tries to find a pattern in data sets without labels associated with it. This could be a clustering of buying behaviour of customers. Features for this would be the household income, age, ... and clusters of different consumers could then be built.
- Classification algorithms look at existing data and predicts what a new data belongs to. Classification is used for spam for years now and these algorithms are more or less maturein classifying something as spam or not. With machine data, it could be used to predict a material quality by several known parameters (e.g. humidity, strength, color, ...).

- **Artificial Neural Networks :**

- To understand the concept of the architecture of an artificial neural network, we have to understand what a neural network consists of. In order to define a neural network that consists of a large number of artificial neurons, which are termed unitsarranged in a sequence of layers.Lets us look at various types of layers available inan artificial neural network.

- > Input Layer

- > Hidden Layer

- > Output layer

- **Convolution Neural Networks :**

- A convolutional neural network, or CNN, is a deep learning neural network sketched for processing structured arrays of data such as portrayals.
- CNN are very satisfactory at picking up on design in the input image, such as lines, gradients, circles, or even eyes and faces.
- CNN can run directly on a underdone image and do not need any preprocessing.

- **Flask:**

- To put it to use in order to predict the new data, we have to deploy it over the internet so that the outside world can use it. In this article, we will talk about how we have trained a machine learning model and created a web application on it using Flask.

Prerequisites :

Anaconda Navigator :

- Anaconda Navigator is a free and open-source distribution of the Python and R programming languages for data science and machine learning-related applications. It can be installed on Windows, Linux, and macOS. Conda is an open-source, cross-platform, package management system. Anaconda comes with so very nice tools like JupyterLab, Jupyter Notebook, QtConsole, Spyder, Glueviz, Orange, Rstudio, Visual StudioCode. For this project, we will be using Jupiter notebook and spyder.

To build Deep learning models you must require the following packages :

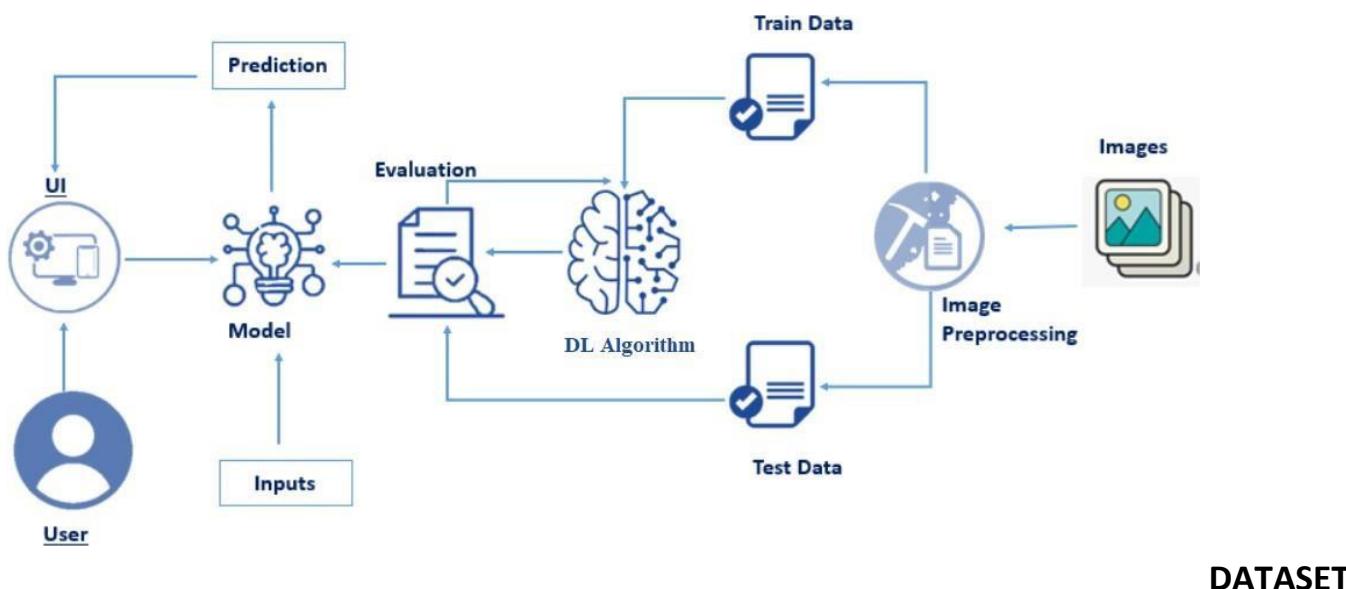
- **Tensor flow:** TensorFlow is an end-to-end open-source platform for machine learning. It has a comprehensive, flexible ecosystem of tools, libraries, and community resources that lets researchers push the state-of-the-art in ML and developers can easily build and deploy ML powered applications.

- **Keras** : Keras leverages various optimization techniques to make high level neural network API easier and more performant. It supports the following features.

1. Consistent, simple and extensible API.
2. Minimal structure - easy to achieve the result without any frills.
3. It supports multiple platforms and backend.
4. It is user-friendly framework that runs on both CPU and GPU.
5. Highly scalability of computation.

- **Flask**: Web framework used for building Web applications.

Technical Architecture :



DATASET

TEAM ID	PNT2022TMID49029
PROJECT NAME	Fertilizer Recommendation System For Disease Prediction
MAXIMUM MARK	4 Marks

Dataset:

<https://drive.google.com/file/d/1fxs7ptl6zh7NTbCOZARKZ7AmYKjnprY/view?usp=sharing>

IBM PROJECT

Fertilizers Recommendation System For Disease Prediction

ABSTRACT

- ✓ Agriculture is the most important sector in today's life. Most plants are affected by a wide variety of bacterial and fungal diseases. 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.

