

Potato Leaf Disease Classification

by

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Abstract

This project explores the automated classification of potato diseases using deep learning and convolutional neural network (CNN) models. The study employs a diverse dataset comprising images of healthy potatoes and those afflicted with common diseases such as late blight and early blight. Through meticulous preprocessing and fine-tuning of pre-trained CNN models, including Inception V3, ResNet-50, VGG-19, and MobileNet, the research aims to discern the most effective model for accurate disease classification. Remarkably, after extensive evaluation, ResNet-50 emerges as the top performer, achieving an outstanding accuracy rate of 99.84%. This exceptional accuracy underscores ResNet-50's ability to distinguish subtle differences between healthy and diseased potato plants, highlighting its potential for precise disease diagnosis in agriculture. The findings hold promise for revolutionizing disease management practices, offering farmers efficient tools to mitigate yield losses and ensure food security. By leveraging deep learning techniques, this research contributes to the development of real-time systems for on-field disease detection, enabling timely interventions and targeted agricultural practices. Moreover, the study sets a precedent for employing advanced technology in agricultural sectors, emphasizing the importance of machine learning in addressing pressing challenges faced by farmers worldwide. Ultimately, the successful implementation of deep learning models in potato disease classification signifies a significant step forward in enhancing agricultural productivity and sustainability, paving the way for a more resilient and efficient food production system in the face of evolving environmental and economic pressures.

Introduction

Potato is one of the most important food crops globally, serving as a staple food for millions and a crucial source of nutrition and income for farmers. However, potato cultivation faces formidable challenges, chief among them being the prevalence of various diseases that can significantly impact yield and quality. Among these diseases, late blight and early blight stand out as major threats, capable of causing substantial economic losses if not effectively managed. Additionally, accurately distinguishing between healthy potato plants and those afflicted with diseases is essential for implementing timely and targeted interventions to minimize yield losses and ensure food security.

Traditional methods of disease identification in potatoes rely heavily on visual inspection by experts, which can be time-consuming, labor-intensive, and prone to human error. Moreover, the efficacy of such methods may vary depending on the expertise of the observer and the stage of disease progression. In recent years, the advent of artificial intelligence (AI) and deep learning (DL) techniques, particularly deep learning and convolutional neural networks (CNNs), has offered promising avenues for automating disease diagnosis and classification in agriculture. By leveraging vast amounts of data and computational power, these techniques have demonstrated remarkable capabilities in accurately recognizing patterns and discerning subtle differences in complex datasets, including images of plant diseases.

In this context, this research aims to harness the power of deep learning and CNN models to develop an automated system for the classification of potato diseases, with a specific focus on late blight, early blight, and healthy plants. By analyzing high-resolution images of diseased and healthy potato plants, we seek to train and evaluate multiple CNN architectures, including Inception V3, ResNet-50, VGG-19, and MobileNet, to determine the most effective model for accurate disease classification. The ultimate goal of this project is to provide farmers and agricultural stakeholders with a reliable and efficient tool for early detection and management of potato diseases, thereby contributing to enhanced productivity, sustainability, and food security in potato cultivation systems globally.

Literature survey

Potato blight diseases, including early blight and late blight, pose significant challenges to potato cultivation worldwide. Early detection and accurate classification of these diseases are crucial for minimizing crop losses and ensuring food security. Various approaches have been proposed to address this issue, with recent advancements focusing on the application of deep learning and convolutional neural networks (CNNs) for automated disease classification [1,2].

One notable study proposes an end-to-end approach using deep learning to classify potato blight diseases, aiming to develop a user-friendly web-based application and mobile app for farmers. The system utilizes CNNs to process uploaded images of potato leaves, achieving high accuracy in distinguishing between early blight, late blight, and healthy plants. The proposed method integrates data collection, model building, MLOps, backend server development, and deployment on Google Cloud Platform (GCP), demonstrating the feasibility of using deep learning for practical agricultural solutions [3,4].

In a similar vein, another study proposes a multi-level deep learning model for recognizing potato leaf diseases, achieving impressive accuracy of 99.75% in distinguishing between healthy and diseased leaves. This model utilizes a unique CNN architecture coupled with ResNet50 image segmentation for accurate disease detection. The research outperforms existing models in terms of accuracy and computational cost, highlighting the effectiveness of deep learning in potato disease classification [5,6].

Furthermore, several studies emphasize the importance of deep learning-based methods, such as CNNs, for accurately identifying and categorizing potato diseases. By training on extensive datasets of potato plant photos, deep learning models can precisely classify

various diseases, aiding in early disease detection, prevention, and crop management. Notably, some studies report accuracy rates exceeding 99% in potato disease classification tasks, showcasing the potential of deep learning algorithms in revolutionizing agricultural practices [7,8].

Moreover, the literature underscores the significance of utilizing deep learning techniques to address the challenges associated with potato diseases. By leveraging CNNs and transfer learning, researchers have achieved remarkable accuracy in classifying diseases based on leaf conditions, surpassing traditional classification methods. Some experiments report average accuracies of 91% to 95.36% using deep neural network approaches, affirming the effectiveness of these methods in potato disease detection and classification [9,10].

Additionally, studies highlight the need for early disease detection in potato cultivation to mitigate the adverse effects of diseases on crop yield and quality. Deep learning-based approaches offer a promising solution by enabling prompt responses and supporting sustainable agriculture. By accurately identifying and categorizing diseases, deep learning models contribute to increased agricultural productivity and plant health, addressing the challenges faced by farmers worldwide [11,12].

