Sri Lanka Institute of Information Technology



Year 4, Semester I, 2024 Assignment

Potato Disease Classification Using CNN Architectures on the Plant Village Dataset

Deep Learning (SE4050)

BSc (Hons) in Information Technology Specializing in Software

Engineering

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GitHub Link

IT21251900/DL-Project (github.com)

YouTube Presentation Link

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Introduction

Problem Statement

Potato crops are a crucial agricultural commodity, feeding millions worldwide. However, they are susceptible to various diseases, which can drastically affect yield and quality. Among the most notorious are Early Blight and Late Blight, both of which can cause significant losses if left unchecked. Identifying these diseases early and accurately is essential for effective intervention and treatment.

In this project, we tackle the problem of classifying potato leaf diseases using image data and deep learning techniques. Specifically, we employ supervised learning methods to classify leaf images into categories such as "Healthy," "Early Blight," and "Late Blight." This project leverages the **Plant Village** dataset, which is widely recognized for plant disease classification research, offering a large and diverse set of annotated images. By building robust deep learning models, the system aims to automate the detection of these diseases from images of potato leaves.

Objective

The goal of this project is to develop a machine learning solution to classify potato leaf diseases. Specifically, we aim to:

- Develop and Compare CNN Models: Train and evaluate four distinct
 Convolutional Neural Networks (CNNs) for accuracy and generalization.
- Provide a Disease Detection API: Deploy a FastAPI backend to process image uploads and provide real-time disease predictions.
- Deploy a React Frontend: Build a user-friendly React interface for easy image uploads and viewing of classification results.

Supervised Learning Approach

Supervised learning was chosen for this project because the problem we are tackling involves **labeled data**, where the input images are associated with known categories (e.g., healthy or diseased classes). In supervised learning, the model is trained to learn the mapping between the input data and its corresponding labels, allowing it to predict the correct class when new, unseen data is presented. This fits well with our objective, which is to classify images into predefined categories based on their features.

The key reasons for choosing supervised learning are:

- Labeled Dataset: Our dataset comes with ground truth labels, meaning each input image is associated with a known class. Supervised learning algorithms are specifically designed to work with labeled data, learning from examples and improving their performance over time.
- 2. High Predictive Accuracy: Supervised learning techniques, especially convolutional neural networks (CNNs), have been widely used in image classification tasks and have shown superior performance. CNNs are highly effective in automatically learning spatial hierarchies of features from images, which are crucial in our classification task.
- 3. **Clear Performance Metrics**: Since supervised learning involves labeled data, it is easier to evaluate the model's performance through established metrics like accuracy, precision, recall, and F1-score. These metrics provide a clear understanding of how well the model generalizes to unseen data.
- 4. **Control Over Training Process**: Supervised learning allows fine-tuning of the model by adjusting hyperparameters and providing more control over the training process. This flexibility ensures that we can iteratively improve the model's performance by adjusting the architecture, learning rate, and other factors.

Background Information on Supervised Learning and CNN

Supervised learning is a machine learning approach where models learn from labeled datasets. In the context of this project, the labels represent different categories of plant diseases, and the input data consists of images. CNNs, a type of deep learning model particularly well-suited for image classification tasks, are employed in this project. CNNs automatically extract features from images, learning patterns such as edges, textures, and colors, which are crucial for identifying plant diseases.

CNNs consist of several layers:

- **Convolutional layers**: Extract features using convolution operations.
- **Pooling layers**: Reduce the spatial dimensions of the data.
- **Fully connected layers**: Combine features to classify images into the correct category.

This architecture is ideal for the classification of plant disease images, where intricate patterns in leaf textures and colors need to be identified to distinguish between healthy plants and various diseases.

Scope

- **Data Preprocessing:** Image cleaning, resizing, normalizing, and augmentation using the Plant Village dataset.
- **Model Development:** Exploring four CNN architectures with techniques like batch normalization, dropout, and transfer learning.
- **API Development:** FastAPI will be used to deploy trained models for real-time image classification.
- **Frontend Development:** A React-based application will allow users to upload images and receive predictions.

Dataset Overview

Dataset Description

The dataset used for this project is the **Plant Village Dataset**, specifically focusing on potato diseases. This dataset is well-suited for tasks like image classification in plant pathology, as it contains high-quality images of plant leaves, labeled for various diseases.

- **Dataset Name:** Plant Village Dataset (Potato Disease Subset).
- **Size:** The subset contains 2152 **images** across **3 classes**. These classes include different plant diseases, but for the purpose of this project, we focus on images of **potato leaves** with three specific labels:
 - 1. Healthy
 - 2. Early Blight
 - 3. Late Blight

Each image represents a close-up shot of a potato leaf, either healthy or affected by one of these two diseases. The large number of images allows for robust training, testing, and validation of the models, reducing the chances of overfitting.

- **Source:** The dataset is publicly available on **Kaggle**. It is one of the most widely used datasets in plant disease classification tasks due to its diversity and quality.
- **Image Format:** All images are colored and provided in JPEG format. They vary in size and quality, necessitating preprocessing steps like resizing and normalization.

Classes of Interest:

- Healthy: These images depict healthy potato leaves without any visible signs of disease.
- Early Blight: Caused by the fungus Alternaria solani, Early Blight appears as small dark brown or black spots on the leaves, often surrounded by a yellow halo.

o **Late Blight:** Caused by the oomycete *Phytophthora infestans*, Late Blight manifests as large dark patches on leaves, with a water-soaked appearance. This disease is particularly devastating and can destroy crops rapidly if left unchecked.