EXISTING PROBLEM

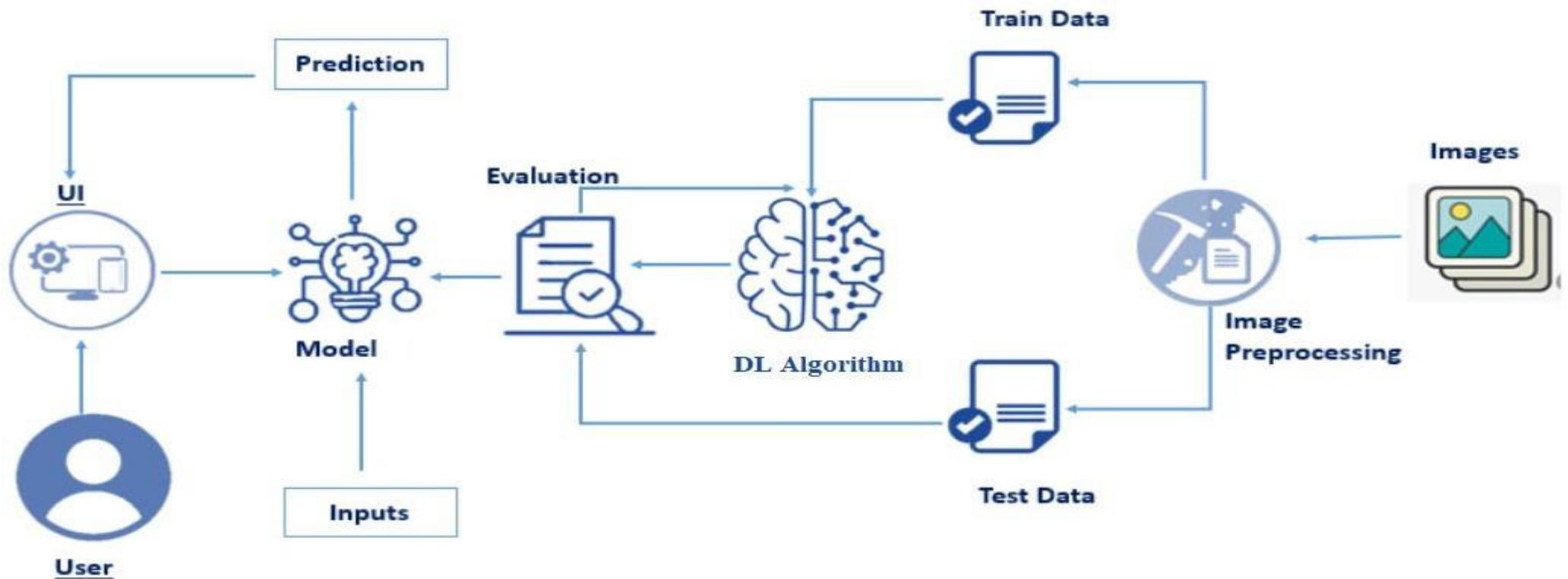
- ✓ Adequate mineral nutrition is central to crop production. However, it can also exert considerable influence on disease development. Fertilizer application can increase or decrease development of diseases caused by different pathogens, and the mechanisms responsible are complex, including effects of nutrients on plant growth, plant resistance mechanisms and direct effects on the pathogen. The effects of mineral nutrition on plant disease and the mechanisms responsible for those effects have been dealt with comprehensively elsewhere. In India, around 40% of land is kept and grown using reliable irrigation technologies, while the rest relies on the monsoon environment for water. Irrigation decreases reliance on the monsoon, increases food security, and boosts agricultural production.
- ✓ Most research articles use humidity, moisture, and temperature sensors near the plant's root, with an external device handling all of the data provided by the sensors and transmitting it directly to an external display or an Android application. The application was created to measure the approximate values of temperature, humidity and moisture sensors that were programmed into a microcontroller to manage the amount of water.

PROPOSED SYSTEM

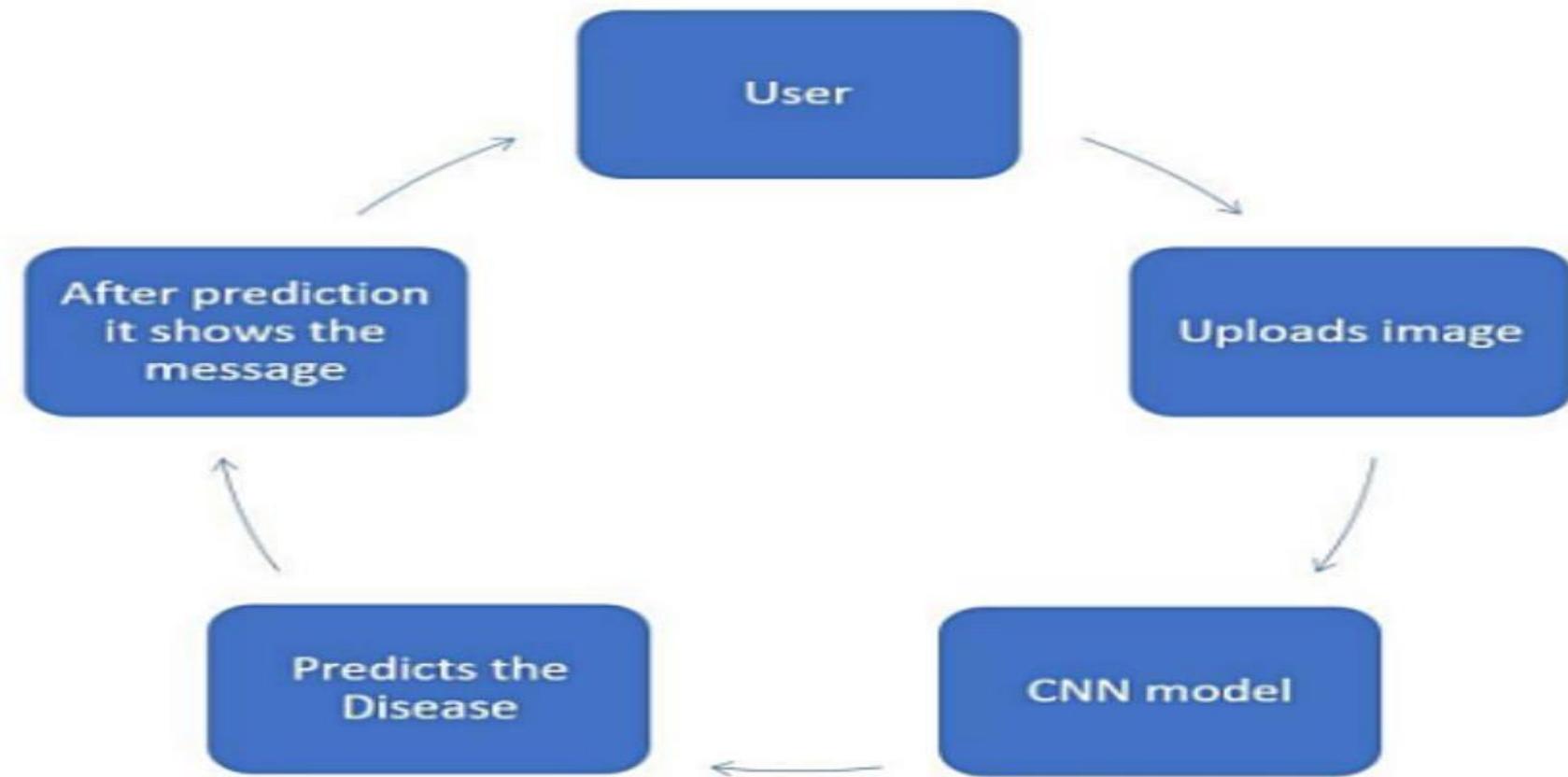
So we have built Web Application where:

- Farmers can easily interact with the application that we have built.
- This web application provides the user interface to upload images of diseased crop or leaf.
- Detection and recognition of plant diseases using machine learning are very efficient in providing symptoms of identifying diseases at its earliest.
- Our web application analysis the disease of the crop or leaf and suggests the farmer with the fertilizers are to be used.
 - It recommends the fertilizer for affected leaves based on severity level.
- This web application makes the farmers to take right decision in selecting the fertilizer for crop disease such that agricultural sector will be developed by innovative idea.

TECHNICAL ARCHITECTURE



FLOW CHART



HARDWARE & SOFTWARE REQUIREMENT

✓ To complete this project, you should have the following software and packages.