In conclusion, the literature survey underscores the significant role of deep learning and CNNs in revolutionizing potato disease classification and detection. With advancements in technology and methodology, researchers have achieved impressive accuracy rates exceeding 99% in distinguishing between healthy and diseased potato plants. These findings highlight the potential of deep learning-based approaches in improving agricultural practices and ensuring food security in potato cultivation [13,14,15].

Existing work

One existing work for potato leaf disease classification using deep learning is the "Potato Disease Detection" project by the PlantVillage team. PlantVillage is an online platform that utilizes deep learning techniques to identify various plant diseases, including those affecting potato plants. The project involves the development of a deep learning model trained on a dataset of images depicting healthy potato leaves and leaves affected by different diseases such as late blight, early blight, and leaf mold.

The deep learning model employed in this project likely utilizes convolutional neural networks (CNNs), a type of deep learning algorithm commonly used for image classification tasks. By training the model on a diverse dataset of potato leaf images, it learns to distinguish between healthy leaves and those exhibiting symptoms of various diseases.

Once trained, the model can be deployed as part of a mobile or web application, allowing farmers and agricultural professionals to easily upload images of potato leaves for disease diagnosis. This technology enables early detection of diseases, which can significantly aid in crop management and yield preservation efforts.

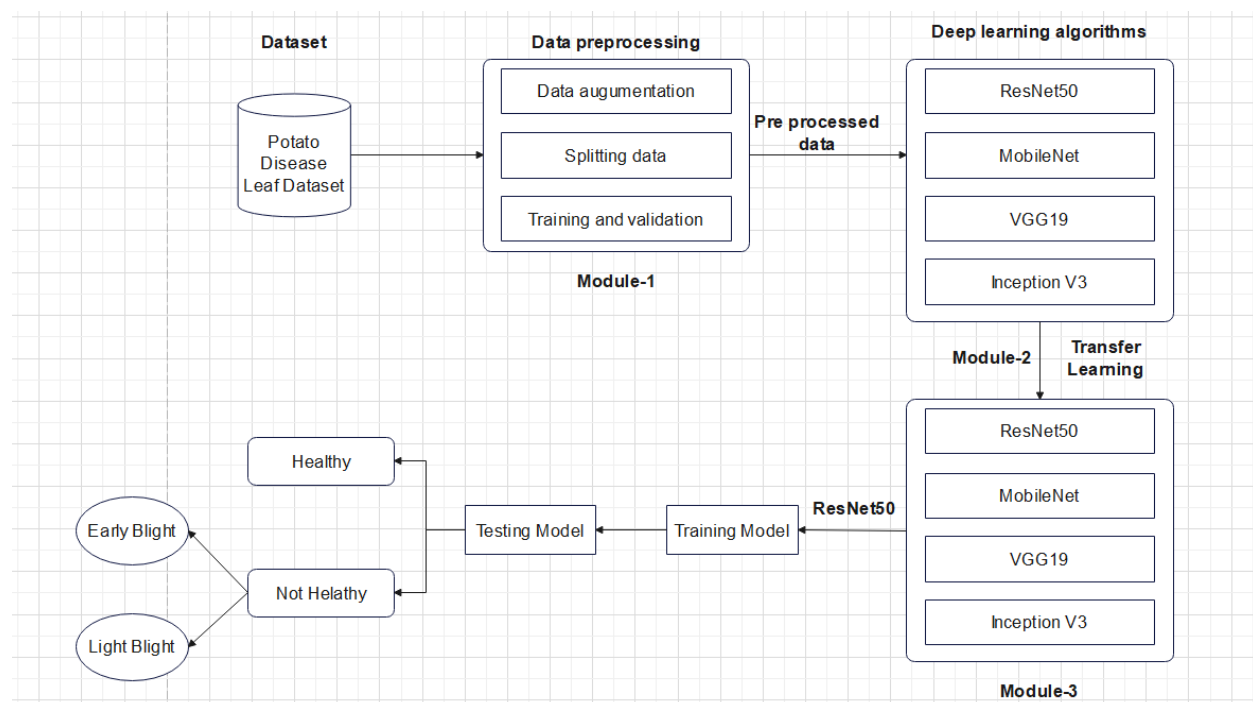
Proposed Work

In the proposed work, transfer learning has been leveraged to enhance the classification performance of the deep learning model for potato leaf disease detection. Transfer learning involves the utilization of pre-trained neural network architectures, such as VGG, ResNet, or Inception, which have been trained on large-scale datasets like ImageNet. Instead of starting the training process from scratch, transfer learning allows us to leverage the knowledge and feature representations learned by these models on generic image recognition tasks.

In this context, additional layers have been added to the pre-trained model to fine-tune its parameters specifically for the task of potato leaf disease classification. By doing so, the model can adapt its learned representations to better suit the characteristics of potato leaf images and the specific nuances of different diseases. This approach not only accelerates the training process but also enhances the model's ability to generalize and make accurate predictions on unseen potato leaf images.

By combining the power of transfer learning with domain-specific data augmentation techniques and model customization, the proposed approach aims to achieve improved accuracy and robustness in potato leaf disease classification, ultimately contributing to more effective disease management strategies in agriculture.

