

# POTATO DISEASE CLASSIFICATION

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# Potato Disease Classification Using Deep Learning

## 1. INTRODUCTION

### 1.1 Project Overview

Globally, the production of potatoes is a major agricultural activity that supports both economic stability and food security. On the other hand, a number of diseases can affect potato plants, which can lower crop quality and output. For efficient illness management, these diseases must be identified and classified as soon as possible. The goal of this research is to create a system that reliably classifies potato diseases using Convolutional Neural Networks (CNN) and deep learning techniques, enabling farmers to make well-informed decisions about disease prevention and treatment.

### 1.2 Purpose

Predicting potato leaf disease as soon as possible is crucial because it can have significant impacts on crop yield and quality. Early detection allows farmers to take prompt action to prevent the spread of the disease and reduce crop damage. Here are some reasons why predicting potato leaf disease early is essential:

1. Minimize crop loss: Potato leaf disease can significantly reduce crop yield and quality. By predicting the disease early, farmers can take timely measures to control its spread, thereby minimizing crop loss.
2. Reduce cost: Early detection of potato leaf disease can help farmers reduce costs associated with disease control measures, such as pesticides and other treatments. By identifying the disease early, farmers can target the specific area of the crop affected, which helps in reducing the overall cost of control measures.
3. Protect the environment: The excessive use of pesticides and other control measures can have negative impacts on the environment. Early detection of potato leaf disease can help farmers target only the affected areas and minimize the use of pesticides, reducing their environmental impact.
4. Improve crop quality: Potato leaf disease can affect the quality of the crop, making it less desirable to buyers. By predicting the disease early, farmers can take appropriate measures to prevent its spread, thereby improving the overall quality of the crop.

## **2. LITERATURE SURVEY**

### **2.1 Existing problem**

Plant disease classification challenges have seen the successful application of deep learning techniques, particularly Convolutional Neural Networks (CNNs), in numerous research. Transfer learning from pretrained models such as VGG19, ResNet50, and Inception has been used by researchers to effectively classify plant illnesses, especially potato infections. These models can identify a wide range of plant diseases with high accuracy, which helps with early identification and management. Large and diversified datasets, data augmentation techniques, and model interpretability are all highlighted in this literature as being vital factors to take into account while creating a potato disease classification system.

High-quality datasets are essential for training deep learning models that classify plant diseases, as previous studies have shown. Model development has benefited greatly from the availability of existing datasets, such as the Plant Village dataset and others. To increase the resilience and generalisation of models, researchers have used data preprocessing techniques such image scaling, normalisation, and data augmentation. In order to address frequent obstacles in plant disease classification, such as uneven class distribution and differences in lighting and background, some research has looked into domain-specific data augmentation techniques.

The literature emphasises the importance of developing accurate models, but it also emphasises the necessity of creating deployment interfaces that are easy to use. The combination of deep learning models with approachable applications to enable farmers to quickly identify agricultural illnesses has been covered in a number of publications. These apps must to work with web browsers, mobile devices, and Internet of Things gadgets, and they ought to give consumers dependable and timely results. In order to guarantee the system's practical applicability in actual agricultural settings, interpretability and user input integration are frequently explored in this domain's literature.

In summary, the body of current research offers a solid framework for creating a deep learning-based system for classifying potato diseases. It emphasises the crucial roles that high-quality datasets, data preparation, model architecture, and user-friendly deployment have in improving crop management in the agriculture industry and helping farmers.

### **2.2 References**

1. <https://www.cambridge.org/core/journals/advances-in-animal-biosciences/article/abs/potato-disease-classification-using-convolution-neural-networks/E9303F667377BD763C3054CB8488D36C>
2. [https://link.springer.com/chapter/10.1007/978-981-13-8406-6\\_37](https://link.springer.com/chapter/10.1007/978-981-13-8406-6_37)
3. <https://ieeexplore.ieee.org/abstract/document/9231784>
4. <https://philpapers.org/rec/ELSPCU>

## 2.3 Problem Statement Definition

An important part of world agriculture is the production of potatoes, which are both a basic food and a source of income for many farmers. On the other hand, a number of diseases that can seriously affect crop quality and productivity can affect potato plants. Effective disease management depends on the quick and precise detection of these conditions. Conventional disease identification techniques frequently call for the manual inspection of agricultural specialists, which can take time and be unavailable to all farmers. Convolutional neural networks (CNNs) and deep learning techniques are being used to create an automated and precise system for classifying potato diseases in order to meet this challenge. By helping farmers make knowledgeable decisions about disease prevention and treatment, this technology will eventually increase crop productivity and food security

Important elements of the issue statement:

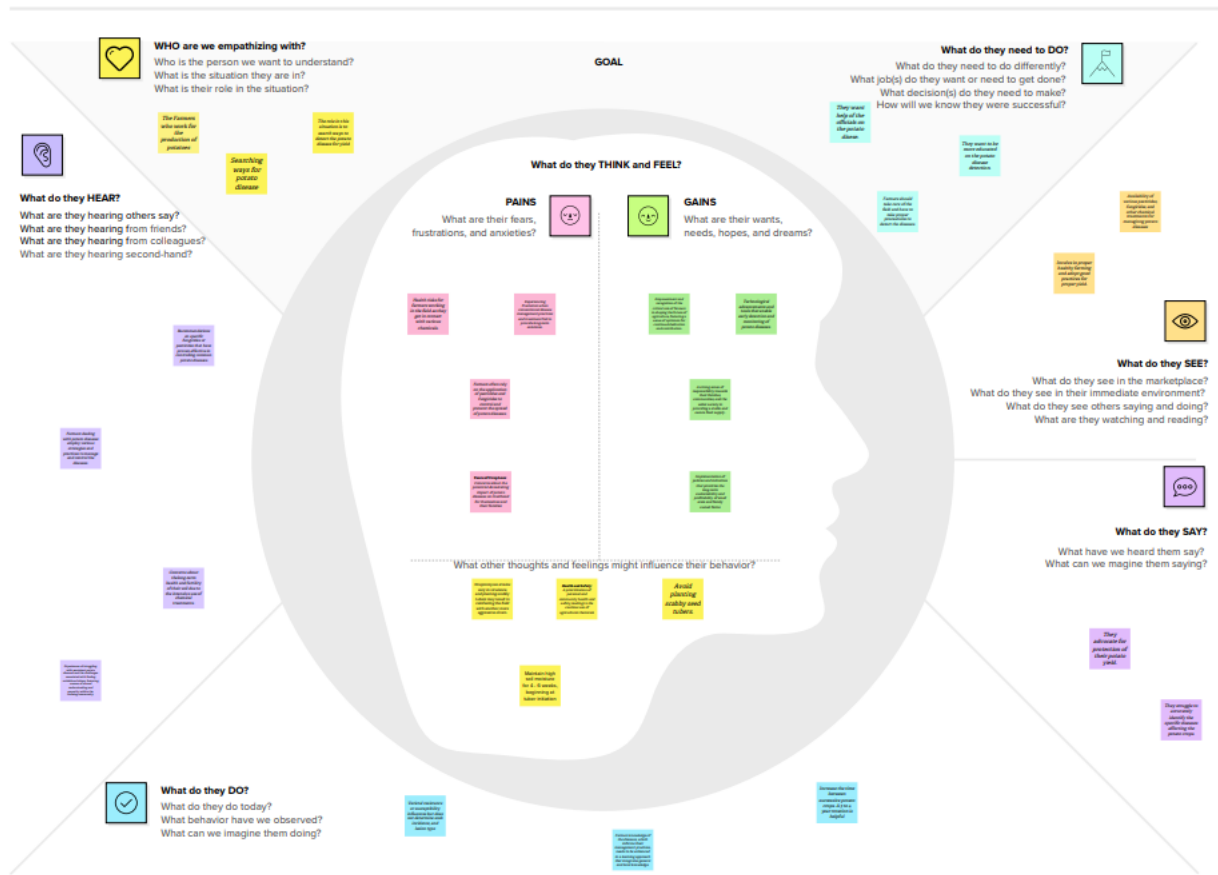
The main challenge is developing a trustworthy system for recognising and categorising the several diseases that impact potato plants, including potato scab, late blight, and early blight. The technology should be able to distinguish between plants that are healthy and those that are ill, and it should give precise details on the kind of disease that is present.

**Timeliness and Precision:** The system should deliver accurate and quick disease classification data so that farmers can begin managing and treating the illnesses that have been recognised right away. This takes care of the requirement for early disease identification, which can cut down on crop losses and excessive pesticide use.