Software's:

- Anaconda Navigator
- py charm
- Visual studio code
- Jupiter notebook
- IBM Watson studio

Packages:

- Tensor flow
- Keras
- Flask
- numpy
- Pandas

OUTPUT

Plant Disease Prediction

Home Predict

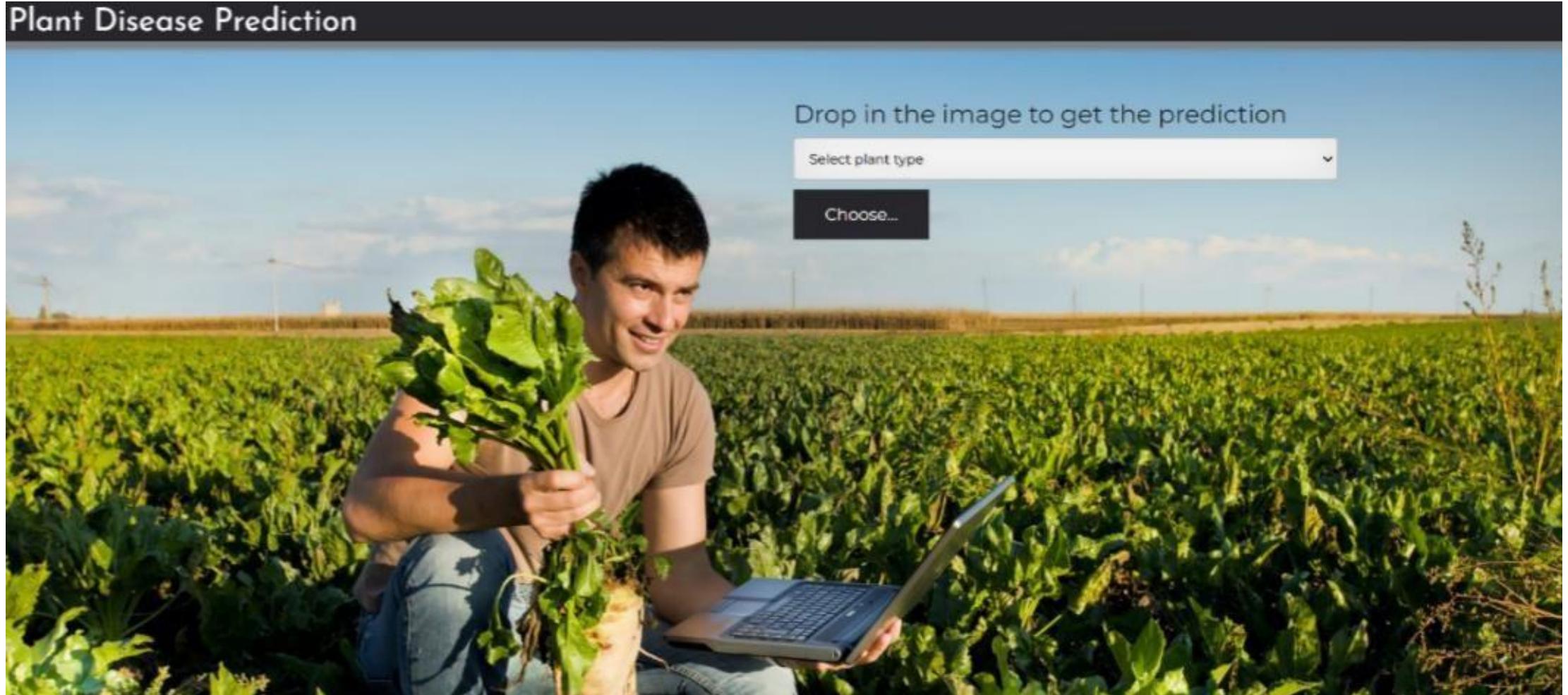
Fertilizers Recommendation System For Disease Prediction!!

Agriculture is one of the major sectors world wide. Over the years it has developed and the use of new technologies and equipment replaced almost all the traditional methods of farming. The plant diseases effect the production. Identification of diseases and taking necessary precautions is all done through naked eye, which requires labour and laboratories. This application helps farmers in detecting the diseases by observing the spots on the leaves, which inturn saves effort and labour costs.



Presented by [redacted]

OUTPUT



OUTPUT

Plant Disease Prediction

Drop in the image to get the prediction

Fruit

Choose...

Prediction: Opps!! Your apple plant is infected by Black Rots. This infection is a fungal infection. To control black rot, remove the cankers by pruning at least 15 inches below the end and burn or bury them. Treating the sites with the antibiotic streptomycin or a copper-based fungicide will be helpful.

APPLICATION

- This web application can be used by farmers or users to check whether their plant is infected or not and can also show the remedy so that the user can take necessary precautions.
- These kind of web applications can be used in the agricultural sector as well as for small house hold plants as well.



Thank you

Presentation title

PROJECT REPORT

Fertilizers Recommendation System for Disease Prediction

Team ID: PNT2022TMID49029

TEAM MEMBERS:

P.NAGAJOTHI (921619106041)

M.BHARATHI PRIYA (921619106009)

T.DIVYADHARSHINI(921619106018)

S.YUVASHREE(921619106067)

ABSTRACT:

- ✓ 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:

- ✓ To Detect and recognize the plant diseases and to recommend fertilizer, it is necessary to provide symptoms in identifying the disease at its earliest.

Hence the authors proposed and implemented new fertilizers
Recommendation System for crop disease prediction.

Presentation title

LITERATURE SURVEY:

Existing Problem:

- ✓ Adequate mineral nutrition is central to crop production. However, it can also exert considerable influence on disease development. Fertilizer application can increase or decrease development of diseases caused by different pathogens, and the mechanisms responsible are complex, including effects of nutrients on plant growth, plant resistance mechanisms and direct effects on the pathogen. The effects of mineral nutrition on plant disease and the mechanisms responsible for those effects have been dealt with comprehensively elsewhere. In India, around 40% of land is kept and grown using reliable irrigation technologies, while the rest relies on the monsoon environment for water. Irrigation decreases reliance on the monsoon, increases food security, and boosts agricultural production.
- ✓ Most research articles use humidity, moisture, and temperature sensors near the plant's root, with an external device handling all of the data provided by the sensors and transmitting it directly to an external display or an Android application. The application was created to measure the approximate values of temperature, humidity and moisture sensors that were programmed into a microcontroller to manage the amount of water.

Presentation title

Proposed solution:

So, we have built Web Application where:

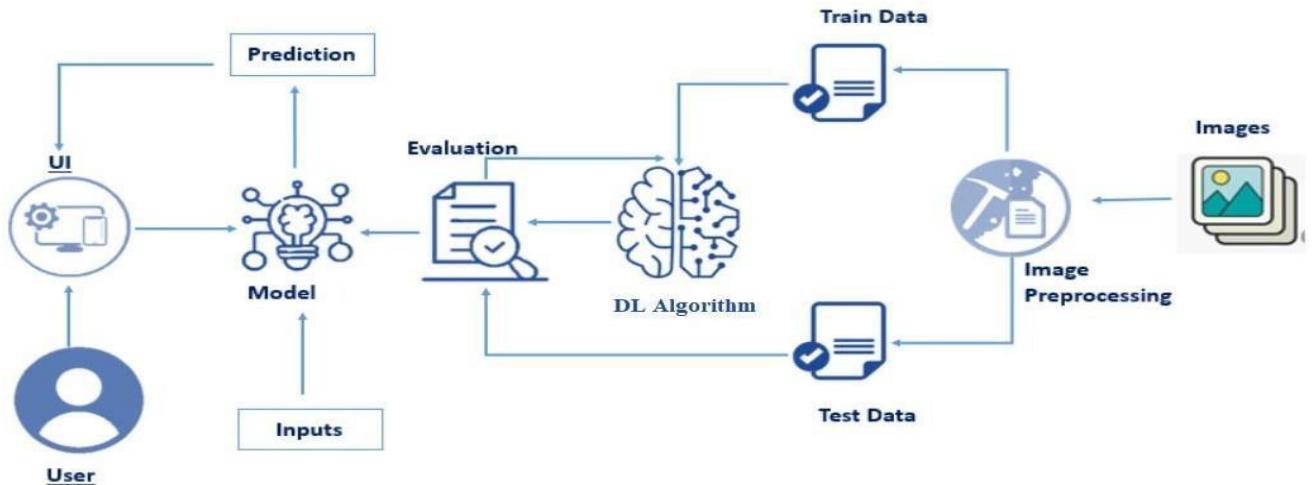
1. Farmers interact with the portal build
2. Interacts with the user interface to upload images of diseased leaf
3. Our model built analyses the Disease and suggests the farmer with fertilizers are to be used