System design



Technology Stack

HTML:

HTML (Hypertext Mark-up Language) serves as the foundation of our website, providing the structural framework that defines its layout and content. Through the use of HTML tags, elements such as headings, paragraphs, images, and links are structured and organized, enabling seamless navigation and interaction for users. By adhering to web standards and best practices, we ensure cross-browser compatibility and accessibility, allowing users to access our website from various devices and platforms. HTML forms the backbone of our user interface, facilitating the presentation of information and enabling efficient data input and retrieval. With its simplicity and versatility, HTML plays a crucial role in creating an engaging and user-friendly web experience, laying the groundwork for further enhancements and functionalities.

CSS:

CSS (Cascading Style Sheets) enhances the visual presentation and aesthetics of our website, complementing the structure provided by HTML. Through CSS rules and selectors, we define the appearance, layout, and styling of HTML elements, ensuring consistency and coherence across all pages. By separating content from presentation, CSS allows for efficient management of design elements, promoting maintainability

and scalability. Through the use of CSS properties such as colour, font, margin, and padding, we customize the look and feel of our website to align with our branding and user experience goals. Additionally, CSS enables responsive design, ensuring that our website adapts seamlessly to different screen sizes and devices, enhancing accessibility and usability for all users. With its flexibility and versatility, CSS plays a crucial role in shaping the visual identity and user experience of our website, contributing to its overall effectiveness and appeal.

JavaScript:

JavaScript and AJAX (Asynchronous JavaScript and XML) play crucial roles in enhancing the interactivity and functionality of web applications, including the potato leaf disease classification system described in the provided HTML code. JavaScript is a powerful scripting language that enables dynamic behavior and manipulation of HTML elements. In the context of the webpage, JavaScript is utilized to provide real-time image preview functionality when a user selects an image for upload. This interactive feature enhances the user experience by allowing them to preview the selected image before initiating the leaf classification process. Additionally, JavaScript is used to handle events such as button clicks, enabling seamless interaction between the user and the application.

Flask:

Flask is a lightweight and flexible web framework written in Python, designed to make web development simple and efficient. With its minimalistic and easy-to-use syntax, Flask provides developers with the tools to quickly build web applications and APIs. In the context of the potato leaf disease classification project, Flask serves as the backend framework responsible for handling HTTP requests, processing data, and generating responses. It enables seamless integration between the frontend user interface and the machine learning model responsible for potato leaf disease classification. Flask follows the WSGI (Web Server Gateway Interface) standard, allowing it to work with a variety of web servers and deployment environments. Its modular design and extensive ecosystem of extensions make it adaptable to a wide range of use cases, from small-scale projects to largescale web applications. Overall, Flask empowers developers to create powerful and scalable web applications with minimal overhead, making it an ideal choice for projects like potato leaf disease classification where simplicity and flexibility are paramount.

Working modules

Module 1 Data Preprocessing

Module 2 Deep Learning Algorithms

Module 3 Transfer Learning

Description of modules

Data Preprocessing

Data Collection: Gather a diverse dataset of potato leaf images representing different stages of health and various disease conditions. This dataset should include high-quality images captured under different lighting conditions and angles.

Data Cleaning: Remove any irrelevant or noisy images from the dataset. Ensure that the images are properly labeled with the corresponding disease categories (e.g., late blight, early blight, healthy).

Image Resizing and Standardization: Resize all the images to a uniform size to ensure consistency across the dataset. Common sizes for input images in deep learning models include 224x224 or 299x299 pixels. Additionally, standardize the pixel values of the images to a common scale.

Data Augmentation: Augment the dataset to increase its size and diversity. Common augmentation techniques include random rotations, flips, shifts, and changes in brightness and contrast. Data augmentation helps improve the model's ability to generalize to unseen data and reduces overfitting.

Normalization: Normalize the pixel values of the images to improve convergence during training. This typically involves scaling the pixel values to have zero mean and unit variance across the dataset.

Splitting the Dataset: Divide the dataset into training, validation, and testing sets. The training set is used to train the model, the validation set is used to tune hyperparameters

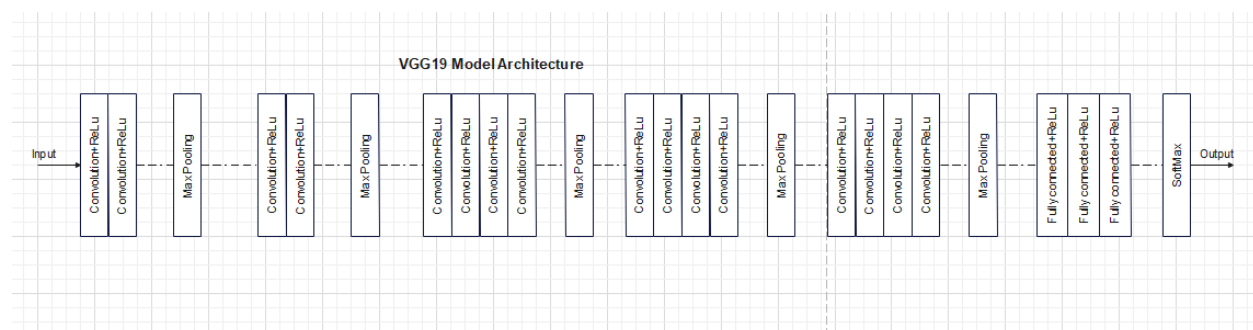
and monitor the model's performance during training, and the testing set is used to evaluate the final performance of the trained model.