• Attributes:

- o **Input**: Images of size 256x256, with 3 color channels (RGB).
- Output: Labels corresponding to disease types.

Feature Selection and Preprocessing Techniques

Feature selection for image data involves the extraction of visual features such as textures, shapes, and colors. In CNNs, this is done automatically by the convolutional layers. The preprocessing steps we applied to the dataset include:

- **Resizing**: All images were resized to a uniform size of 256x256 pixels.
- **Normalization**: Pixel values were scaled to the range [0, 1] to improve model convergence during training.
- **Data Augmentation**: Techniques such as random rotation, flipping, and zooming were applied to artificially expand the dataset and improve model generalization.

Challenges

Despite the benefits of the Plant Village dataset, several challenges arose during model training:

- Class Imbalance: The "Healthy" class is overrepresented, leading to biased models. This was addressed with data augmentation and a tuned loss function.
- Image Quality: Variations in lighting and resolution affected consistency.

 Normalization and preprocessing were applied to mitigate these issues.
- **Leaf Occlusion:** Some images had partially occluded leaves or overlapping patterns, complicating the identification of disease symptoms.

Model Architectures

Model 1

This model uses a **Sequential CNN** architecture with six Conv2D layers, which is effective for image classification tasks. CNNs are chosen for their ability to automatically detect important features such as edges, textures, and patterns in images.

- Layers Chosen: Multiple Conv2D layers followed by MaxPooling layers enable downsampling and feature extraction. The model flattens before the Dense layers to connect to the final classification task.
- Activation Functions: ReLU is used for all convolutional layers because of its
 ability to mitigate the vanishing gradient problem. The Softmax function is
 applied at the output layer for multi-class classification.

```
### TR21302862 - Model Architecture

We use CRX coupled with a Softmax activation in the corpor layer. We also add the initial layers for reasizing, commission and Data Augmentation.

First Conv20 Layer

A 2D convolution layer with 25 filters, such of size 201. The ReLU activation function remoduces non-linearity, This layer is reagonable for learning basic image features like adjust and stances. The layer is reagonable for learning basic image features like adjust and stances. The layer layer is reagonable for learning basic image features like adjusts of the proper basic image features and adjusts of the model of the proper basic image features. But adjusts on the previous one. Using 64 filters allows the network to capture a higher vertey of features.

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Fourth ConvOID layer with 64 filters, using the same kernel size (3x3). As the model deeports, it extracts higher-level features such as shapes and extracts.

Fifth ConvOID layer, gain with 64 filters, using the same kernel size (3x3). As the model deeports, it extracts higher-level features such as shapes and extracts.

Fifth ConvOID layer, gain with 64 filters, which allows the model to capture increasingly complex features.

Fifth ConvOID layer, gain with 64 filters, which allows the model and capture elevance.

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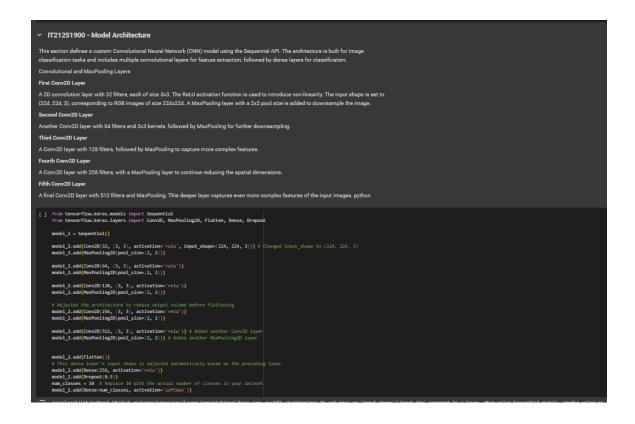
First ConvOID layer, gain with 64 filters, the first continues to deepen the model and capture elevance.

First ConvOID layer, gain with 64 filters, the first continues to deepen the model and capture elevance.

First ConvOID layer, gain with 64 filters, the first continues to the filters of the filters of the filters of
```

This model is a more complex CNN with five Conv2D layers, using **Adam** as the optimizer for faster convergence.

- Layers Chosen: With increasing filter sizes, each convolutional layer captures
 progressively more complex patterns. MaxPooling downscales feature maps to
 reduce computational complexity.
- Activation Functions: Again, ReLU for hidden layers and Softmax for the final output layer due to the classification task.



This model includes Batch Normalization, which stabilizes learning and improves performance in deep architectures.

- **Layers Chosen**: Conv2D layers capture image features, and Batch Normalization ensures stable and faster training.
- Activation Functions: ReLU for hidden layers, Softmax for output.



This model focuses on generalization using four convolutional layers and dropout to prevent overfitting.