- Detection and recognition of plant diseases using machine learning are very efficient in providing symptoms of identifying diseases at its earliest.
 - It recommends the fertilizer for affected leaves based on severity level.
- This web application makes the farmers to take right decision in selecting

the fertilizer for crop disease such that agricultural sector will be developed by innovative idea.

Presentation title

TECHNICAL ARCHITECTURE:



HARDWARE / SOFTWARE REQUIREMENTS:

To complete this project, you should have the following software and packages.

Software's:

- Anaconda Navigator
 - py charm
 - Visual studio code
 - Jupiter notebook
 - IBM Watson studio

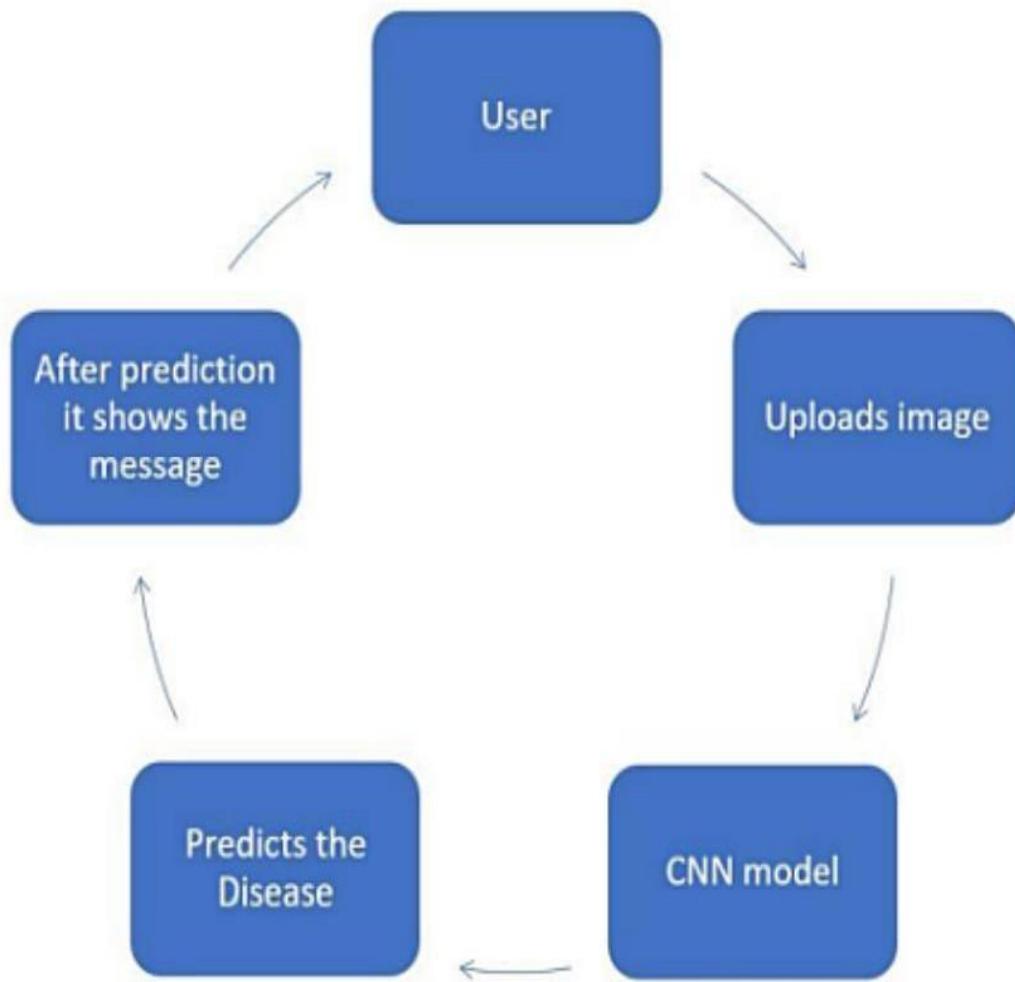
Presentation title

Packages:

- Tensor flow
 - Keras
 - Flask
 - numpy
 - Pandas

By using the above listed software's and packages, we built this application to take the input as image from the Farmer and detects whether the plant is infected or not. Here we use Deep learning techniques and give the output to the user as Farmer.

FLOWCHART:



Presentation title

To accomplish the above task,
you must complete the below
activities and tasks:

- Download the dataset.
- Classify the dataset into train and test sets.
- Add the neural network layers.
- Load the trained images and fit the model.

- Test the model.
- Save the model and its dependencies.
- Build a Web application using a flask that integrates with the model built.

Presentation title

OUTPUT:

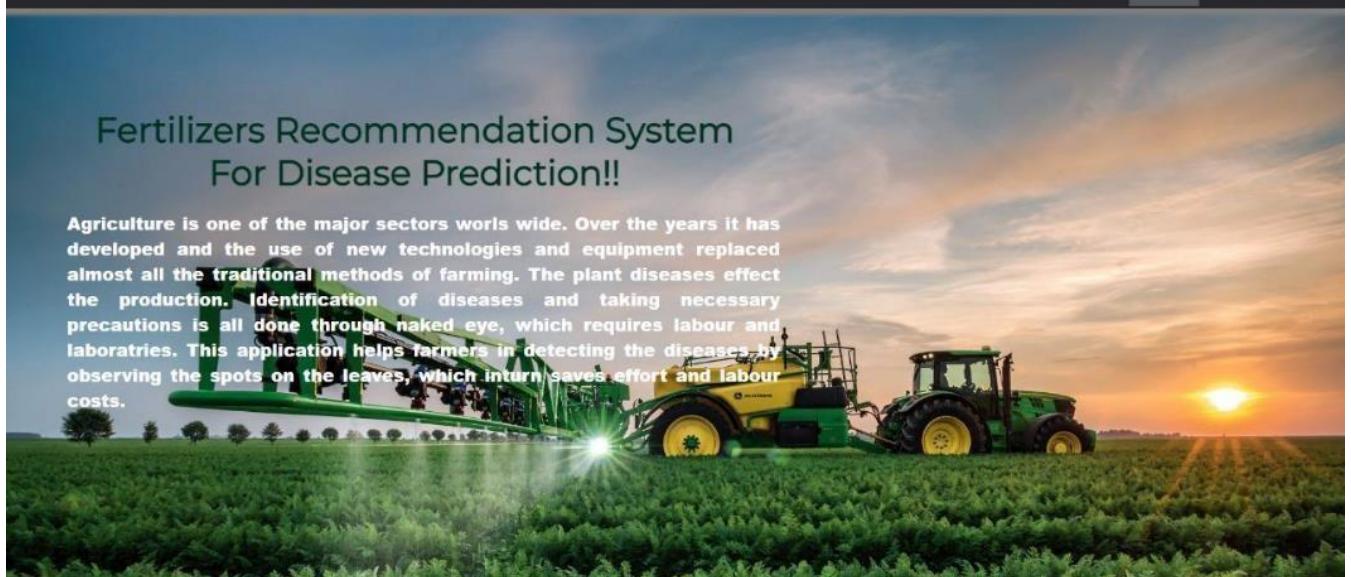
Home Page:

Plant Disease Prediction

Home Predict

Fertilizers Recommendation System For Disease Prediction!!