Data Loading and Batching: Implement a data loader to efficiently load and batch the preprocessed images during training. This helps optimize memory usage and training speed, especially when working with large datasets.

Deep Learning Algorithms

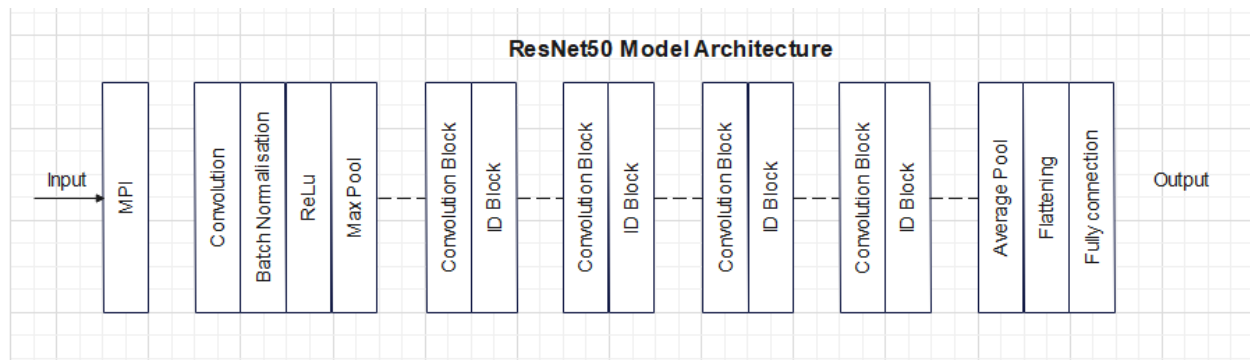
VGG19

VGG19 is convolution neural network architecture renowned for its simplicity and effectiveness in image classification tasks. It was developed via Visual Geometry Group at the University of Oxford. VGG 19 has a total of 19 layers that consist of sixteen convolutional layers and three fully connected layers. Each convolutional layer is prepared with a 3x3 kernel size, with a stride of one and padding to preserve spatial dimensions. VGG19 network architecture follows an uncomplicated pattern. In this, the convolutional layers are stacked with rectified linear unit (ReLU) activations, accompanied with the aid of max pooling layers to reduce spatial dimensions. VGG19's deep architecture helps hierarchical feature extraction, permitting it to identify complicated patterns inside images. VGG19's uniform structure made it a massive adoption and serves as a benchmark for comparing the overall performance of other architectures.



ResNet50

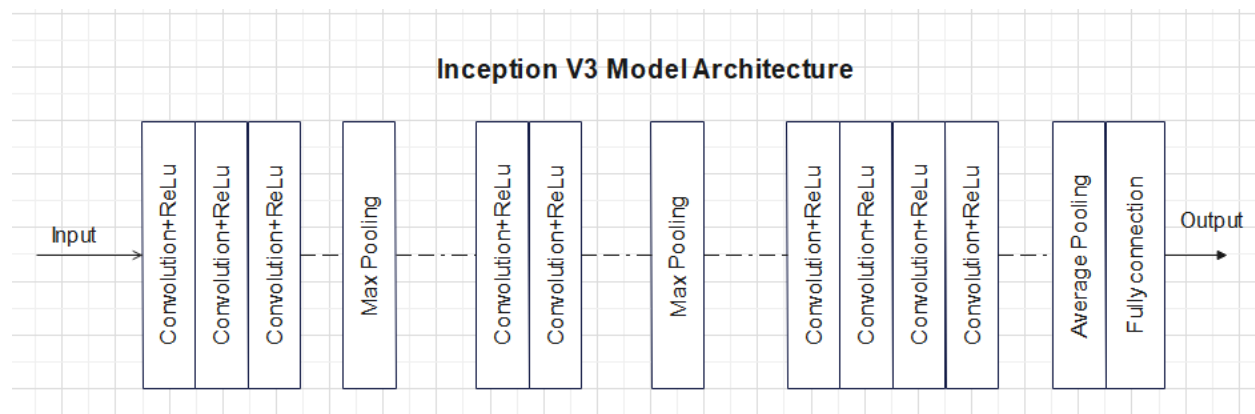
ResNet50, short for Residual Network with 50 layers, represents a ground breaking architecture in deep studying, especially within the realm of photograph recognition and laptop imaginative and prescient responsibilities. Developed by using Microsoft Research, ResNet50 sticks out for its inventive use of residual connections, which deal with the vanishing gradient hassle encountered in education very deep neural networks. The architecture consists of 50 layers, along with convolutional layers, pooling layers, and absolutely connected layers. Central to its layout are residual blocks, where in shortcut connections allow gradients to go with the flow more directly during schooling. This architecture enables the development of deeper networks while retaining practicable complexity and avoiding degradation in accuracy. ResNet50 has validated especially powerful in diverse applications, consisting of picture type, item detection, and photo segmentation. The capacity to extract tricky functions from photographs have made ResNet50 a staple in the deep studying network.