- **Layers Chosen**: Each Conv2D block is followed by MaxPooling, batch normalization, and Dropout for regularization.
- Activation Functions: ReLU and Softmax for classification.

```
V IT21360428 - Model Architecture
 This code defines a convolutional neural network (CNN) using Keras for image classification.
After flattening the feature maps, a fully connected layer with dropout is used for regularization, followed by an output layer with softma activation for multi-class classification.
 The model is designed to preprocess images by resizing and rescaling pixel values to the range [0, 1].
[ ] # Define the image size and number of channels
IMAGE_SIZE = 256
CHANNELS = 3
         # You need to manually define class names based on your dataset directory
class_names = ['healthy', 'diseased_class1', 'diseased_class2', 'diseased_class3'] # Modify this according to your dataset
          # Define input shape
input_shape = (IMAGE_SIZE, IMAGE_SIZE, CHANNELS)
          # Define the resize and rescale layers
resize_and_rescale = tf.keras.Sequential([
tf.keras.layers.Resizing[IMAG_SIZE, IMAGE_SIZE),
tf.keras.layers.Rescaling(1./255),
}
               Define the model

del_4 = models.Sequential([
    resize_and_rescale, # Use the resize and rescale layer
                    First Convolutional Block

syers.Conv20(32, (3, 3), activation='relu', padding='same', input_shape=input_shape),

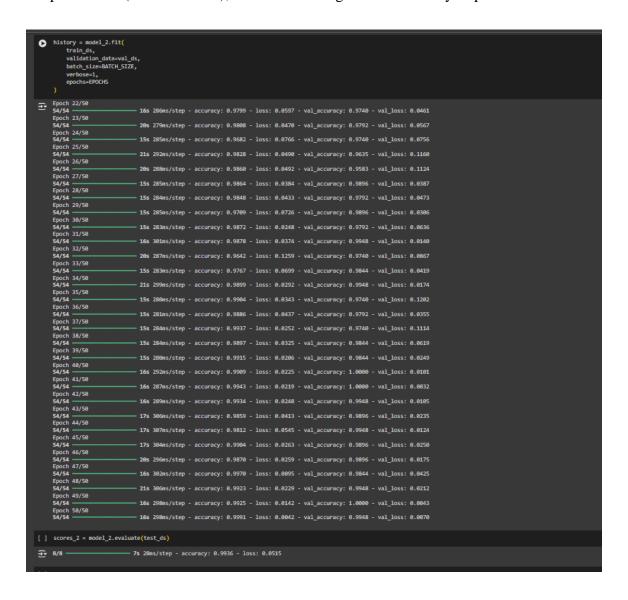
syers.BatcNormalization(),

syers.MaxPooling2D((2, 2)),
                  # Second Convolutional Block
layers.Conv2D(64, (3, 3), activation='relu', padding='same'),
layers.BatchWornalization(),
layers.MaxPooling2D((2, 2)),
                  % Third Convolutional Block
layers.Conv20(128, (3, 3), activation='relu', padding='same'),
layers.BatchWormalization(),
layers.MaxPooling20((2, 2)),
                  # Fourth Convolutional Block
layers.Conv2D(256, 3, 3), activation='relu', padding='same'),
layers.Batkhormalization(),
layers.MaxPooling2D((2, 2)),
                 # Flatten the data
layers.Flatten(),
                 # Dense layers with Dropout for regularization
layers.Dense(512, activation='relu'),
layers.Dense(10.5),
layers.Dense(n_classes, activation='softmax'), # Output layer with the correct number of classes
              Build and summarize the model 
kdel_4.build(input_shape=(None, IMAGE_SIZE, IMAGE_SIZE, CHANNELS)) ## Build the model with the correct input shap
```

Results

- **Test Accuracy**: Achieved a test accuracy of **97.63%** after 50 epochs...
- Performance: The model showed strong performance due to multiple Conv2D layers
 that captured diverse image features effectively. Loss decreased steadily during
 training, indicating good model convergence.
- **Challenges**: There was a noticeable improvement in validation accuracy across epochs, with minimal overfitting thanks to the inclusion of data augmentation.

- Test Accuracy: Achieved a test accuracy of 99.36% after 50 epochs...
- Performance: Model 2 performed the best among the four models, likely due to its
 deeper architecture with five Conv2D layers. The increasing filter sizes and more
 complex feature extraction likely contributed to this high accuracy.
- **Challenges**: The model required a longer training time due to the larger number of parameters (over 4 million), but this led to significant accuracy improvements.