Agriculture is one of the major sectors world wide. Over the years it has developed and the use of new technologies and equipment replaced almost all the traditional methods of farming. The plant diseases effect the production. Identification of diseases and taking necessary precautions is all done through naked eye, which requires labour and laboratories. This application helps farmers in detecting the diseases by observing the spots on the leaves, which in turn saves effort and labour costs.



Prediction Page:

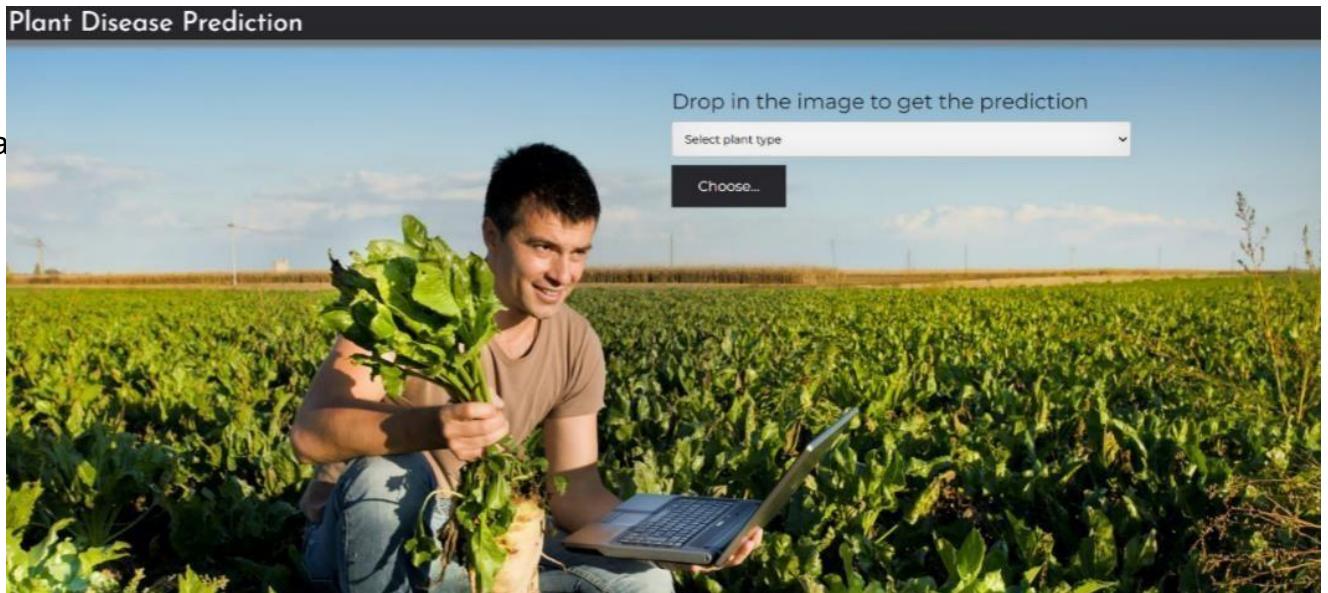
Plant Disease Prediction

Presenta

Drop in the image to get the prediction

Select plant type

Choose...



Result Page:

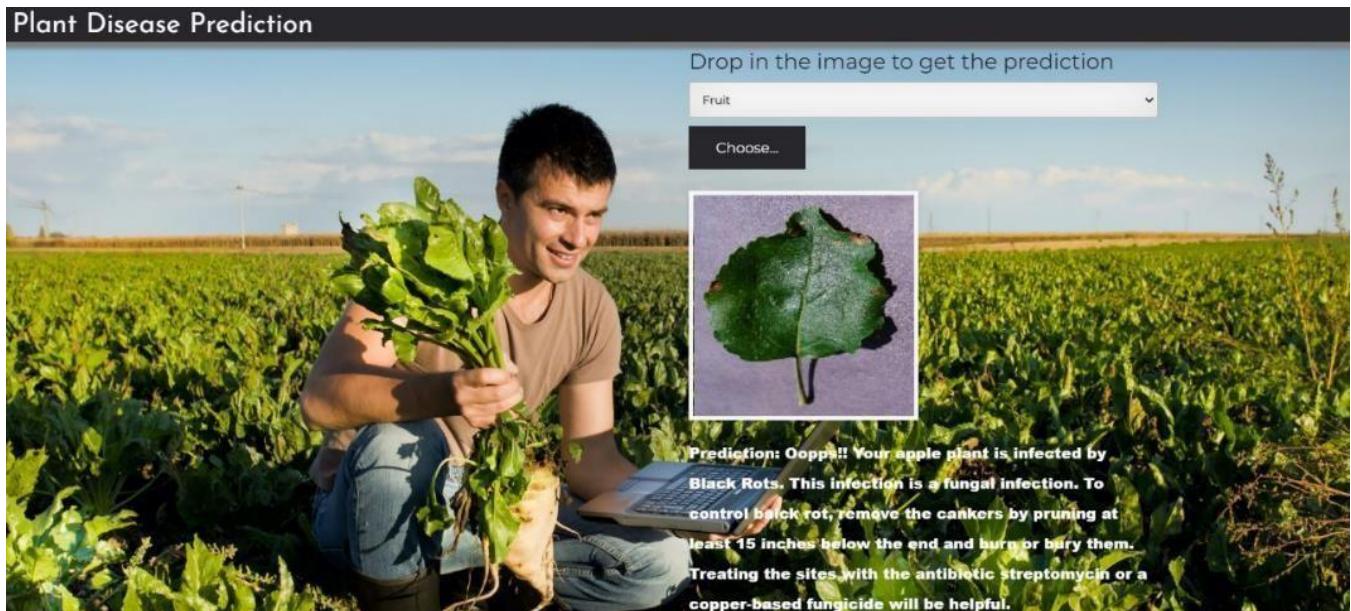
⇒ **Fruit:**

Plant Disease Prediction

Drop in the image to get the prediction

Fruit

Choose...