User-Friendly Interface: To solve this problem effectively, a user-friendly interface or application should be developed, allowing farmers to easily upload images of their potato plants and receive disease classification results. The system should be intuitive and adaptable to the technological and educational background of the users.

### 3. IDEATION & PROPOSED SOLUTION

#### 3.1 Empathy Map Canvas





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### Group ideas

Take turns sharing your ideas while clustering similar or related notes as you go. Once all sticky notes have been grouped, give each cluster a sentence-like label. If a cluster is bigger than six sticky notes, try and see if you can break it up into smaller sub-groups.

🕒 20 minutes

#### TIP

Add customizable tags to sticky notes to make it easier to find, browse, organize, and categorize important ideas as themes within your mural.

**Data Analysis and Machine Learning:**  
Implement machine learning algorithms that analyze historical data on weather patterns, soil conditions, and disease occurrences.

*collaboration among farmers, agricultural experts, and researchers to share knowledge and best practices for disease prediction*

*Educate the farmers about the issue on potatoes diseases.*

*Train farmers and agricultural workers to recognize early symptoms of common potato leaf diseases.*

4

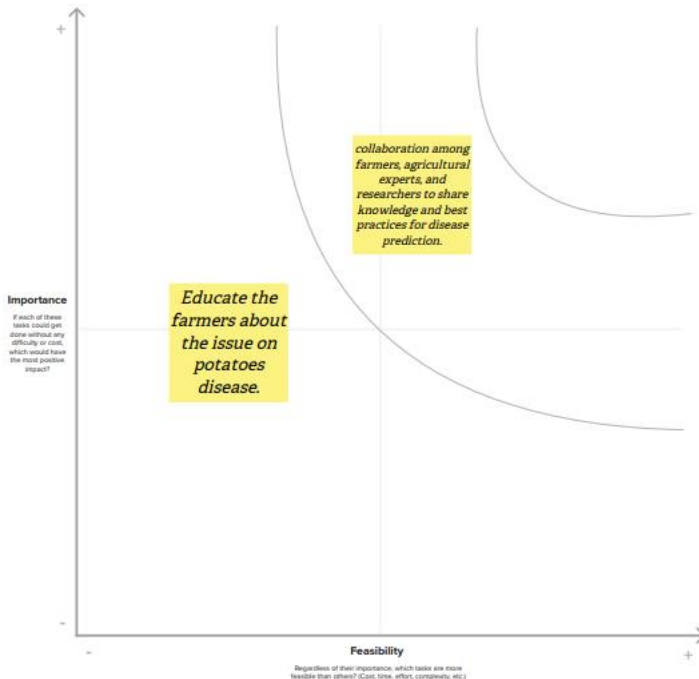
### Prioritize

Your team should all be on the same page about what's important moving forward. Place your ideas on this grid to determine which ideas are important and which are feasible.

🕒 20 minutes

#### TIP

Participants can use their cursor to point at where sticky notes should go on the grid. The facilitator can confirm the spot by using the laser pointer holding the H key on the keyboard.



## **4. REQUIREMENT ANALYSIS**

### **4.1 Functional requirement**

**Image Upload and Processing:** In order to classify diseases, users (farmers) should be able to upload photos of potato plants to the system. To guarantee that these photographs are compatible with the deep learning model, it should preprocess them, resizing and normalising them as needed.

**Disease Classification:** The system's main job is to correctly identify potato illnesses from the supplied photos. It should be able to distinguish between distinct disease states and healthy plant states, giving each case a unique disease identifier.

**User Interface:** A user-friendly interface that is compatible with PCs, tablets, and smartphones must be provided by the system. It ought to offer a simple and intuitive user interface to suit consumers with varying degrees of technological expertise.

**Model Interpretability:** The system ought to provide justifications or illustrations of the model's decision-making procedure, giving users an understanding of the rationale behind the classification of a certain ailment. This feature improves user comprehension and trust.

**Scalability and Integration:** To enable wider acceptance and influence within the agricultural sector, make sure the system is scalable and compatible with other agricultural technology and information systems.



## 4.2 Non-Functional requirements

**Accuracy and Precision:** For farmers to receive trustworthy findings, the disease classification system needs to attain a high degree of accuracy. It should also minimise false positives and false negatives by precisely identifying diseases.

**Response Time:** In order to enable farmers to take rapid action, the system should provide disease categorization results as soon as possible, ideally in real-time or within a few seconds of image input.

**Security and Privacy:** Put security measures in place to guard user information and guarantee the privacy of photos that users post. Images and personal data should be treated carefully and in accordance with applicable data protection laws.

**Robustness and Reliability:** Strong and dependable, the system should be able to adapt to changes in disease stages, lighting, and image quality. In the event of an unplanned outage, failover procedures ought to be included.

**Scalability and Performance:** Make sure the system is capable of managing a sizable volume of user requests and expanding to accommodate a rising user base without experiencing performance issues. Optimization and load balancing ought to be implemented.

## **5. PROJECT DESIGN**

### **5.1 Data Flow Diagrams & User Stories**

**Data Collection:** Obtaining a wide range of potato leaf image datasets from image archives, field surveys, and other sources is the task of this phase. Following collection, the photos are kept in a raw data repository.

**Data Pre-processing:** Images of raw potato leaves are pre-processed in order to get them ready for model training. To improve dataset diversity, this can involve resizing photographs, normalising pixel values, and using data augmentation techniques.

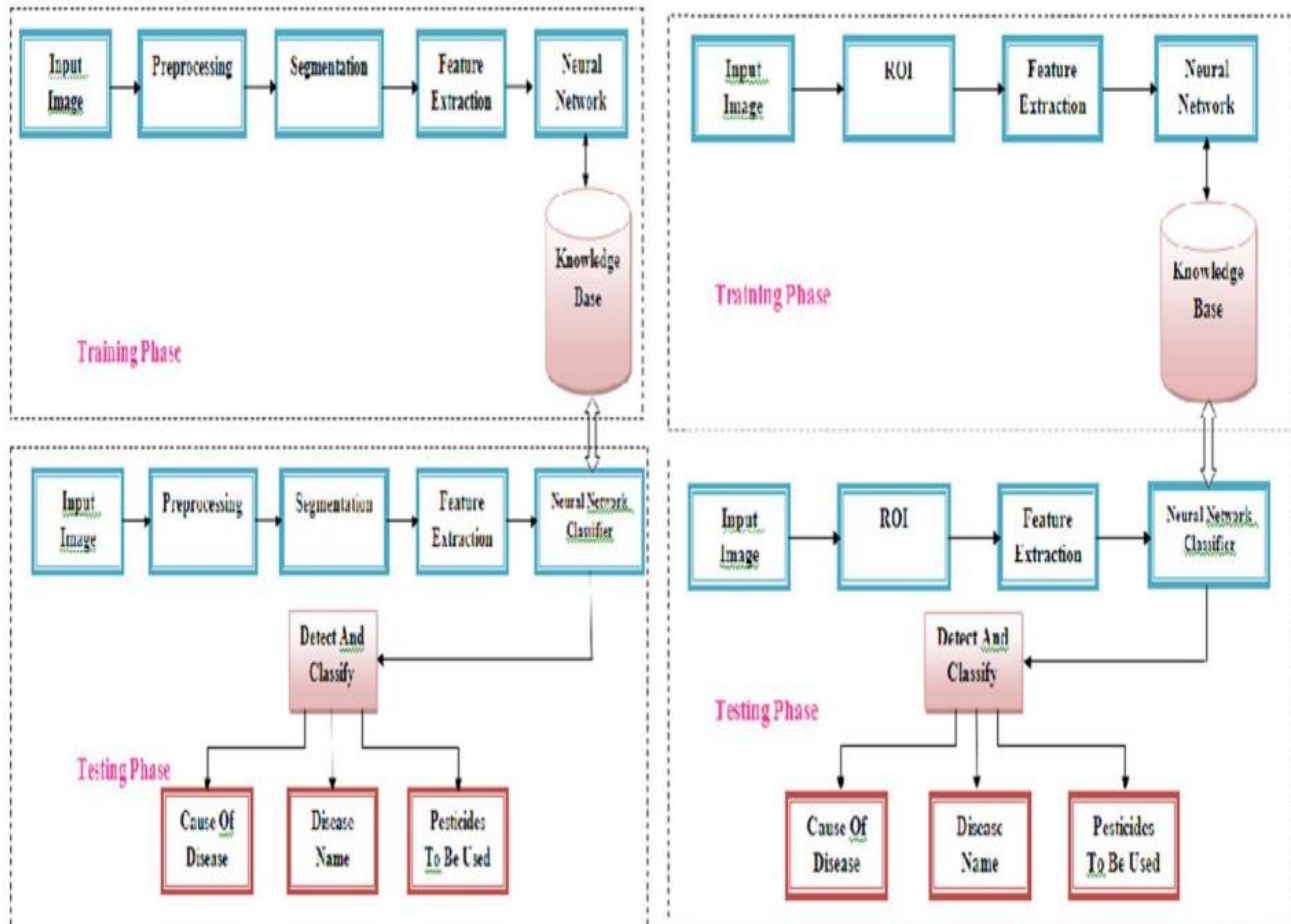
**Model Training:** This step involves using the pre-processed data to train a deep learning model so that it can identify different potato leaf diseases. For later use, the trained model is stored.