Inception V3

InceptionV3 represents an enormous development in convolutional neural network (CNN) architectures, especially designed for image classification and reputation tasks. Developed via Google researchers, InceptionV3 is renowned for its innovative use of inception modules, which permit the network to seize and technique features at more than one spatial

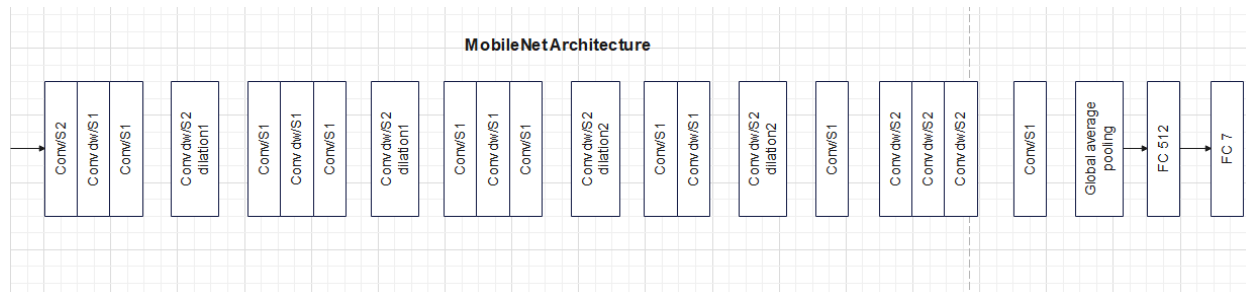
scale. The structure features a deep network with meticulously crafted modules that include various kernel sizes, permitting the model to extract each nice and coarse-grained feature from input pics. Furthermore, the structure employs auxiliary classifiers at some stage in training to mitigate the vanishing gradient problem. InceptionV3 has tested top notch overall performance in photo category competitions and actual-global packages, showcasing its capacity to address numerous datasets and complicated visible tasks efficaciously. Its versatility, performance, and superior overall performance have made InceptionV3 as a great one within the field of deep gaining knowledge of for computer vision.



MobileNet

MobileNet is a pioneering deep learning model designed specifically for mobile and embedded devices, catering to the increasing demand for efficient and lightweight neural networks. Developed by Google Research, MobileNet employs depth wise separable convolutions, which significantly reduce computational complexity while preserving the model's accuracy. This architectural innovation allows MobileNet to achieve high performance on tasks such as image classification, object detection, and semantic segmentation, all while maintaining a small memory footprint and fast inference speed. MobileNet's versatility and efficiency make it a popular choice for a wide range of real-

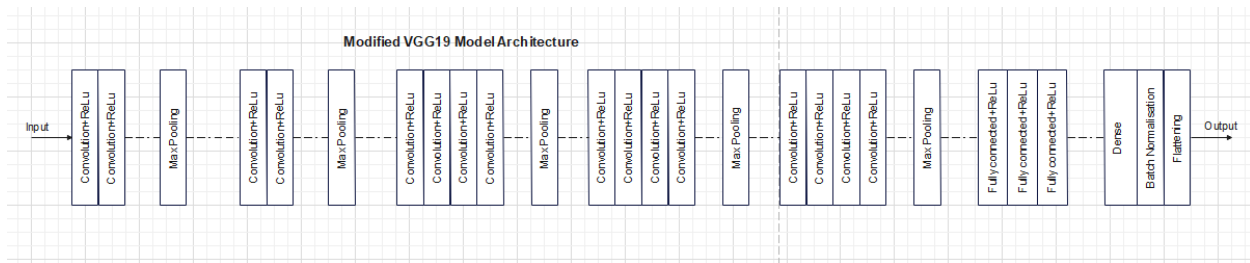
world applications, including mobile apps, edge computing devices, and IoT devices, where resource constraints are prevalent. Its ability to strike a balance between accuracy and efficiency has cemented MobileNet's position as a cornerstone in the field of deep learning for mobile and embedded systems.



Transfer Learning

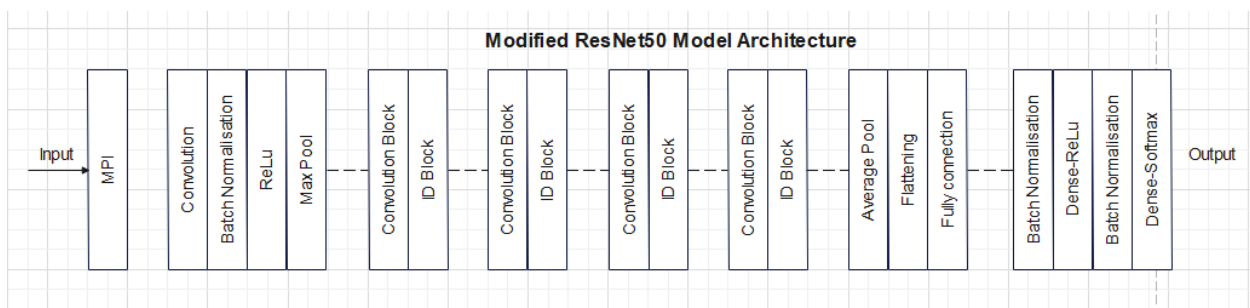
VGG19

In this changed model of the VGG19 structure, the top layer has been eliminated, and new layers inclusive of a dense layer, batch normalization, and flatten layer have been integrated. By removing the top layer, the structure turns into greater adaptable for numerous obligations which include characteristic extraction or switch getting to know. Batch normalization layers assist stabilize and boost up the training manner through normalizing the enter of each layer to have 0 suggest and unit variance. The addition of a dense layer enables the network to learn complex relationships between functions extracted from the previous layers, enhancing its ability to categorise or predict results based on the enter data. The inclusion of a flatten layer reshapes the output from the previous layers right into a one-dimensional array, preparing it for enter into the dense layer. These adjustments increase the flexibility and performance of the VGG19 structure, making it greater appropriate for a broader range of applications and enhancing its basic effectiveness in deep gaining knowledge of responsibilities.