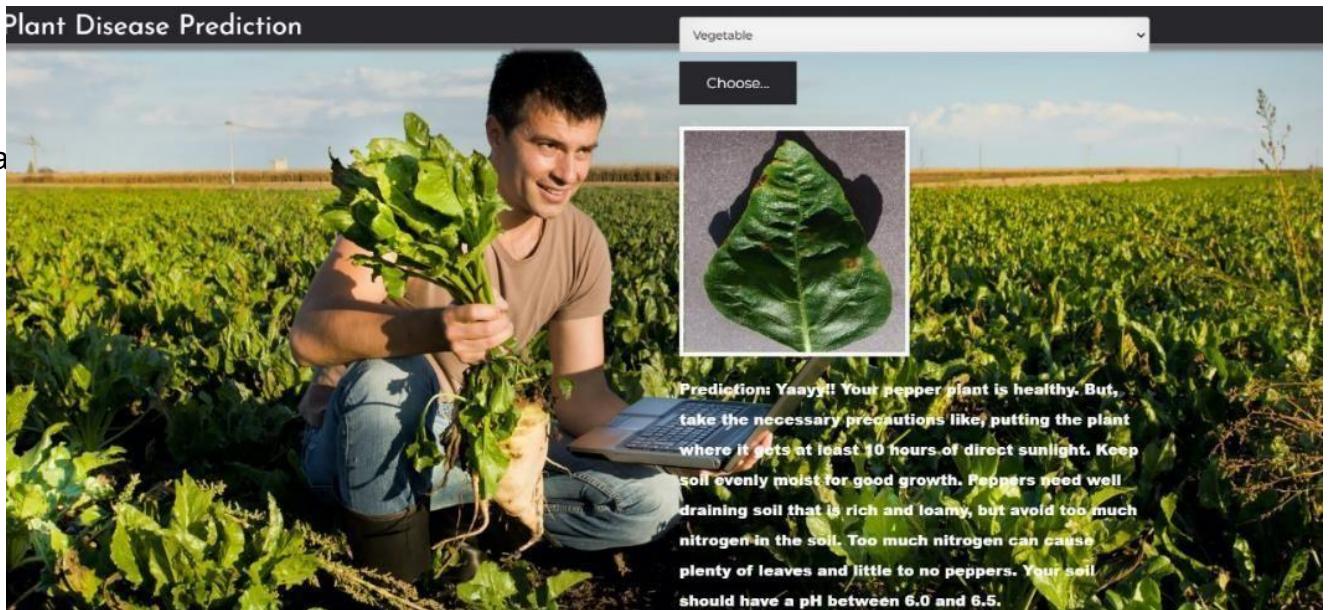
Prediction: Oopp!! Your apple plant is infected by Black Rots. This infection is a fungal infection. To control black rot, remove the cankers by pruning at least 15 inches below the end and burn or bury them. Treating the sites with the antibiotic streptomycin or a copper-based fungicide will be helpful.

⇒ **Vegetable:**

Plant Disease Prediction

Vegetable

Choose...



Prediction: Yaay!! Your pepper plant is healthy. But, take the necessary precautions like putting the plant where it gets at least 10 hours of direct sunlight. Keep soil evenly moist for good growth. Peppers need well draining soil that is rich and loamy, but avoid too much nitrogen in the soil. Too much nitrogen can cause plenty of leaves and little to no peppers. Your soil should have a pH between 6.0 and 6.5.

ADVANTAGES:

- The proposed model could predict the disease just from the image of a particular plant.
 - Easy to use UI.
- Model has some good accuracy in detecting the plant just by taking the input(leaf).

APPLICATIONS:

- This web application can be used by farmers or users to check whether their plant is infected or not and can also show the remedy so that the user can take necessary precautions.
- These kind of web applications can be used in the agricultural sector as well as for small house hold plants as well.

CONCLUSION:

- 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.

Presentation title

- 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. Usage of such applications could help the farmers to necessary precautions so that they don't face any loss as such.

FUTURE SCOPE:

- 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.

Presentation title

- This can be also done in Mobile applications like android, ios. It helps in many ways to improve the agriculture in cultivation of crops and predict the correct fertilizers to the crops.

APPENDIX:

Source Code:

```
In [1]: ls
```

```
In [2]: pwd
```

```
Out[2]: '/home/wsuser/work'
```

```
In [8]: !pip install keras==2.7.0  
!pip install tensorflow==2.5.0
```

```
Collecting keras==2.7.0
  Using cached keras-2.7.0-py2.py3-none-any.whl (1.3 MB)
Installing collected packages: keras
  Attempting uninstall: keras
    Found existing installation: Keras 2.2.4
    Uninstalling Keras-2.2.4:
      Successfully uninstalled Keras-2.2.4
Successfully installed keras-2.7.0
Requirement already satisfied: tensorflow==2.5.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (2.5.0)
Requirement already satisfied: protobuf>=3.9.2 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (3.19.1)
Requirement already satisfied: h5py~=3.1.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (3.1.0)
Requirement already satisfied: astunparse~=1.6.3 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (1.6.3)
Requirement already satisfied: keras-nightly~=2.5.0.dev in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (2.5.0.dev2021032900)
Requirement already satisfied: termcolor~=1.1.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (1.1.0)
Requirement already satisfied: flatbuffers~=1.12.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (1.12)
Requirement already satisfied: wrapt~=1.12.1 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (1.12.1)
Requirement already satisfied: six~=1.15.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0)
```

```

Requirement already satisfied: typing-extensions~=3.7.4 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow<2.5.0) (3.7.4.3)
Requirement already satisfied: keras-preprocessing~=1.1.2 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow<2.5.0) (1.1.2)
Requirement already satisfied: absl-py~=0.10 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (0.12.0)
Requirement already satisfied: grpcio~=1.34.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (1.34.1)
Requirement already satisfied: numpy~=1.19.2 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (1.19.5)
Requirement already satisfied: google-pasta~=0.2 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (0.2.0)
Requirement already satisfied: wheel~=0.35 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (0.37.0)
Requirement already satisfied: opt-einsum~=3.3.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (3.3.0)
Requirement already satisfied: gast==0.4.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (0.4.0)
Requirement already satisfied: tensorboard~=2.5 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow==2.5.0) (2.7.0)
Requirement already satisfied: google-auth<3,>=1.6.3 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorflow<2.5->tensorflow==2.5.0) (1.23.0)
Requirement already satisfied: markdown>=2.6.8 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~>2.5->tensorflow==2.5.0) (3.3.3)
Requirement already satisfied: werkzeug>=0.11.15 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~>2.5->tensorflow==2.5.0) (2.0.2)
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~>2.5->tensorflow==2.5.0) (0.6.1)
Requirement already satisfied: tensorboard-plugin-wit>=1.6.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~>2.5->tensorflow==2.5.0) (1.6.0)
Requirement already satisfied: setuptools>=41.0.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~>2.5->tensorflow==2.5.0) (58.0.4)
Requirement already satisfied: requests<3,>=2.21.0 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from tensorboard~>

```

Image Augmentation

```

In [9]: from tensorflow.keras.preprocessing.image import ImageDataGenerator

In [10]: train_datagen=ImageDataGenerator(rescale=1./255,zoom_range=0.2,horizontal_flip=True,vertical_flip=False)

In [11]: test_datagen=ImageDataGenerator(rescale=1./255)

In [12]: ls

In [13]: pwd
Out[13]: '/home/wsuser/work'

```

```

In [14]:
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_4ff9f1114db24196a9abd4f5c1f0b60a = ibm_boto3.client(service_name='s3',
    ibm_api_key_id='j4lNXssktSSxQiDx3pbMR_eFi1SMCDE6MFnBQ_EmNCDM',
    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')