**Model Evaluation:** The performance of the trained model is assessed using a separate dataset not used in training, to gauge its accuracy, sensitivity, and specificity in disease classification.

**Model Deployment:** This step involves deploying the trained model to make it accessible for real-world disease classification applications, either on local devices or in the cloud.

**User Interaction:** Through an intuitive application or API, end users engage with the deployed model, allowing them to upload photographs of potato leaves for disease identification and obtain timely results.

## Data Flow Diagram:



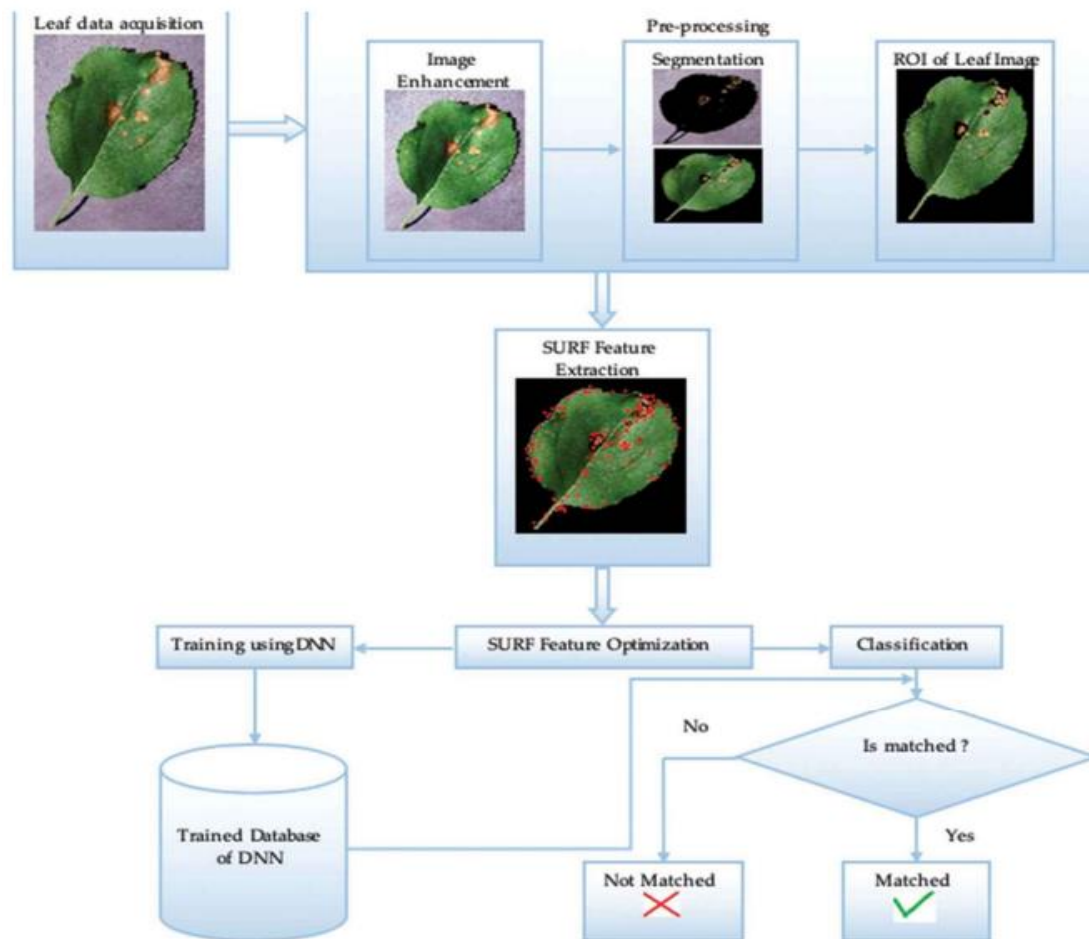
## User Stories:

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

User Type	Functional Requirement (Epic)	User Story Number	User Story / Task	Acceptance criteria	Priority	Release
potatofarmer	Farming	USN-1	I want a potato disease classification system that can quickly identify the specific diseases affecting my crop, so that I can implement timely and targeted treatments to prevent further damage and ensure a successful harvest.	access the disease classification system viaa user-friendly interface, either through a web application or a mobile app.	High	Sprint-1
research scientist	Research On Plants	USN-2	I need a robust potato disease classification tool that can accurately differentiate between various types of diseases affecting potato crops, enablingme to conduct in-depth analyses and develop effective strategies for disease management and prevention.	potato disease classification system should be equipped withan extensive database that covers a wide rangeof potato diseases, including both common and rare occurrences, toprovide comprehensive information for research purposes.	High	Sprint-1
agronomist		USN-3	I require a user-friendly potato disease classification application that can be easily accessed and utilized in the field, allowingme to provide real-time guidance and support to farmers in diagnosing and addressing disease issues, thus improvingoverall crop health and yield.	The potato disease classification system should provide timely andaccurate information on various potato diseases, including their symptoms,causes, and appropriate management strategies, enabling agronomists to make informed decisions and provide effective guidance to farmers.	Medium	Sprint-2
government agricultural officer		USN-4	I am responsible for monitoring and controlling potato diseases at a regional level. I need an advanced potato disease classification system that can efficiently process large volumes of data, enabling me to make informed decisions and implement preventive measures to safeguard the agricultural industry and	It should have the capability to handle a large volume of data and provide real-time analysis, enabling government agricultural officers to make informed decisions and implement timely interventions for disease	Medium	Sprint-1

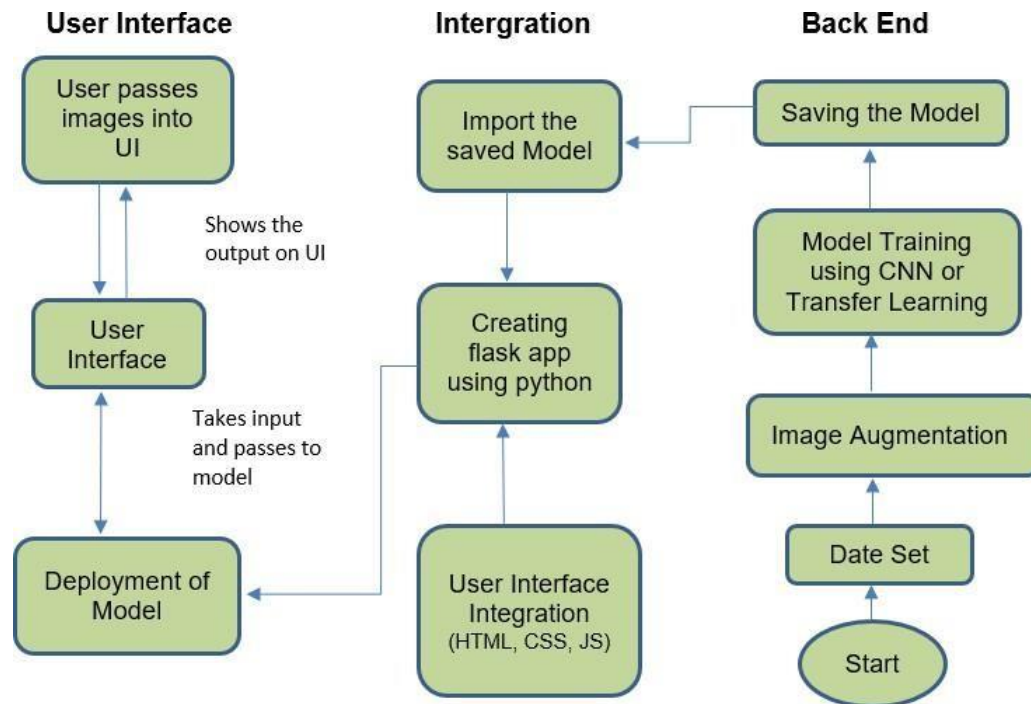
			ensure food security within the region.	control and prevention at a regional or national level.		
student studying plant pathology		USN-5	I am interested in learning about various potato diseases and their classifications. I would benefit from an interactive and educational potato disease classification platform that provides comprehensive information , helping me to understand the complexities of potato diseases and their impact on global agriculture.	The potato disease classification system should provide comprehensive and detailed information about various potato diseases, including their classifications, etiology, symptoms, and management strategies, enabling students to deepen their understanding of plant pathology.	High	Sprint-1