- **Test Accuracy**: Achieved a test accuracy of **76.42%** after 50 epochs.
- **Performance**: Although the architecture was simpler than Model 2, the inclusion of Batch Normalization improved training stability. However, the performance was lower compared to Model 2, likely due to underfitting or insufficient feature extraction in the deeper layers.
- Challenges: The model struggled to generalize on the validation set, suggesting that
 additional tuning or a more complex architecture might be required for better
 performance.

```
[ ] history = model_3.fit(
    train_ds,
    batch_size=BATCH_SIZE,
    validation_data=val_ds
    verbose=1,
    epochs=EPOCHS,
Epoch 1/58

58 549ms/step - accuracy: 0.8021 - loss: 4.8974 - val_accuracy: 0.4479 - val_loss: 40.0144

Epoch 1/50

50 549ms/step - accuracy: 0.9085 - loss: 0.5371 - val_accuracy: 0.4479 - val_loss: 68.2467
      34/94 195 39385/5tep - accuracy: 8.9885 - 1055: 8.53/4 - Val_accuracy: 8.4479 - Val_loss: 88.8459
54/54 195 3558s/step - accuracy: 8.9119 - 1055: 8.3546 - Val_accuracy: 8.4479 - Val_loss: 88.8459
      54/54 135 33.35; step accuracy: 6.9231 - loss: 6.3743 - val_accuracy: 6.4479 - val_loss: 69.6785
54/54 21s 368ms/step - accuracy: 6.9231 - loss: 6.3743 - val_accuracy: 6.4479 - val_loss: 69.6785

        54/54
        21s 368ms/step - accuracy; 8,9231 - 10ss; 8,3743
        Val_ecturacy; 8,5000

        Fpoot 5/56
        19s 349ms/step - accuracy; 8,9275 - 10ss; 8,2420 - Val_accuracy; 8,4479 - Val_10ss; 54,5698

      54/54 - 152 SMS,/Step - accuracy: 8.9277 - loss: 8.2783 - val_accuracy: 8.4479 - val_loss: 42.5919
54/54 - 21s 383ms/step - accuracy: 8.9277 - loss: 8.2783 - val_accuracy: 8.4479 - val_loss: 42.5919
54/54 - 19s 356ms/step - accuracy: 8.9211 - loss: 8.2676 - val_accuracy: 8.4479 - val_loss: 27.7518
      54/54 19s 354ms/step - accuracy: 0.9479 - loss: 0.1801 - val_accuracy: 0.6615 - val_loss: 7.9593
     54/54
Epoch 22/50
54/54
Epoch 23/50
54/54
                       19s 359ms/step - accuracy: 0.9634 - loss: 0.1230 - val_accuracy: 0.6146 - val_loss: 5.8954
20s 361ms/step - accuracy: 0.9430 - loss: 0.2192 - val_accuracy: 0.8966 - val_loss: 0.6860
      Fpoch 24/50
54/54
       54/54
Epoch 25/50
54/54
Epoch 26/50
54/54
                       193 1358ms/step - accuracy: 8.9686 - loss: 8.1245 - val_accuracy: 8.4896 - val_loss: 27.4846

193 136ms/step - accuracy: 8.9555 - loss: 8.1299 - val_accuracy: 8.9271 - val_loss: 8.2916

193 136ms/step - accuracy: 8.9656 - loss: 8.1744 - val_accuracy: 8.8385 - val_loss: 2.2625
8/8 6s 28ms/step - accuracy: 0.7642 - 10ss: 9.8739
Test accuracy: 0.73828125
```

- Training Accuracy: Achieved a test accuracy of 97.33% after 50 epochs.
- Performance: This model showed potential but demonstrated underfitting, with the
 validation accuracy lagging behind the training accuracy by a significant margin. This
 suggests that either the model requires more epochs to train or more advanced data
 augmentation techniques to improve generalization.
- Challenges: The model's simplicity compared to Models 1 and 2 may have contributed to its lower accuracy. Additionally, it might have benefitted from more complex feature extraction or regularization methods to reduce overfitting.

Comparison of Results

Model 2 achieved the highest test accuracy of **99.36%**, making it the best performer. Its deeper architecture with five Conv2D layers and more complex feature extraction likely contributed to this superior performance. Despite the larger number of parameters and longer training time, it delivered the most accurate results.

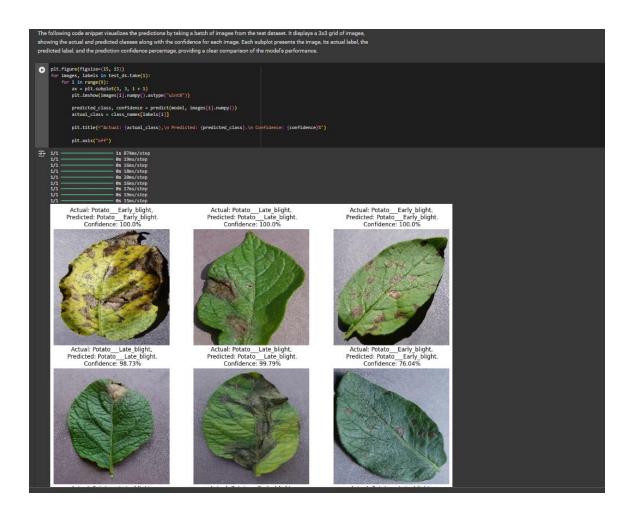
Model 1 followed closely with a test accuracy of **97.63%**. It showed strong performance with minimal overfitting, due to effective data augmentation techniques. Its six Conv2D layers allowed it to capture diverse image features efficiently.

Model 4 achieved a test accuracy of **97.33%**, slightly lower than Model 1. This model demonstrated potential but exhibited signs of underfitting, where the validation accuracy lagged behind the training accuracy. It may have benefitted from more training epochs or advanced data augmentation to improve generalization.

Model 3 had the test accuracy of 76.42% which is lower than comparing with other 3 models. While Batch Normalization improved training stability, the simpler architecture resulted in insufficient feature extraction, leading to underfitting. This model struggled to generalize well compared to the others, indicating the need for architectural improvements or additional tuning.

Training & Validation Accuracy and Loss

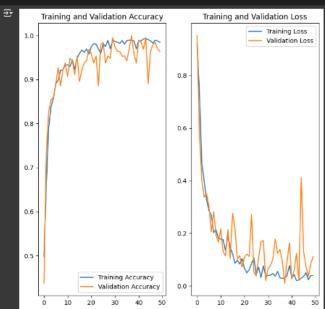
- Model Performance: It helps assess how well the model generalizes to unseen data.
- Overfitting Detection: A large gap between training and validation metrics signals overfitting, where the model performs well on training data but poorly on new data.
- **Hyperparameter Tuning**: Tracking these metrics informs adjustments to improve model performance, such as tuning learning rate or batch size.
- **Early Stopping**: Monitoring allows for early stopping to prevent overfitting and save computational resources.