```

Presenta

```
streaming_body_1 = client_4ff9f1114db24196a9abd4f5c1f0b60a.get_object(Bucket='trainmodel-donotdelete-pr-cbqe37eh8gzes', Key='fru  
  
# 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 [15]: from io import BytesIO  
import zipfile  
unzip = zipfile.ZipFile(BytesIO(streaming_body_1.read()), "r")  
file_paths = unzip.namelist()  
for path in file_paths:  
    unzip.extract(path)
```

```
In [16]: pwd
```

```
Out[16]: '/home/wsuser/work'
```

```
In [17]: import os  
filenames = os.listdir('/home/wsuser/work/fruit-dataset/train')
```

Presentatio

```
In [18]: x_train=train_datagen.flow_from_directory("/home/wsuser/work/fruit-dataset/train",target_size=(128,128),class_mode='categorical')
```

```
Found 5384 images belonging to 6 classes.
```

```
In [ ]:
```

```
In [19]: x_test=test_datagen.flow_from_directory(r"/home/wsuser/work/fruit-dataset/test",target_size=(128,128),  
class mode='categorical',batch size=24)
```

```
Found 1686 images belonging to 6 classes.
```

```
In [20]: x_train.class_indices  
Out[20]: {'Apple__Black_rot': 0,  
          'Apple__healthy': 1,  
          'Corn_(maize)___Northern_Leaf_Blight': 2,  
          'Corn_(maize)___healthy': 3,  
          'Peach__Bacterial_spot': 4,  
          'Peach__healthy': 5}
```

CNN

```
In [21]: from tensorflow.keras.models import Sequential  
from tensorflow.keras.layers import Dense, Convolution2D, MaxPooling2D, Flatten  
  
In [24]: model=Sequential()  
  
In [25]: model.add(Convolution2D(32,(3,3),input_shape=(128,128,3),activation='relu'))  
  
In [26]: model.add(MaxPooling2D(pool_size=(2,2)))  
  
In [27]: model.add(Flatten())  
  
In [28]: model.summary()  
  
Model: "sequential_1"  
-----  
Layer (type)           Output Shape        Param #  
=====  
conv2d_1 (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
```

Presentatio

```
In [29]: 32*(3*3*3+1)
```

```
Out[29]: 896
```

Hidden Layers

```
In [30]: model.add(Dense(300,activation='relu'))  
model.add(Dense(150,activation='relu'))
```

Output Layer

```
In [31]: model.add(Dense(6,activation='softmax'))
```

```
In [32]: model.compile(loss='categorical_crossentropy',optimizer='adam',metrics=['accuracy'])
```

```
In [33]: len(x_train)
Out[33]: 225

In [34]: 1238/24
Out[34]: 51.58333333333336

In [35]: model.fit_generator(x_train,steps_per_epoch=len(x_train),validation_data=x_test,validation_steps=len(x_test),epochs=10)
          /tmp/wsuser/ipykernel_164/1582812018.py:1: 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=len(x_train),validation_data=x_test,validation_steps=len(x_test),epochs=10)

          Epoch 1/10
          225/225 [=====] - 118s 520ms/step - loss: 0.8920 - accuracy: 0.8094 - val_loss: 0.2273 - val_accuracy: 0.9235
          Epoch 2/10
          225/225 [=====] - 116s 515ms/step - loss: 0.2367 - accuracy: 0.9179 - val_loss: 0.2056 - val_accuracy: 0.9324
          Epoch 3/10
          225/225 [=====] - 116s 517ms/step - loss: 0.1970 - accuracy: 0.9337 - val_loss: 0.4972 - val_accuracy: 0.8754
          Epoch 4/10
          225/225 [=====] - 117s 521ms/step - loss: 0.1688 - accuracy: 0.9422 - val_loss: 0.2279 - val_accuracy: 0.9217
          Epoch 5/10
          225/225 [=====] - 116s 516ms/step - loss: 0.1438 - accuracy: 0.9487 - val_loss: 0.1685 - val_accuracy: 0.9484
          Epoch 6/10
          225/225 [=====] - 117s 518ms/step - loss: 0.1362 - accuracy: 0.9556 - val_loss: 0.1176 - val_accuracy: 0.9662
          Epoch 7/10
          225/225 [=====] - 116s 515ms/step - loss: 0.1282 - accuracy: 0.9590 - val_loss: 0.5466 - val_accuracy: 0.951
          Epoch 8/10
          225/225 [=====] - 116s 514ms/step - loss: 0.1282 - accuracy: 0.9597 - val_loss: 0.1194 - val_accuracy: 0.9620
          Epoch 9/10
          225/225 [=====] - 116s 514ms/step - loss: 0.1141 - accuracy: 0.9616 - val_loss: 0.1478 - val_accuracy: 0.9508
          Epoch 10/10
          225/225 [=====] - 116s 516ms/step - loss: 0.0927 - accuracy: 0.9695 - val_loss: 0.0772 - val_accuracy: 0.9751

Out[35]: <keras.callbacks.History at 0x7f71e8184070>
```

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Saving Model

```
In [36]: ls
fruit-dataset/

In [37]: model.save('fruit.h5')

In [40]: !tar -zcvf Train-model_new.tgz fruit.h5
fruit.h5

In [39]: ls -1
fruit-dataset/
fruit.h5
Train-model_new.tgz
```

IBM Cloud Deployment Model

```
In [41]: !pip install watson-machine-learning-client --upgrade
Collecting watson-machine-learning-client
  Downloading watson_machine_learning-client-1.0.391-py3-none-any.whl (538 kB)
    |████████| 538 kB 21.2 MB/s eta 0:00:01
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: certifi in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (2022.9.24)
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: 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: lomond in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (0.3.3)
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: urllib3 in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from watson-machine-learning-client) (1.26.7)
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: 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: 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 boto3core<1.22.0,>=1.21.21->boto3->watson-machine-learning-client) (2.8.2)

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) (1.19.5)
Installing collected packages: watson-machine-learning-client
Successfully installed watson-machine-learning-client-1.0.391
```

```
In [43]: from ibm_watson_machine_learning import APIClient
wml_credentials = {
    "url": "https://us-south.ml.cloud.ibm.com",
    "apikey": "0P3XkyCFYqABnc48BNG2ReoGAJy-oDXDRuULl4Y_zFxa"
}
client = APIClient(wml_credentials)

In [44]: client = APIClient(wml_credentials)

In [45]: def guid_from_space_name(client, space_name):
    space = client.spaces.get_details()
    return(next(item for item in space['resources'] if item['entity'][name]==space_name)[metadata]['id'])

In [46]: space_uid = guid_from_space_name(client, 'Trainmodel')
print("Space UID = " + space_uid)

Space UID = 616c7d74-e99b-4c09-9922-27394a62c2d0

In [47]: client.set.default_space(space_uid)

Out[47]: 'SUCCESS'

In [48]: client.software_specifications.list()
```

Presenta

NAME	ASSET_ID	TYPE
default_py3.6	0062bc9-8b7d-44a0-a9b9-46c416adcbd9	base
kernel-spark3.2-scala2.12	020d9ce-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-b5f6fcc6471	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	10ac12d6-6b30-4cc0-8392-3e922c096a92	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-4d5ee5abb85	base
default_r3.6	1b70aec3-ab34-4b87-8aa0-a4a3c8296a36	base
pytorch-onnx_1.3-py3.6	1bc6029a-cc97-56da-b8e0-39c3880dbbe7	base
kernel-spark3.3-r3.6	1c9e5454-f216-59d4-a20e-474a5cdf5988	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-d466306ce658	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
kernel-spark3.3-py3.9	2b7961e2-e3b1-5a8c-a491-482c8368839a	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