## 5.2 Solution Architecture



## 6. PROJECT PLANNING & SCHEDULING

### 6.1 Technical Architecture



### 6.2 Sprint Planning & Estimation

Use the below template to create product backlog and sprint schedule

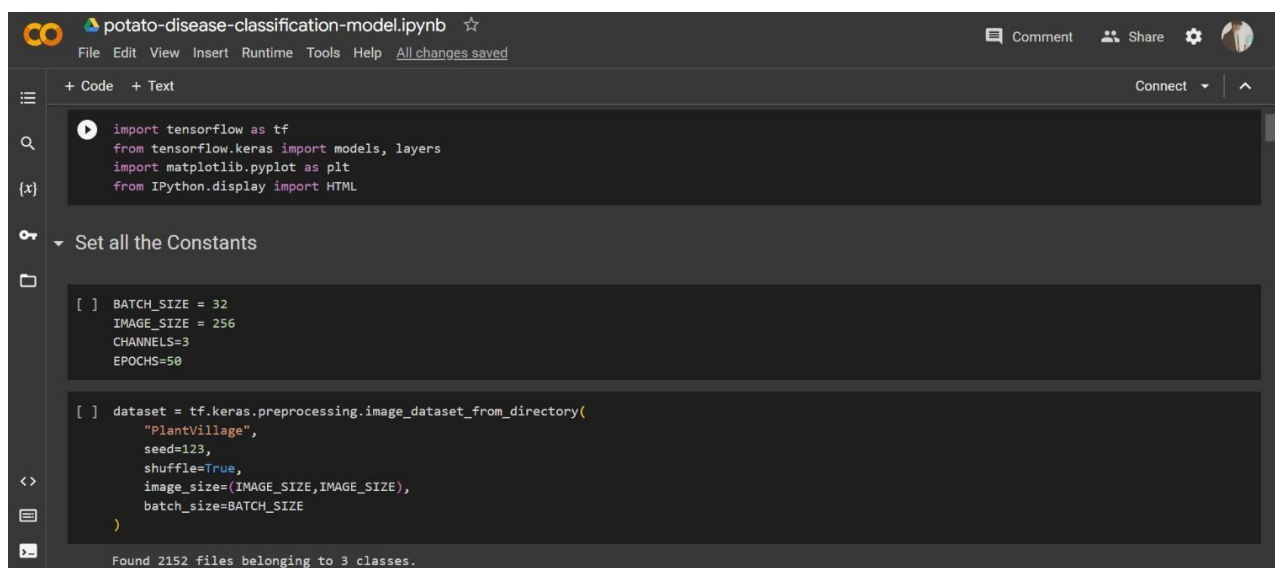
Sprint	Functional Requirement (Epic)	User Story Number	User Story / Task	Story Points	Priority	Team Members
Sprint-1	Disease Classification	USN-1	As a user, I can upload images of potato plants affected by various diseases for classification.	3	High	EDE RENUKA MADHAV
Sprint-1	Disease Classification	USN-2	As a user, I can view the predicted disease class and its probability for the uploaded potato plant image.	2	High	V.P.PRANEETH REDDY
Sprint-2	Disease Classification	USN-3	As a user, I can access historical data and analysis of previously classified potato diseases.	2	Medium	SAI SANTHOSH
Sprint-2	Disease Classification	USN-4	As a user, I can provide feedback on the accuracy of the disease classification for continuous model improvement.	1	Medium	V.P.PRANEETH REDDY
Sprint-3	Disease Classification	USN-5	As a user, I can receive recommendations for disease management strategies based on the classified potato disease.	3	Low	EDE RENUKA MADHAV

## 6.3 Sprint Delivery Schedule

Sprint	Total Story Points	Duration	Sprint Start Date	Sprint End Date (Planned)	Story Points Completed (as on Planned End Date)	Sprint Release Date (Actual)
Sprint-1	20	5 Days	23 Oct 2022	27 Oct 2022	20	27 Oct 2023
Sprint-2	20	5 Days	23 Oct 2022	27 Oct 2022	20	27 Oct 2023
Sprint-3	20	5 Days	23 Oct 2022	27 Oct 2022	20	27 Oct 2023
Sprint-4	20	5 Days	23 Oct 2022	27 Oct 2022	20	27 Oct 2023

## 7. CODING & SOLUTIONING

### Activity 1.1: Importing the libraries



```
potato-disease-classification-model.ipynb ☆
File Edit View Insert Runtime Tools Help All changes saved

+ Code + Text
Connect ^

import tensorflow as tf
from tensorflow.keras import models, layers
import matplotlib.pyplot as plt
from IPython.display import HTML

Set all the Constants

[ ] BATCH_SIZE = 32
    IMAGE_SIZE = 256
    CHANNELS=3
    EPOCHS=50

[ ] dataset = tf.keras.preprocessing.image_dataset_from_directory(
    "PlantVillage",
    seed=123,
    shuffle=True,
    image_size=(IMAGE_SIZE, IMAGE_SIZE),
    batch_size=BATCH_SIZE
)

Found 2152 files belonging to 3 classes.
```

## Activity 2: Data Preparation

As we have understood how the data is, let's pre-process the collected data.

The download data set is not suitable for training the deep learning model. We have to extract class names of the data and let's try to visualise the data with labels.

Extracting class names.

- Visualising the data

### Activity 2.1: Extracting class names

- Let's find the class names of our dataset first. To find the class names of the dataset, we can use `.class_names`.



```
Activity 2: Data Preparation

[ ] class_names = dataset.class_names
class_names

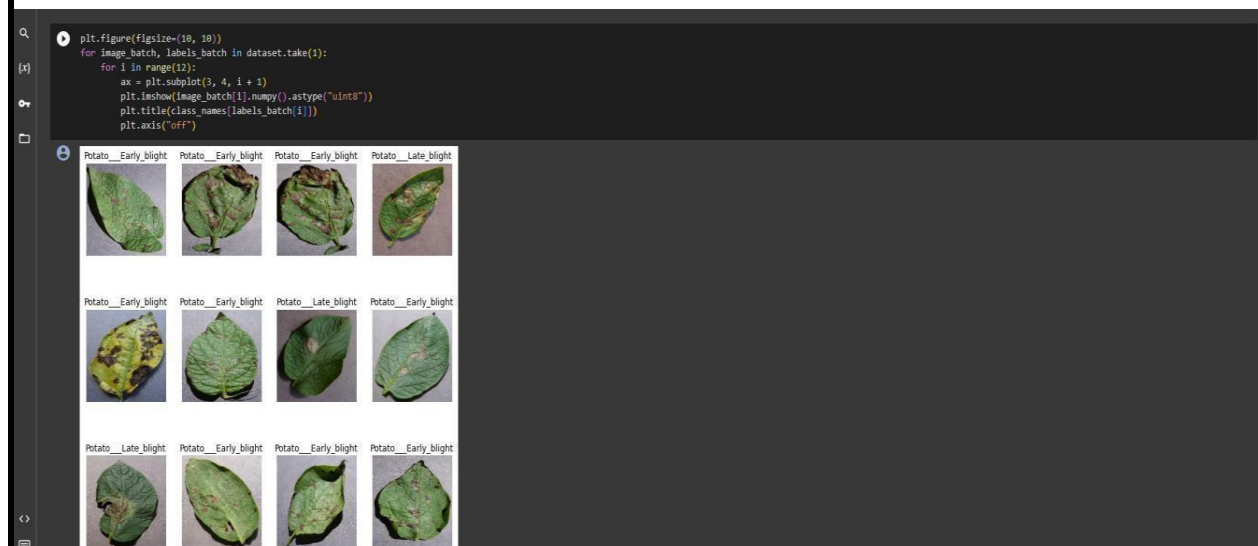
['Potato__Early_blight', 'Potato__Late_blight', 'Potato__healthy']

for image_batch, labels_batch in dataset.take(1):
    print(image_batch.shape)
    print(labels_batch.numpy())

(32, 256, 256, 3)
[1 1 1 0 0 0 0 0 1 1 1 1 0 1 0 1 1 1 0 1 0 1 0 0 1 0 0 1 1 2 0 0]
```



## Activity 2.2: Visualising the data



## **Milestone 3: Exploratory Data Analysis**

### Activity 1: Descriptive statistical

Descriptive analysis is to study the basic features of data with the statistical process. With this describe function we can understand the unique, top and frequent values of categorical features. And we can find mean, std, min, max and percentile values of continuous features. Which are not suitable for our dataset.