ResNet50

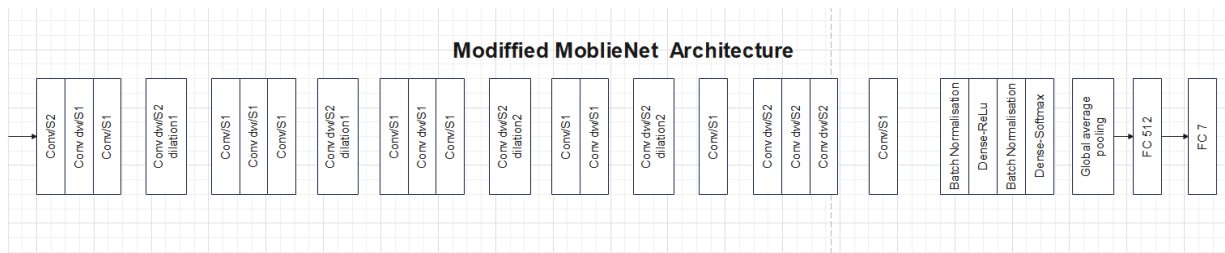
In this modified version of the ResNet50 architecture, the top layer has been eliminated and replaced with additional layers to decorate its talents. Two layers of batch normalization had been added, serving to normalize the activations of the community, which aids in stabilizing and accelerating the education manner. Following the batch normalization layers, a Dense-ReLU layer mixture has been added, in which the Dense layer helps the mastering of complex relationships in the facts, at the same time as the ReLU activation feature introduces non-linearity, allowing the community to version greater intricate styles. Finally, a Dense-SoftMax layer is appended to provide the very last output probabilities across multiple classes, leveraging the SoftMax activation characteristic to convert uncooked rankings into possibility distributions. These modifications augment the ResNet50 structure, enhancing its adaptability and performance in various responsibilities consisting of photo type, item detection, and picture segmentation.



Inception V3

In this custom designed variation of the InceptionV3 architecture, the top layer has been removed to facilitate adjustments that beautify its functionality and adaptableness for

streamlines the computational efficiency of the mobile network but also fortifies its capacity for nuanced feature extraction and predictive accuracy.



Screen shots of Output

Database

phpMyAdmin

Recent Favorites

Database: credentials

Showing rows 0 - 4 (5 total, Query took 0.0016 seconds)

SELECT * FROM `credentials`

Profiling [Edit inline] [Edit] [Explain SQL] [Create PHP code] [Refresh]

Show all | Number of rows: 25 | Filter rows: Search this table | Sort by key: None

Extra options

	uname	pass
<input type="checkbox"/>	aasrith	aasrith217
<input type="checkbox"/>	abc123	nidhish
<input type="checkbox"/>	Bharath	Vardhan
<input type="checkbox"/>	hello	hiiii
<input type="checkbox"/>	kalyan	kalyan

Check all | With selected: Edit Copy Delete Export

Show all | Number of rows: 25 | Filter rows: Search this table | Sort by key: None