Plotting Training and Validation Accuracy The first subslet displays the training accuracy and validation

The first subplot displays the training accuracy and validation accuracy over epochs. The x-axis represents the number of epochs, while the y-axis represents accuracy values. acc refers to the list of training accuracy values, and val_acc refers to the validation accuracy values.

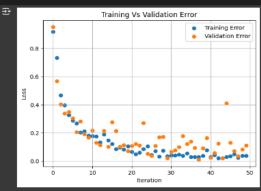
```
[] plt.figure(figsize=(8, 8))
plt.subplot(1, 2, 1)
plt.plot(range(len(acc)), acc, label='Training Accuracy')
plt.plot(range(len(val_acc)), val_acc, label='Validation Accuracy')
plt.legend(loc='lower right')
plt.title('Training and Validation Accuracy')
plt.subplot(1, 2, 2)
plt.plot(range(len(loss)),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



Scatter Plot: Training vs Validation Error

This plot visualizes the loss (error) during training and validation across different iterations (epochs). The x-axis represents the number of epochs, while the y-axis shows the loss values. history.history['10ss'] contains the training loss values, and history.history['val_loss'] contains the validation loss. The scatter plot allows easy comparison between training and validation errors to assess how well the model is generalizing.

```
[ ] plt.scatter(x=history.epoch,y=history.history['loss'],label='Training Error')
plt.scatter(x=history.epoch,y=history.history['val_loss'],label='Validation Error')
plt.grid(True)
plt.xlabel('tloration')
plt.ylabel('loss')
plt.title('Training Vs Validation Error')
plt.tiegend()
plt.show()
```





```
# Plot training and validation accuracy
plt.plot(history.history['accuracy'], label='Training Accuracy')
plt.plot(history.history['val_accuracy'], label='Validation Accuracy')
plt.vlabel('Epochs')
plt.vlabel('Accuracy')
plt.legend(loc='lower right')
plt.title('Training and Validation Accuracy')
plt.show()
8 Plot training and validation loss
plt.plot(history.history['loss'], label='Training Loss')
plt.plot(history.history['val_loss'], label='Validation Loss')
plt.vlabel('Epochs')
plt.ylabel('Loss')
plt.tlegend(loss'upper right')
plt.tlegend('Training and Validation Loss')
plt.show()
                                                       Training and Validation Accuracy
          1.0
          0.9
   Accuracy
0.7
          0.6
          0.5
                                                                                                                                Training Accuracy
                                                                                                                               Validation Accuracy
                         ò
                                                     10
                                                                                                                                                                     50
                                                                                                            30
                                                                                                                                          40
                                                           Training and Validation Loss

    Training Loss
    Validation Loss

          80
    SS01 40
          20
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                                                   10
                                                                                20
                                                                                                                                        40
                                                                                                                                                                    50
                                                                                      Epochs
```