### Activity 2: Visual analysis

Visual analysis is the process of using visual representations, such as charts, plots, and graphs, to explore and understand data. It is a way to quickly identify patterns, trends, and outliers in the data, which can help to gain insights and make informed decisions.

This is not needed for our current dataset as it is an image dataset. Visual analysis is mostly used for numerical data.

### Activity 3: Splitting data into train and test and validation sets

Now let's split the Dataset into train, test and validation sets. First split the dataset into train and test sets.

The split will be in 8:1:1 ratio : train : test : validation respectively.

```
Activity 3: Splitting data into train and test and validation sets

[ ] len(dataset)
68

[ ] train_size = 0.8
len(dataset)*train_size
54.400000000000006

[ ] train_ds = dataset.take(54)
len(train_ds)
54

[ ] test_ds = dataset.skip(54)
len(test_ds)
14

[ ] val_size=0.1
len(dataset)*val_size
6.800000000000001

[ ] val_ds = test_ds.take(6)
len(val_ds)
6

[ ] test_ds = test_ds.skip(6)
len(test_ds)
8
```

```
[ ] def get_dataset_partitions_tf(ds, train_split=0.8, val_split=0.1, test_split=0.1, shuffle=True, shuffle_size=10000):
    assert (train_split + test_split + val_split) == 1

    ds_size = len(ds)

    if shuffle:
        ds = ds.shuffle(shuffle_size, seed=12)

    train_size = int(train_split * ds_size)
    val_size = int(val_split * ds_size)

    train_ds = ds.take(train_size)
    val_ds = ds.skip(train_size).take(val_size)
    test_ds = ds.skip(train_size).skip(val_size)

    return train_ds, val_ds, test_ds

train_ds, val_ds, test_ds = get_dataset_partitions_tf(dataset)

[ ] len(train_ds)
54

[ ] len(val_ds)
6

[ ] len(test_ds)
8
```

## Activity 4: Optimizing, Resize, Rescale and Augmentation of the data

```
Activity 4: Optimizing, Resize, Rescale and Augmentation of the data

[ ] train_ds = train_ds.cache().shuffle(1000).prefetch(buffer_size=tf.data.AUTOTUNE)
    val_ds = val_ds.cache().shuffle(1000).prefetch(buffer_size=tf.data.AUTOTUNE)
    test_ds = test_ds.cache().shuffle(1000).prefetch(buffer_size=tf.data.AUTOTUNE)

resize_and_rescale = tf.keras.Sequential([
    layers.experimental.preprocessing.Resizing(IMAGE_SIZE, IMAGE_SIZE),
    layers.experimental.preprocessing.Rescaling(1./255),
])

[ ] data_augmentation = tf.keras.Sequential([
    layers.experimental.preprocessing.RandomFlip("horizontal_and_vertical"),
    layers.experimental.preprocessing.RandomRotation(0.2),
])
```

Prefetching and Caching saves a lot of time in accessing the data from disks. Refer this [link](#) for better understanding.

Resize and Rescale is used to maintain the uniformity among the data.

Augmentation helps you to increase the amount of data and helps your model to learn better.

## Milestone 4: Model Building

### Activity 1: Building a model

Now our data is ready and it's time to build the model.

Here we are going to build our model layer by layer using .Sequential() from

keras. Let's go !

### Model Building

```
input_shape = (BATCH_SIZE, IMAGE_SIZE, IMAGE_SIZE, CHANNELS)
n_classes = 3

model = models.Sequential([
    resize_and_rescale,
    layers.Conv2D(32, kernel_size = (3,3), activation='relu', input_shape=input_shape),
    layers.MaxPooling2D((2, 2)),
    layers.Conv2D(64, kernel_size = (3,3), activation='relu'),
    layers.MaxPooling2D((2, 2)),
    layers.Conv2D(64, kernel_size = (3,3), activation='relu'),
    layers.MaxPooling2D((2, 2)),
    layers.Conv2D(64, (3, 3), activation='relu'),
    layers.MaxPooling2D((2, 2)),
    layers.Conv2D(64, (3, 3), activation='relu'),
    layers.MaxPooling2D((2, 2)),
    layers.Conv2D(64, (3, 3), activation='relu'),
    layers.MaxPooling2D((2, 2)),
    layers.Flatten(),
    layers.Dense(64, activation='relu'),
    layers.Dense(n_classes, activation='softmax'),
])

model.build(input_shape=input_shape)

[ ] model.summary()

Model: "sequential_2"
```

```
model.summary()

Model: "sequential_2"

Layer (type)                Output Shape                Param #
-----
sequential (Sequential)      (32, 256, 256, 3)          0
conv2d (Conv2D)              (32, 254, 254, 32)         896
max_pooling2d (MaxPooling2D) (32, 127, 127, 32)         0
conv2d_1 (Conv2D)            (32, 125, 125, 64)         18496
max_pooling2d_1 (MaxPooling2 (32, 62, 62, 64)           0
conv2d_2 (Conv2D)            (32, 60, 60, 64)          36928
max_pooling2d_2 (MaxPooling2 (32, 30, 30, 64)           0
conv2d_3 (Conv2D)            (32, 28, 28, 64)          36928
max_pooling2d_3 (MaxPooling2 (32, 14, 14, 64)           0
conv2d_4 (Conv2D)            (32, 12, 12, 64)          36928
max_pooling2d_4 (MaxPooling2 (32, 6, 6, 64)            0
conv2d_5 (Conv2D)            (32, 4, 4, 64)            36928
max_pooling2d_5 (MaxPooling2 (32, 2, 2, 64)            0
flatten (Flatten)            (32, 256)                  0
dense (Dense)                (32, 64)                   16448
dense_1 (Dense)              (32, 3)                    195
-----
Total params: 183,747
Trainable params: 183,747
Non-trainable params: 0
```

## Activity 2: Compiling the model

```
Compiling the Model

[ ] model.compile(
    optimizer='adam',
    loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=False),
    metrics=['accuracy']
)
```

## Activity 3: Training the model

```
Training the Model

history = model.fit(
    train_ds,
    batch_size=BATCH_SIZE,
    validation_data=val_ds,
    verbose=1,
    epochs=50,
)

Epoch 1/50 ----- 28s 255ms/step - loss: 0.8882 - accuracy: 0.5341 - val_loss: 0.8462 - val_accuracy: 0.5338
Epoch 2/50 ----- 11s 190ms/step - loss: 0.6831 - accuracy: 0.7396 - val_loss: 0.6225 - val_accuracy: 0.6979
Epoch 3/50 ----- 9s 172ms/step - loss: 0.3647 - accuracy: 0.9483 - val_loss: 0.3865 - val_accuracy: 0.8882
Epoch 4/50 ----- 18s 178ms/step - loss: 0.2776 - accuracy: 0.8999 - val_loss: 0.2782 - val_accuracy: 0.8758
Epoch 5/50 ----- 18s 179ms/step - loss: 0.2448 - accuracy: 0.8953 - val_loss: 0.1857 - val_accuracy: 0.9862
Epoch 6/50 ----- 9s 174ms/step - loss: 0.2828 - accuracy: 0.9144 - val_loss: 0.2387 - val_accuracy: 0.9115
Epoch 7/50 ----- 18s 185ms/step - loss: 0.1751 - accuracy: 0.9288 - val_loss: 0.1854 - val_accuracy: 0.9375
Epoch 8/50 ----- 18s 188ms/step - loss: 0.1436 - accuracy: 0.9444 - val_loss: 0.2273 - val_accuracy: 0.9167
Epoch 9/50 ----- 18s 179ms/step - loss: 0.1128 - accuracy: 0.9583 - val_loss: 0.1425 - val_accuracy: 0.9479
Epoch 10/50 ----- 18s 179ms/step - loss: 0.1218 - accuracy: 0.9549 - val_loss: 0.2318 - val_accuracy: 0.9115
Epoch 11/50 ----- 18s 179ms/step - loss: 0.1524 - accuracy: 0.9398 - val_loss: 0.4774 - val_accuracy: 0.9688
Epoch 12/50 ----- 18s 188ms/step - loss: 0.1862 - accuracy: 0.9578 - val_loss: 0.1767 - val_accuracy: 0.9427
Epoch 13/50 ----- 9s 172ms/step - loss: 0.1299 - accuracy: 0.9549 - val_loss: 0.4929 - val_accuracy: 0.9531
Epoch 14/50 ----- 9s 169ms/step - loss: 0.4971 - accuracy: 0.9681 - val_loss: 0.1238 - val_accuracy: 0.9531
Epoch 15/50 ----- 9s 173ms/step - loss: 0.4967 - accuracy: 0.9609 - val_loss: 0.4884 - val_accuracy: 0.9635
Epoch 16/50 ----- 9s 172ms/step - loss: 0.4764 - accuracy: 0.9676 - val_loss: 0.1125 - val_accuracy: 0.9531
Epoch 17/50 ----- 9s 174ms/step - loss: 0.1157 - accuracy: 0.9543 - val_loss: 0.2288 - val_accuracy: 0.9219
Epoch 18/50 ----- 18s 178ms/step - loss: 0.4947 - accuracy: 0.9659 - val_loss: 0.1852 - val_accuracy: 0.9271
Epoch 19/50 ----- 9s 174ms/step - loss: 0.4737 - accuracy: 0.9711 - val_loss: 0.4923 - val_accuracy: 0.9583
Epoch 20/50 ----- 9s 173ms/step - loss: 0.4518 - accuracy: 0.9815 - val_loss: 0.4678 - val_accuracy: 0.9688
Epoch 21/50 ----- 9s 172ms/step - loss: 0.4873 - accuracy: 0.9826 - val_loss: 0.4516 - val_accuracy: 0.9748
Epoch 22/50 ----- 9s 173ms/step - loss: 0.4518 - accuracy: 0.9883 - val_loss: 0.3843 - val_accuracy: 0.8958
Epoch 23/50 ----- 9s 178ms/step - loss: 0.4518 - accuracy: 0.9792 - val_loss: 0.2573 - val_accuracy: 0.9882
```