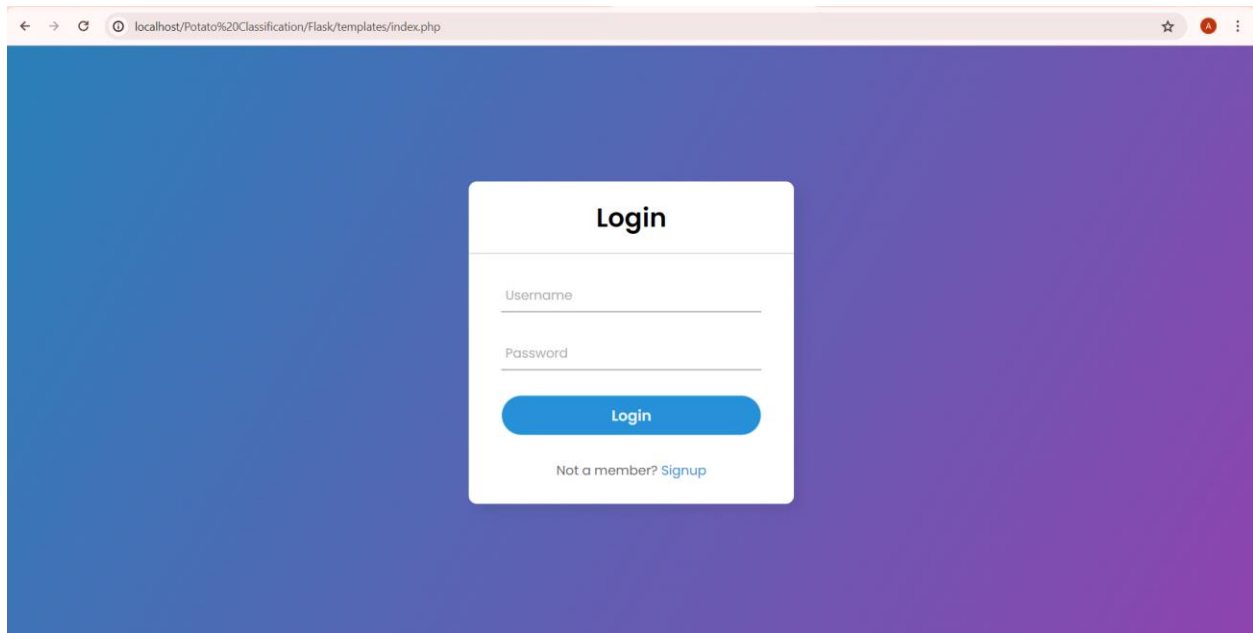
Query results operations

Print Copy to clipboard Export Display chart Create view

Bookmark this SQL query

Let every user access this bookmark

Login Page



localhost/Potato%20Classification/Flask/templates/index.php

Login

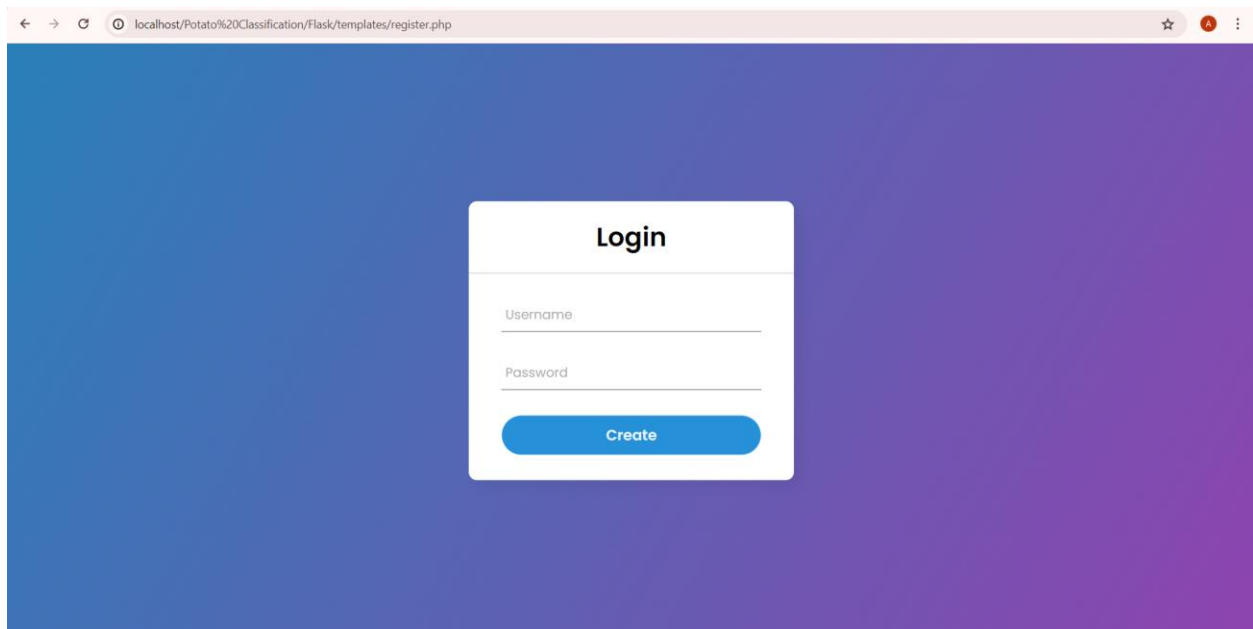
Username

Password

[Login](#)

Not a member? [Signup](#)