```
Plotting Training and Validation Accuracy
The first subplot displays the training accuracy and validation accuracy over epochs. The x-axis represents the number of epochs, while the y-
axis represents accuracy values. acc refers to the list of training accuracy values, and val_acc refers to the validation accuracy values.
plt.figure(figsize=(8, 8))
plt.subplot(1, 2, 1)
plt.plot(range(len(acc)), acc, label='Training Accuracy')
plt.plot(range(len(val_acc)), val_acc, label='Validation Accuracy')
plt.legend(loc='lower right')
plt.title('Training and Validation Accuracy')
       plt.subplot(1, 2, 2)
plt.plot(range(len(loss)), loss, label='Training Loss')
plt.plot(range(len(val_loss)), val_loss, label='Validation Loss')
plt.legend(loc='upper right')
plt.title('Training and Validation Loss')
plt.show()
≆
                 Training and Validation Accuracy
                                                                                          Training and Validation Loss
                                                                                                                     Training Loss
         1.0
                                                                                                              --- Validation Loss
                                                                             700
         0.9
                                                                             600
                                                                             500
         0.8
                                                                             400
         0.7
                                                                             300
         0.6
                                                                             200
                                                                             100
         0.5
                                          Training Accuracy
                                     20
[] plt.scatter(x=history_4.epoch,y=history_4.history['loss'],label='Training Error')
plt.scatter(x=history_4.epoch,y=history_4.history['val_loss'],label='Validation Error')
        plt.scatter(x=nistory_4.epocn,y=nistory_4.plt.grid(True)
plt.xlabel('Iteration')
plt.ylabel('Loss')
plt.title('Training Vs Validation Error')
plt.legend()
plt.show()
₹
                                                        Training Vs Validation Error
                                                                                                                Training Error
                700
                                                                                                                Validation Error
                600
                500
          s 400
                300
                200
                100
                                                     •
                    0
                                                                                                                 40
                                                 10
                                                                      20
                                                                                            30
                                                                           Iteration
```

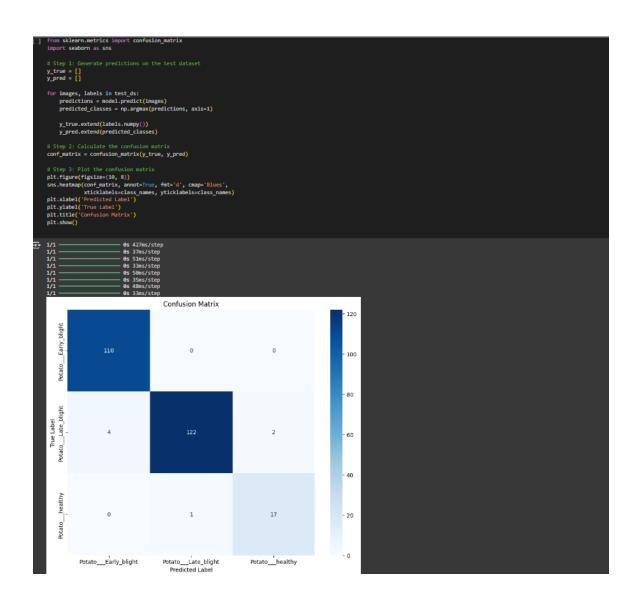
Confusion Matrix

Visualizing training and validation metrics plays a crucial role in understanding how a machine learning model evolves during training. To ensure effective learning and avoid overfitting, it's essential to monitor these metrics closely through graphical representations.

In this project, Matplotlib was used to create detailed visualizations that illustrate training accuracy, validation accuracy, training loss, and validation loss over time. These plots provide a clear visual summary of the model's performance, making it easier to spot trends, identify overfitting or underfitting, and understand the impact of hyperparameter adjustments.

All team members relied on these visualizations to assess and improve the performance of their individual models designed to classify potato leaf diseases. The graphical representations served as an indispensable tool for evaluating model convergence, guiding model adjustments, and ensuring that each model maintained a healthy balance between accuracy and generalization.

- **Potato Early Blight**: Predicted 110 correctly, 4 misclassified as Potato Late Blight, 0 misclassified as Healthy.
- **Potato Late Blight**: Predicted 122 correctly, 2 misclassified as Healthy, 0 misclassified as Early Blight.
- Potato Healthy: Predicted 17 correctly, 1 misclassified as Late Blight.



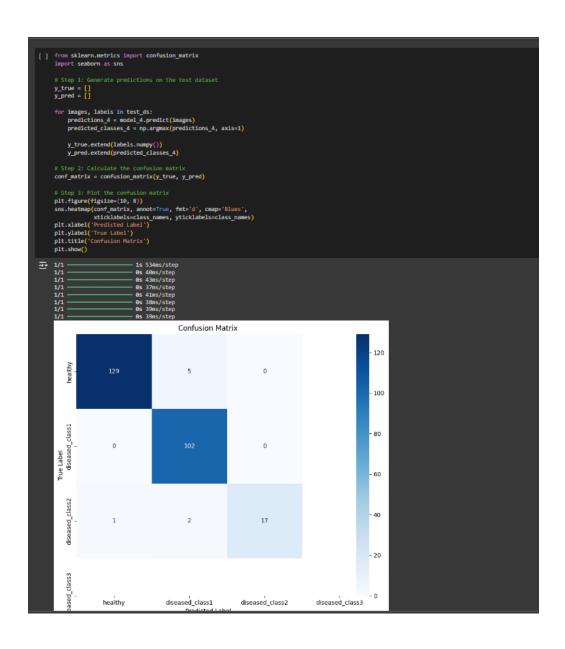
- **Potato Early Blight**: Predicted 110 correctly, 0 misclassified as Potato Late Blight, 0 misclassified as Healthy.
- **Potato Late Blight**: Predicted 128 correctly, 1 misclassified as Healthy, 0 misclassified as Early Blight.
- **Potato Healthy**: Predicted 17 correctly, 1 misclassified as Late Blight.



- Potato Early Blight: Predicted 134 correctly, 48 misclassified as Potato Late Blight,
 6 misclassified as Healthy.
- **Potato Late Blight**: Predicted 44 correctly, 10 misclassified as Healthy, 48 misclassified as Early Blight.
- **Potato Healthy**: Predicted 11 correctly, 3 misclassified as Late Blight, 6 misclassified as Early Blight.