The verbose option specifies that you want to display detailed processing information on your screen

If your laptop is functioning slow for 100 epochs, then you can try Google Colab Notebooks.

You just have to follow the below steps:

- Go to GoogleColab and create a new notebook.
- Click on Files and mount GoogleDrive.
- Upload your dataset onto GoogleDrive.
- After uploading the dataset go to runtime and click on “Change runtime type”.
- Select “gpu” and choose “standard” and click on save.
- Booyah! Now you can run the whole code.

## Milestone 5: Model Deployment

### Activity 1: Model Evaluation

As we have the record of accuracy stored in the history variable. We can try visualising the performance of our model.

```
Model Deployment
[ ] scores = model.evaluate(test_ds)
8/8 [=====] - 1s 14ms/step - loss: 0.0063 - accuracy: 1.0000
[ ] scores
[0.00625185994431765, 1.0]
[ ] history
<tensorflow.python.keras.callbacks.History at 0x7d88437e50>
[ ] history.params
{'verbose': 1, 'epochs': 50, 'steps': 54}
[ ] history.history.keys()
dict_keys(['loss', 'accuracy', 'val_loss', 'val_accuracy'])
[ ] type(history.history['loss'])
list
[ ] len(history.history['loss'])
50
[ ] history.history['loss'][:5] # show loss for first 5 epochs
[0.00818462392586769,
0.0031139228820088,
0.0044925388794893,
0.02788617189825879,
0.04488397999206652]
```

```
▶ history.history['accuracy']
```

```
↳ [0.5214120149612427,  
    0.7685185074806213,  
    0.8449074029922485,  
    0.8894675970077515,  
    0.9224537014961243,  
    0.9068287014961243,  
    0.9259259104728699,  
    0.9259259104728699,  
    0.9357638955116272,  
    0.9461805820465088,  
    0.9560185074806213,  
    0.9502314925193787,  
    0.9542824029922485,  
    0.9502314925193787,  
    0.9589120149612427,  
    0.9554398059844971,  
    0.9513888955116272,  
    0.9600694179534912,  
    0.9554398059844971,  
    0.9618055820465088,  
    0.9728009104728699,  
    0.9716435074806213,  
    0.9629629850387573,  
    0.9641203880310059,  
    0.9710648059844971,  
    0.9577546119689941,  
    0.9751157164573669,  
    0.9820601940155029,  
    0.9722222089767456,  
    0.9826388955116272,  
    0.9826388955116272,  
    0.9803240895271301,  
    0.9785879850387573,  
    0.9768518805503845,  
    0.9774305820465088,  
    0.9768518805503845,  
    0.984375,  
    0.9780092835426331,  
    0.9855324029922485,  
    0.9707452000000000]
```

history.history['accuracy'] contains the track of accuracies achieved while you were training your model.

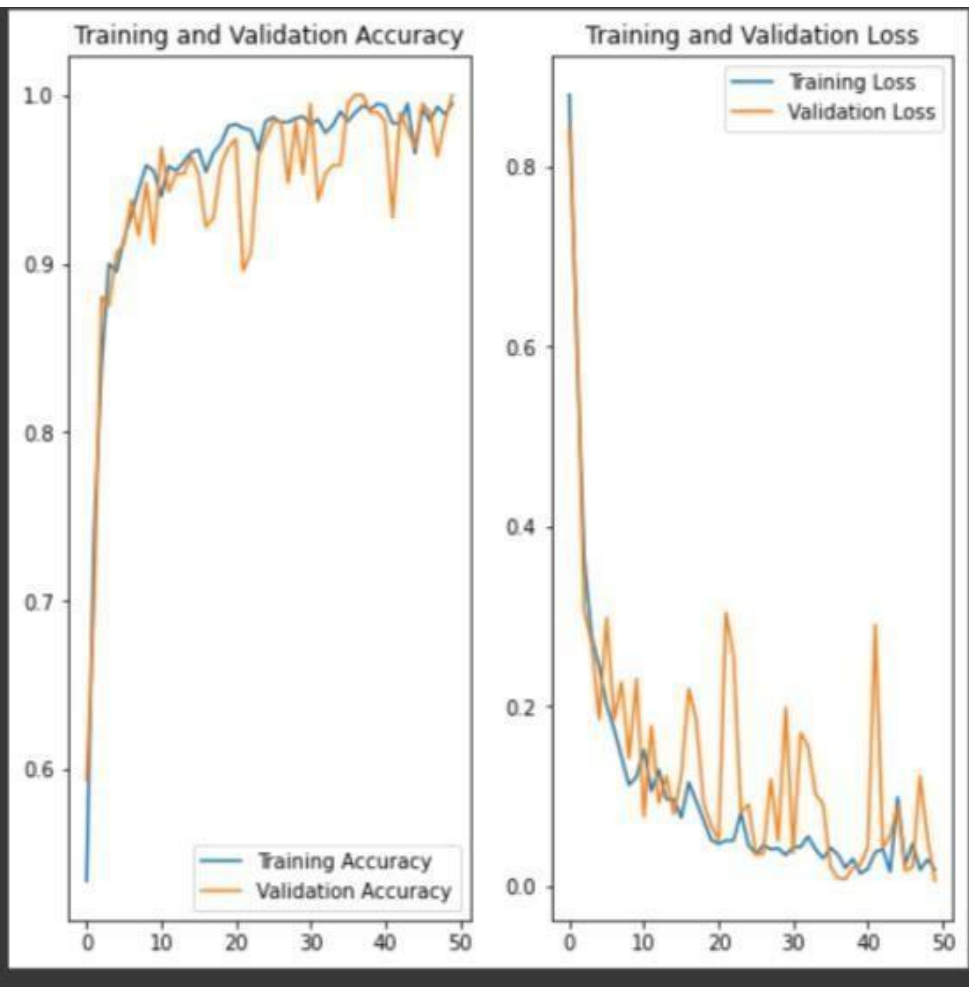
```
acc = history.history['accuracy']
val_acc = history.history['val_accuracy']

loss = history.history['loss']
val_loss = history.history['val_loss']

[ ] plt.figure(figsize=(8, 8))
plt.subplot(1, 2, 1)
plt.plot(range(EPOCHS), acc, label='Training Accuracy')
plt.plot(range(EPOCHS), val_acc, label='Validation Accuracy')
plt.legend(loc='lower right')
plt.title('Training and Validation Accuracy')

plt.subplot(1, 2, 2)
plt.plot(range(EPOCHS), loss, label='Training Loss')
plt.plot(range(EPOCHS), val_loss, label='Validation Loss')
plt.legend(loc='upper right')
plt.title('Training and Validation Loss')
plt.show()
```

The above code plots two graphs, one graph b/w Training and Validation Accuracy and the other graphs b/w Training and Validation Loss.





## Activity 2: Model Evaluation with Visualisation

First we'll select one image from the test dataset and try to predict using the trained model. Then we'll try to print the actual class and predicted class of that particular image.