Note: Only first 50 records were displayed. To display more use 'limit' parameter.

```
In [51]: software_space_uid = client.software_specifications.get_uid_by_name("tensorflow_rt22.1-py3.9")
software_spec_uid
```

```
Out[51]: '1eb25b84-d6ed-5dde-b6a5-3fbdf1665666'
```

```
In [54]: ls
```

```
fruit-dataset/ fruit.h5 Train-model_new.tgz
```

Presentatio

```
In [56]: model_details = client.repository.store_model(model= 'Train-model_new.tgz',
meta_props={
    client.repository.ModelMetaNames.NAME:"CNN",
    client.repository.ModelMetaNames.TYPE:"tensorflow_2.7",
    client.repository.ModelMetaNames.SOFTWARE_SPEC_UID:software_space_uid
})
```

```
In [57]: model_id = client.repository.get_model_id(model_details)
```

```
In [58]: model_id
```

```
Out[58]: 'd0aeb6a2-e89c-4f8d-bf2f-a28ca4ea3cca'
```

```
In [60]: ls
```

```
fruit-dataset/ fruit.h5 Train-model_new.tgz
```

Test The Model

```
In [54]: import numpy as np
from tensorflow.keras.models import load_model
from tensorflow.keras.preprocessing import image

In [55]: model=load_model('fruit.h5')

In [68]: img=image.load_img(r"C:\Users\Sree Ram\Desktop\ibm\Dataset Plant Disease\fruit-dataset\fruit-dataset\test\Apple__healthy\0adc1c5f55334344a2e0333333333333.jpg")

In [69]: img
Out[69]: 
```



```
In [70]: img=image.load_img(r"C:\Users\Sree Ram\Desktop\ibm\Dataset Plant Disease\fruit-dataset\fruit-dataset\test\Apple__healthy\0adc1c5f55334344a2e0333333333333.jpg")
img
Out[70]: 
```


Presenta In [71]: x=image.img_to_array(img)

```
In [72]: x
Out[72]: array([[[ 99.,  86., 106.],
   [101.,  88., 108.],
   [118., 105., 125.],
   ...,
   [ 92.,  83., 102.],
   [ 93.,  84., 103.],
   [ 89.,  80.,  99.]],

   [[ 96.,  83., 103.],
   [ 87.,  74.,  94.],
   [102.,  89., 109.],
   ...,
   [ 88.,  79.,  98.],
   [ 89.,  80.,  99.],
   [ 83.,  74.,  93.]]],
```

```
...,
[ 88.,  79.,  98.],
[ 89.,  80.,  99.],
[ 83.,  74.,  93.]],

[[ 86.,  73.,  93.],
[ 88.,  75.,  95.],
[ 98.,  85., 105.],
...,
[107.,  98., 117.],
[ 96.,  87., 106.],
[ 96.,  87., 106.]],

...,

[[172., 175., 194.],
[173., 176., 195.],
[175., 178., 197.],
...,
[179., 180., 198.],
[184., 185., 203.],
[179., 180., 198.]],

[[172., 175., 194.],
[170., 173., 192.],
[173., 176., 195.],
...,
[178., 179., 197.],
[182., 183., 201.],
[178., 179., 197.]],

[[169., 172., 191.],
[166., 169., 188.],
[168., 171., 190.],
...,
...,
[187., 188., 206.],
[185., 186., 204.],
[186., 187., 205.]]], dtype=float32)
```

In [73]: `x=np.expand_dims(x,axis=0)`

In [74]: `x`

```
Out[74]: array([[[[ 99.,  86., 106.],
[101.,  88., 108.],
[118., 105., 125.],
...,
[ 92.,  83., 102.],
[ 93.,  84., 103.],
[ 89.,  80.,  99.]],

[[ 96.,  83., 103.],
[ 87.,  74.,  94.],
[102.,  89., 109.],
...,
[ 88.,  79.,  98.],
[ 89.,  80.,  99.],
[ 83.,  74.,  93.]],

[[ 86.,  73.,  93.],
[ 88.,  75.,  95.],
[ 98.,  85., 105.],
...,
[107.,  98., 117.],
[ 96.,  87., 106.],
[ 96.,  87., 106.]],

...,
```

Presenta

```

    ...,
[187., 188., 206.],
[185., 186., 204.],
[186., 187., 205.]]], dtype=float32)

In [75]: y=np.argmax(model.predict(x),axis=1)
          1/1 [=====] - 0s 105ms/step

In [76]: x_train.class_indices
Out[76]: {'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 [77]: index=['Apple__Black_rot','Apple__healthy','Corn_(maize)___Northern_Leaf_Blight','Corn_(maize)___healthy','Peach__Bacterial_sp
          ↴
          ↵

In [78]: index[y[0]]
Out[78]: 'Apple__healthy'

In [82]: img=image.load_img(r"C:\Users\Sree Ram\Desktop\ibm\Dataset Plant Disease\fruit-dataset\fruit-dataset\test\Peach__healthy\0a2ed4e
x=image.img_to_array(img)
x=np.expand_dims(x,axis=0)
y=np.argmax(model.predict(x),axis=1)
index=['Apple__Black_rot','Apple__healthy','Corn_(maize)___Northern_Leaf_Blight','Corn_(maize)___healthy','Peach__Bacterial_sp
index[y[0]]
          ↴
          ↵

          1/1 [=====] - 0s 26ms/step
Out[82]: 'Corn_(maize)___healthy'

In [83]: import os
from tensorflow.keras.models import load_model
from tensorflow.keras.preprocessing import image
from flask import Flask,render_template,request
          ↴

In [61]: app=Flask(__name__)

model=load_model("fruit.h5")

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

@app.route('/predict',methods=['GET','POST'])
def upload():
    if request.method=='POST':
        f=request.files['image']
        basepath=os.path.dirname('__file__')
        filepath=os.path.join(basepath,'uploads',f.filename)
        f.save(filepath)
        img=image.load_img(filepath,target_size=(128,128))
        x=image.img_to_array(img)
        x=np.expand_dims(x,axis=0)
        pred=np.argmax(model.predict(x),axis=1)
        index=['Apple__Black_rot','Apple__healthy','Corn_(maize)___Northern_Leaf_Blight','Corn_(maize)___healthy','Peach__Bacterial_sp
        text="The Classified Fruit disease is : "+str(index[pred[0]])
    return text
if __name__=='__main__':
    app.run(debug=False)
          ↴
          ↵

```

Presenta

REFERENCE:

- ⇒ [Fertilizers Recommendation System For Disease Prediction In Tree Leave | SemanticScholar](#)

- ⇒ [Soil Based Fertilizer Recommendation System for Crop Disease Prediction System\(ijetajournal.org\)](#)

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- ⇒ [A nutrient recommendation system for soil fertilization based on evolutionarycomputation - ScienceDirect](#)

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 - ⇒ [2204.11340.pdf \(arxiv.org\)](#)
 - ⇒ [371-376,Tesma405,IJEAST.pdf](#)

- ⇒ [CROFED - Crop and Fertilizer Recommendation and Disease diagnosis system usingMachine Learning and Internet of Things. \(ijirt.org\)](#)
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