- **Potato Early Blight**: Predicted 129 correctly, 5 misclassified as Potato Late Blight, 0 misclassified as Healthy.
- Potato Late Blight: Predicted 102 correctly, 0 misclassified as Healthy, 0 misclassified as Early Blight.
- **Potato Healthy**: Predicted 17 correctly, 2 misclassified as Late Blight, 1 misclassified as Early Blight.



Overall Observation

- Model 1 and Model 2 performed consistently well, with minimal misclassifications across all classes.
- **Model 3** struggled significantly with Potato Late Blight predictions, misclassifying many cases as Early Blight.
- Model 4 had generally good performance, though it had some misclassifications for Early Blight and Late Blight.

In conclusion, **Model 2** seems to offer the most balanced performance across all categories, while **Model 3** may need further refinement to handle the Late Blight class more effectively.

Critical Analysis and Discussion

How Accuracy Could Be Improved

- **Hyperparameter Tuning**: Adjusting learning rates, batch sizes, and the number of epochs could further improve accuracy. For instance, increasing the number of epochs for Model 4 could help prevent underfitting.
- More Data Augmentation: Increasing the variety of augmentations such as zoom, brightness, and contrast adjustments could help improve generalization, especially for underperforming models like Model 4.
- **Regularization**: Implementing techniques like **L2 regularization** could help prevent overfitting in high-capacity models like Model 2.
- Pre-trained Models: Using a pre-trained architecture such as VGG16 or ResNet
 could improve performance, especially in complex classification tasks with limited
 training data.

Possible Future Work

- **Transfer Learning**: Implementing transfer learning with pre-trained models could yield higher accuracy with fewer training epochs.
- **Ensemble Methods**: Combining predictions from multiple models in an ensemble could improve overall performance by leveraging the strengths of each individual model.
- Optimization Algorithms: Experimenting with different optimizers such as RMSprop or SGD could fine-tune model performance and convergence.
- Further Hyperparameter Optimization: Automated techniques like Grid Search or Random Search could be used to systematically explore a wider range of hyperparameters.

Potato Disease Identification API

Overview

The **Potato Disease Identification API** is a web-based API developed using **FastAPI**. This API utilizes a machine learning model to identify diseases in potato leaves by analyzing uploaded images. The API can classify potato leaf images into three categories: **Early Blight**, **Late Blight**, or **Healthy**. The application is designed to provide quick and accurate predictions, assisting farmers and agricultural professionals in managing crop health.

How It Works

- 1. **Image Upload**: Users can upload an image of a potato leaf to the /predict endpoint.
- 2. **Preprocessing**: The uploaded image is resized and normalized to meet the input requirements of the machine learning model.
- 3. **Model Prediction**: A pre-trained TensorFlow model processes the image and predicts the class of disease or indicates that the plant is healthy.
- 4. **Response**: The API returns the predicted disease class along with a confidence score.

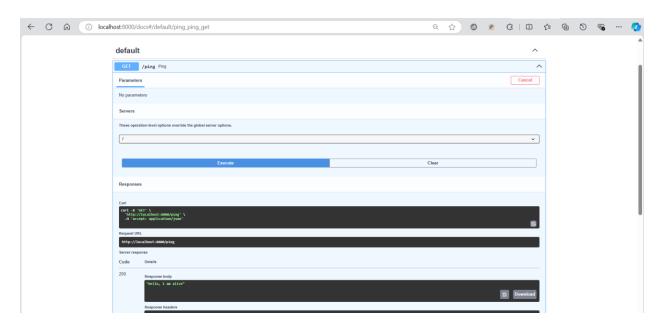
API Endpoints

1. Health Check

Endpoint: /pingMethod: GET

• **Description**: A simple health check endpoint to verify if the API is running.

Response:



2. Predict Disease

• **Endpoint**: /predict

• Method: POST

• **Description**: Upload an image of a potato leaf to receive a prediction about its health status.

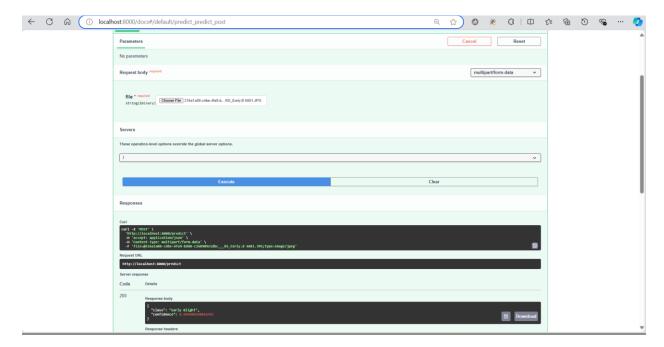
Request:

• Form Data:

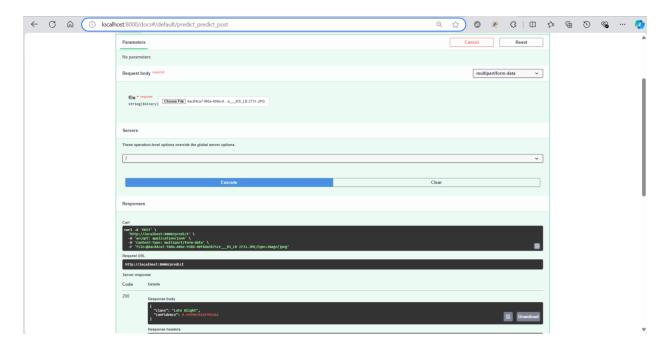
o file: The image file of the potato leaf (must be in a supported image format).

Response:

Early Blight Disease Identification.



Late Blight Disease Identification



React Application

The React application interacts with the FastAPI to facilitate the image upload process and display the results. Below are the key functionalities implemented in the React app:

Features

1. User Interface:

The app includes a user-friendly interface for uploading images of potato leaves.

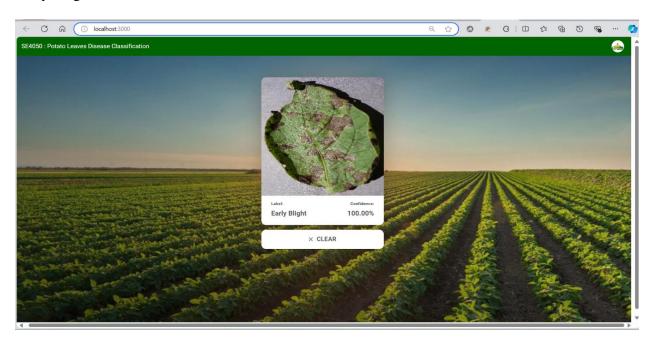
2. API Calls:

- The React application makes asynchronous calls to the FastAPI endpoints using Axios or Fetch API.
- When the user uploads an image, the app sends a POST request to the /predict endpoint.

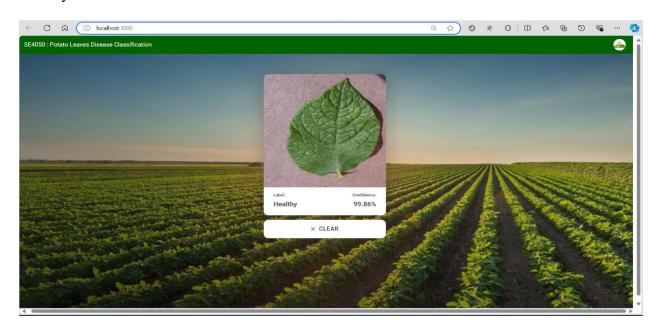
3. **Displaying Results**:

- Upon receiving the response from the API, the React application displays the predicted class and confidence score to the user.
- The results are shown in a structured format, allowing users to understand the health status of their potato plants at a glance.

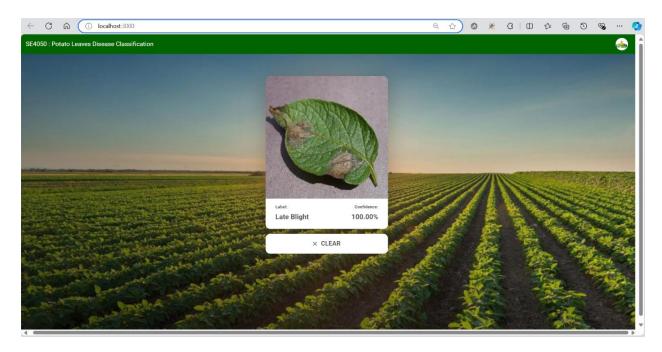
Early Blight Disease Identification.



Healthy Plants Identification



Late Blight Disease Identififcation



Workflow

- 1. **Upload Image**: The user selects a potato leaf image and clicks the "Upload" button.
- 2. **Send Request**: The React app sends the image to the /predict endpoint.
- 3. **Receive Response**: The API processes the image and returns the prediction.
- 4. **Display Prediction**: The React application displays the predicted disease class and confidence score.

To run the application

- 1. run the backend server(FasterAPI potato-disease-application-python-api)
- 2. set the environment