### Run prediction on a sample image

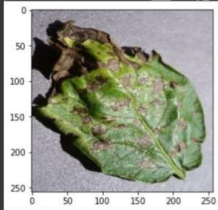
```
import numpy as np
for images_batch, labels_batch in test_ds.take(1):

    first_image = images_batch[0].numpy().astype('uint8')
    first_label = labels_batch[0].numpy()

    print("first image to predict")
    plt.imshow(first_image)
    print("actual label:", class_names[first_label])

    batch_prediction = model.predict(images_batch)
    print("predicted label:", class_names[np.argmax(batch_prediction[0])])
```

```
first image to predict
actual label: Potato_Early_blight
predicted label: Potato_Early_blight
```



Each time you run this code, you'll get a different image to predict.

Now let us write a function for prediction, which returns the predicted class and the confidence of its prediction.

Write a function for inference

```
] def predict(model, img):  
    img_array = tf.keras.preprocessing.image.img_to_array(images[i].numpy())  
    img_array = tf.expand_dims(img_array, 0)  
  
    predictions = model.predict(img_array)  
  
    predicted_class = class_names[np.argmax(predictions[0])]  
    confidence = round(100 * (np.max(predictions[0])), 2)  
    return predicted_class, confidence
```

Actual: Potato\_\_healthy,  
Predicted: Potato\_\_healthy.  
Confidence: 98.23%



Actual: Potato\_\_Early\_blight,  
Predicted: Potato\_\_Early\_blight.  
Confidence: 99.51%



Actual: Potato\_\_Early\_blight,  
Predicted: Potato\_\_Early\_blight.  
Confidence: 99.91%



Actual: Potato\_\_Early\_blight,  
Predicted: Potato\_\_Early\_blight.  
Confidence: 89.89%



Actual: Potato\_\_Late\_blight,  
Predicted: Potato\_\_Late\_blight.  
Confidence: 100.0%



Actual: Potato\_\_Late\_blight,  
Predicted: Potato\_\_Late\_blight.  
Confidence: 97.07%



Actual: Potato\_\_Late\_blight,  
Predicted: Potato\_\_Late\_blight.  
Confidence: 99.64%



Actual: Potato\_\_Late\_blight,  
Predicted: Potato\_\_Late\_blight.  
Confidence: 99.98%



Actual: Potato\_\_Early\_blight,  
Predicted: Potato\_\_Early\_blight.  
Confidence: 99.94%



### Activity 3: Save the model

```
model.save(f"..../Models/potato.h5")
```

This will allow us to save the .h5 format of the model.

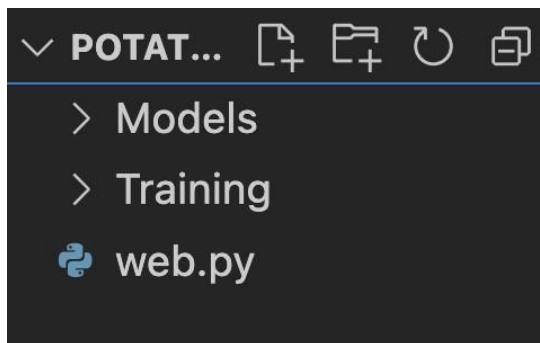
## Activity 4: Integrate with Web Framework

In this section, we will be building a web application that is integrated to the model we built.

We will be using the streamlit package for our website development.

Streamlit is a free and open-source framework to rapidly build and share beautiful machine learning and data science web apps.

### Activity 4.1: Create a web.py file and import necessary packages:



```
import streamlit as st
import tensorflow as tf
from PIL import Image, ImageOps
import numpy as np
from io import BytesIO
```

### Activity 4.2: Defining class names and loading the model:

```
st.cache(allow_output_mutation=True)
CLASS_NAMES = ["Early Blight" , "Late Blight" , "Healthy"]
def load_model():
    model=tf.keras.models.load_model('./Models/potato.h5')
    return model
model = load_model()
```

### Activity 4.3: Accepting the input from user and prediction

```
st.write("""# Potato Leaf Disease Classification""")

file = st.file_uploader("Please upload an brain scan file", type=["jpg", "png"])

def import_and_predict(image_data, model):
    size = (255,255)
    image = ImageOps.fit(image_data, size, Image.ANTIALIAS)
    image = np.asarray(image)
    img_reshape = np.expand_dims(image,0)
    prediction = model.predict(img_reshape)
    return prediction
```

```
if file is None:
    st.text("Please upload an image file")
else:
    image = Image.open(file)
    st.image(image, use_column_width=True)
    predictions = import_and_predict(image, model)
    index = np.argmax(predictions[0])
    predicted_class = CLASS_NAMES[index]
    confidence = np.max(predictions[0])
    st.write(predicted_class)
    st.write(confidence)
    print(
        "This image most likely belongs to {} with a {:.2f} percent confidence."
        .format(CLASS_NAMES[np.argmax(confidence)], 100 * np.max(confidence))
    )
```



You can now view your Streamlit app in your browser.

Local URL: `http://localhost:8501`

Network URL: `http://192.168.1.11:8501`

For better performance, install the Watchdog module:

```
$ xcode-select --install
```

```
$ pip install watchdog
```

## **Milestone 5: Building Flask application**

After the model is built, we will be integrating it to a web application so that normal users can also use it. The new users need to initially register in the portal. After registration users can login to browse the images to detect the condition of potatoes. Activity 1: Build a python application

Step 1: Load the required packages.

```
from tensorflow.keras.models import load_model
from tensorflow.keras.preprocessing import image
from flask import Flask, render_template, request
import os
import numpy as np
import pickle
```

Step 2: Initialize the flask app, load the model and configure the html pages Instance of Flask is created and the model is loaded using load\_model from keras.

```
app = Flask(__name__)
model = load_model(r"Potato_model.h5", compile = False)

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

@app.route('/predict', methods = ['GET', 'POST'])
```

Step 3: Pre-process the frame and run

```

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 =(240,240))
        x = image.img_to_array(img)
        x = np.expand_dims(x,axis = 0)
        pred =np.argmax(model.predict(x),axis=1)
        index =['EarlyBlight','Healthy','LateBlight']
        text="The potato is : "+index[pred[0]]
    return text

if __name__=='__main__':
    app.run(debug=True)

```

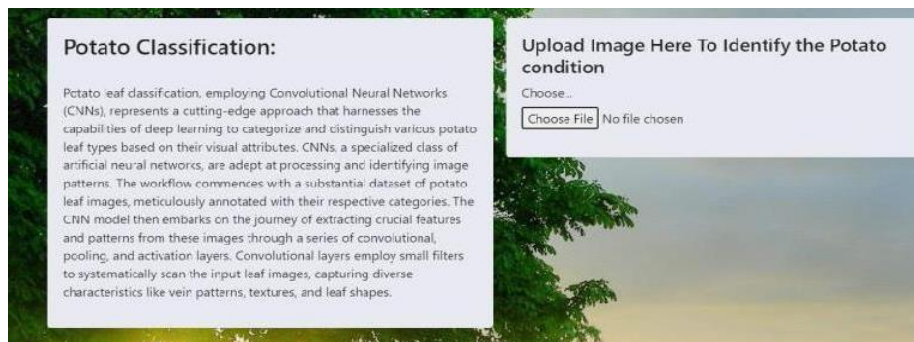
Run the flask application using the run method. By default, the flask runs on port 5000. If the port is to be changed, an argument can be passed and the port can be modified.

### Activity 2: Build the HTML page and execute

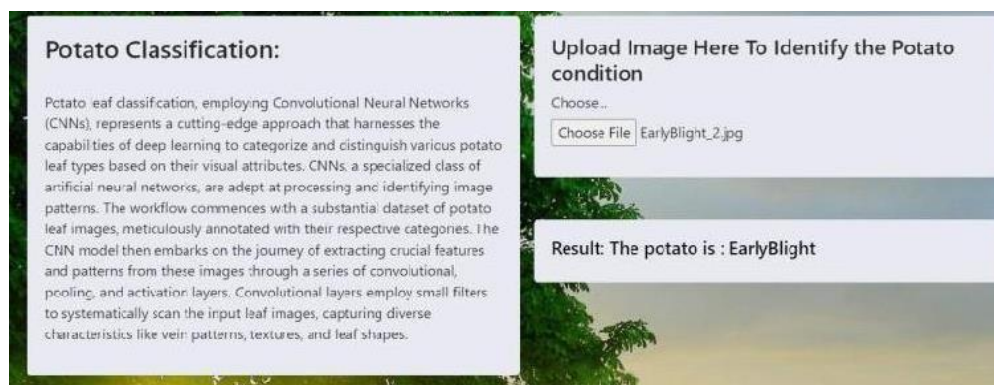
Build the UI where a home page will have details about the application and prediction page where a user is allowed to browse an image and get the predictions of images. Step 1: Run the application

In the anaconda prompt, navigate to the folder in which the flask app is present. When the python file is executed, the localhost is activated on port 5000 and can be accessed through it.

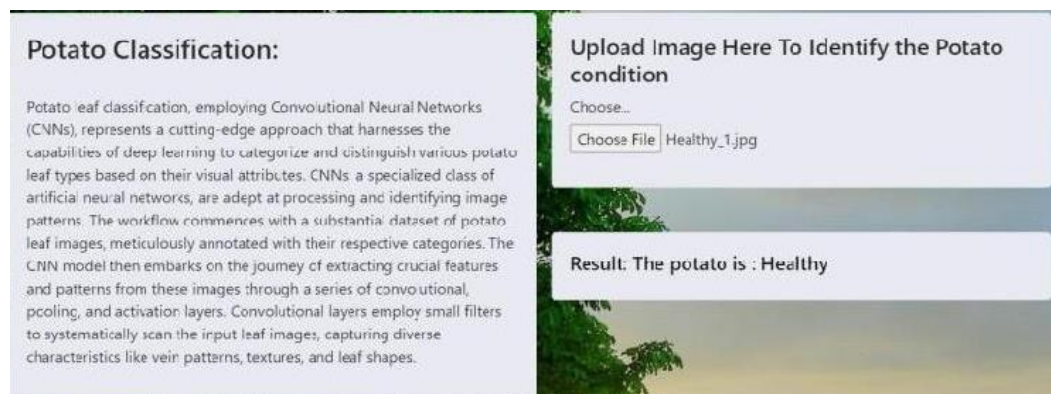
Step 2: Open the browser and navigate to localhost:5000 to check your application The home page looks like this:



Click on choose the image to upload and select an image to be classified.Input-1:

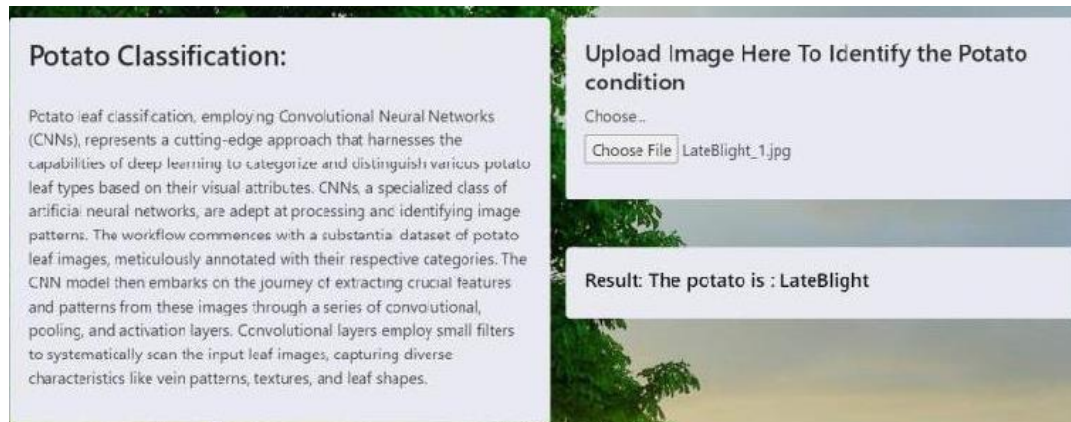


Input-2:





### Input-3:



## **8. ADVANTAGES & DISADVANTAGES**

### **Advantages:**

**Early Disease Detection:** The technology makes it easier to identify potato illnesses early, enabling farmers to respond quickly to minimise crop damage. Higher crop yields and fewer financial losses may result from this.

**Accurate Classification:** When taught and maintained appropriately, deep learning models can yield remarkably accurate illness classification outcomes. Making educated judgements about illness prevention and treatment depends on this accuracy.

**User-Friendly:** The system's user interface and feedback mechanisms enable a broad spectrum of users, even those with low technical competence, to utilise it. It gives farmers the ability to control their crops with cutting-edge technologies.

**Continuous Improvement:** The accuracy and robustness of the model can be continuously enhanced by utilising user feedback. As more data and feedback are collected, the system becomes increasingly reliable and effective.

**Scalability:** The system can be scaled to accommodate a growing number of users and adapt to the specific needs of different agricultural regions and crops.

## **Disadvantages:**

**Data Requirements:** Obtaining big and diverse datasets for training is a challenge for deep learning models, particularly for rare or regionally unique diseases. Such datasets can be expensive and time-consuming to obtain and annotate.

**Model Complexity:** Deep learning models, particularly CNNs, can be computationally intensive and may require high-performance hardware for training and inference. This could be a barrier for small-scale farmers with limited resources.

**Interpretability:** Even while deep learning models are capable of great accuracy, they are frequently regarded as "black boxes," which makes it challenging to provide an explanation for a particular categorization conclusion. The difficulty of interpreting a model might affect user acceptability and trust.

**Maintenance and Updates:** Continuous model maintenance and updates are required to keep the system effective. This includes retraining the model with new data and addressing evolving disease patterns.

**Privacy and Security:** Handling user-generated data, including images, requires robust security measures to protect user privacy. Information misuse or data breaches may have moral and legal ramifications.

**Dependency on Technology:** The technology infrastructure—which includes internet connection and suitable devices—is necessary for the system to function. The effectiveness of the system can be compromised in areas with inadequate connectivity or restricted device access.

**False Positives and Negatives:** Like all machine learning models, false positives and false negatives can occur. Misclassifications can lead to unnecessary pesticide use or failure to address actual diseases, potentially causing harm or economic losses.

## **9. CONCLUSION**

An encouraging approach for agriculture is the creation of a deep learning system, specifically Convolutional Neural Networks (CNNs), for the categorization of potato diseases. The prompt and precise identification of illnesses in potato plants by this technology allows for early detection, which can result in higher crop yields, lower financial losses, and improved food security. A wide range of users, including non-technical people, can utilise this technology because of its user-friendly interface and feedback systems. To guarantee the system's efficacy, however, issues with data quality, model complexity, interpretability, and continuing maintenance must be properly handled. As the project moves forward, accuracy and relevance must be maintained by regular model changes and the incorporation of user feedback. To sum up, this project has the potential to greatly help farmers and the agricultural sector as a whole when implemented responsibly and thoughtfully.

## **10. FUTURE SCOPE**

**Expanded Crop Coverage:** The underlying deep learning architecture can be extended to classify diseases in numerous other crops, while the current focus is on potato diseases. Numerous agricultural techniques can profit from its scalability, which also helps with crop health management.

**AI-Driven Precision Agriculture:** By integrating this system with more comprehensive precision agriculture techniques, resource allocation may be optimised, pesticide use can be decreased, and sustainable farming can be encouraged. The deep learning model is able to adjust to particular crop varieties and local conditions.

**IoT Integration:** The capabilities of the system can be improved by integration with Internet of Things (IoT) devices, such as automated irrigation systems and sensors. By gathering data in real time, these gadgets can enhance illness detection and prevention even further.

**AI-Enhanced Recommendations:** In addition to categorising diseases, the system has the capacity to develop recommendations for disease management tactics, such as the choice of suitable pesticides, organic treatments, and preventive measures.

**Mobile Apps and Accessibility:** The creation of specialised mobile applications can improve accessibility and make it possible for farmers in isolated locations to use the technology for managing and diagnosing diseases.

## **11. APPENDIX**

The drive link for the ipynb file with the potato classification model using Resnet\_50

[https://drive.google.com/file/d/16ewXg9IGIHdGm2UILM35V22\\_WisV6nnk/view?usp=sharing](https://drive.google.com/file/d/16ewXg9IGIHdGm2UILM35V22_WisV6nnk/view?usp=sharing)

The github link for the ipynb file

[https://github.com/smartinternz02/SI-GuidedProject-600905-1697560237/blob/main/DEVELOPMENT%20PHASE/potato\\_diseases\\_classification\\_model.ipynb](https://github.com/smartinternz02/SI-GuidedProject-600905-1697560237/blob/main/DEVELOPMENT%20PHASE/potato_diseases_classification_model.ipynb)

The github link for the flask source code

<https://github.com/smartinternz02/SI-GuidedProject-600905-1697560237/blob/main/DEVELOPMENT%20PHASE/Flask/app.py>

Project Demo Video link

<https://drive.google.com/file/d/152PUd2KeQgdyN0SAL0429TLy4lC948a2/view?usp=sharing>