```
$env:NODE_OPTIONS="--openssl-legacy-provider"
```

3. Install the dependencies

```
npm install
```

Conclusion

In this study of potato disease identification using various CNN architectures, **Model 2** demonstrated the highest accuracy at **99.36%**, showcasing the effectiveness of deeper architectures and complex feature extraction. **Model 1** and **Model 4** also performed well with accuracies of **97.63%** and **97.33%**, respectively, although Model 4 exhibited signs of underfitting. In contrast, **Model 3** underperformed with a test accuracy of **76.42%**, indicating a need for further architectural enhancement. Overall, these results highlight the importance of model complexity and the balance between training efficiency and generalization in achieving high performance in image classification tasks.

References

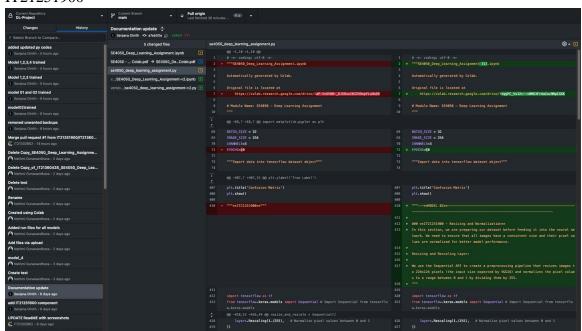
Plant Village Dataset:

https://www.kaggle.com/datasets/arjuntejaswi/plant-village

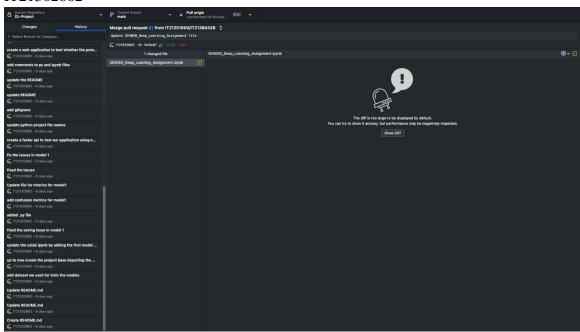
Appendix

Contribution

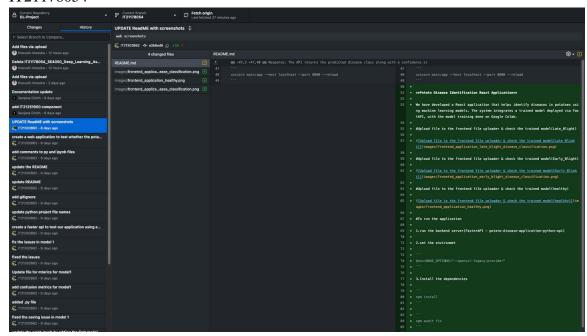
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