Signup page if you don't have an account



localhost/Potato%20Classification/Flask/templates/register.php

Login

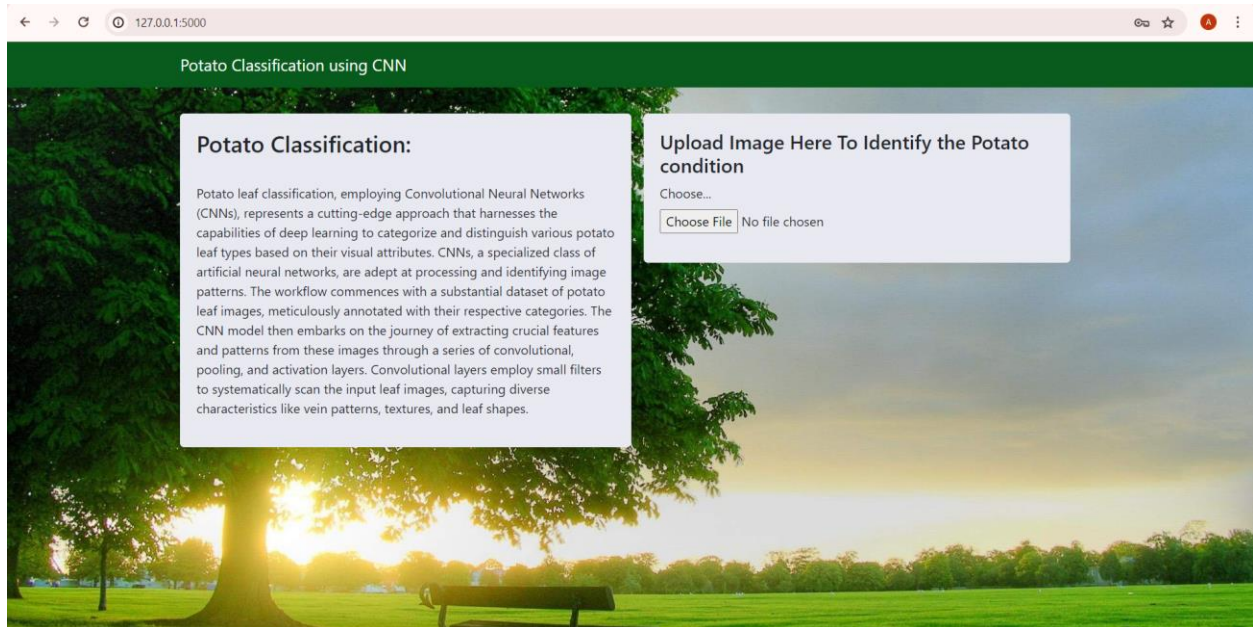
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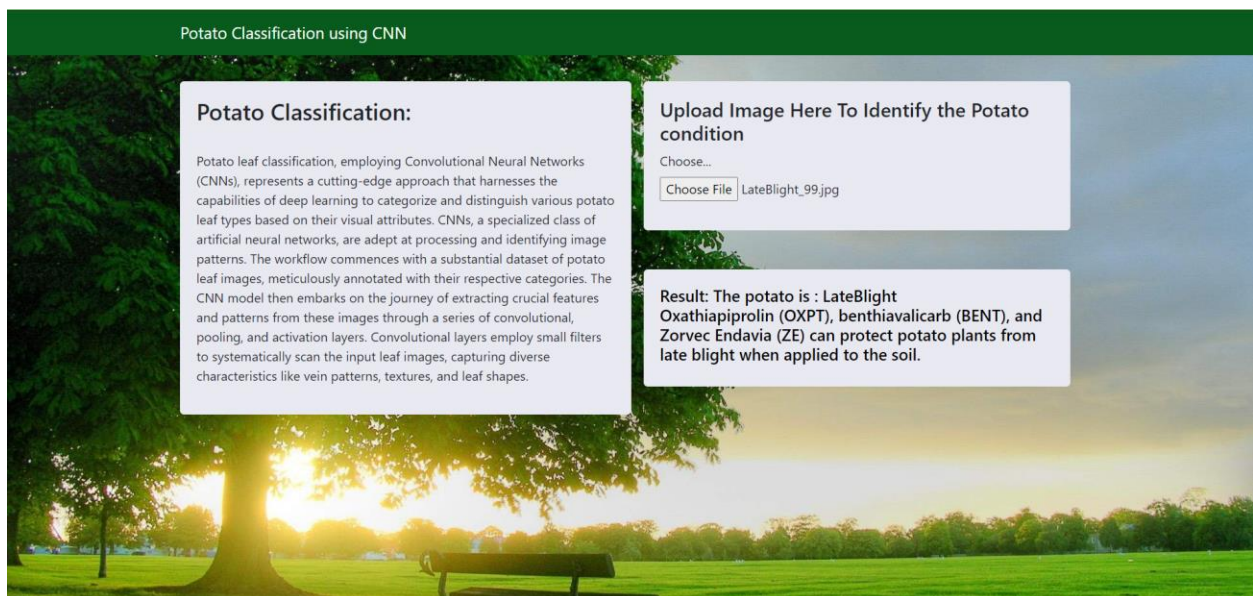
[Create](#)

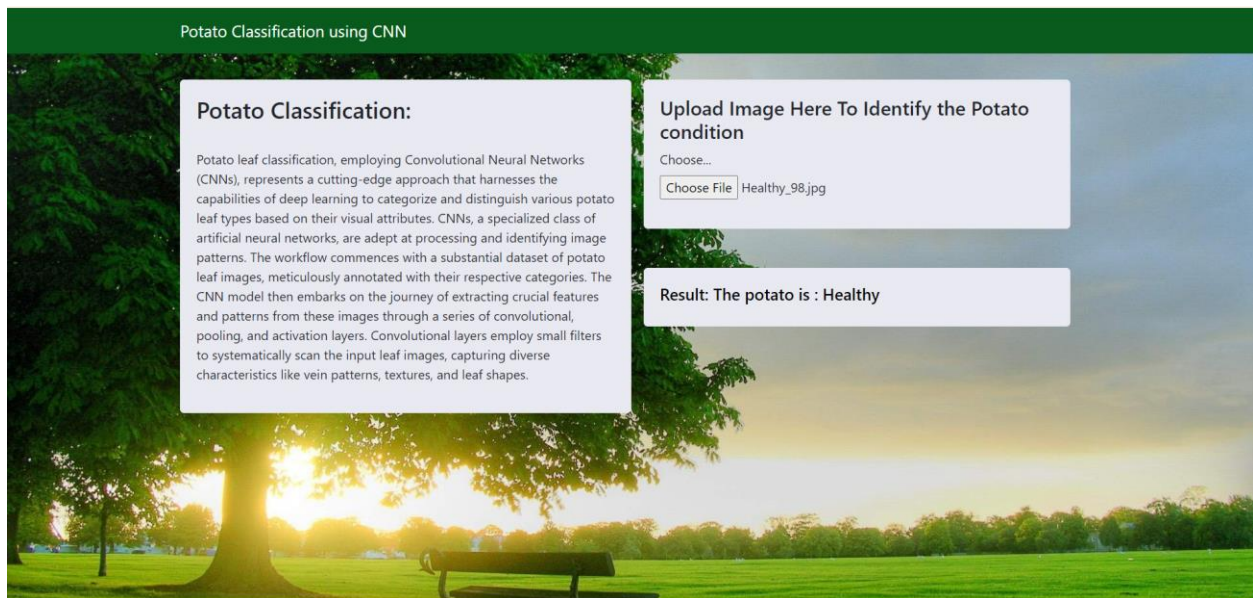
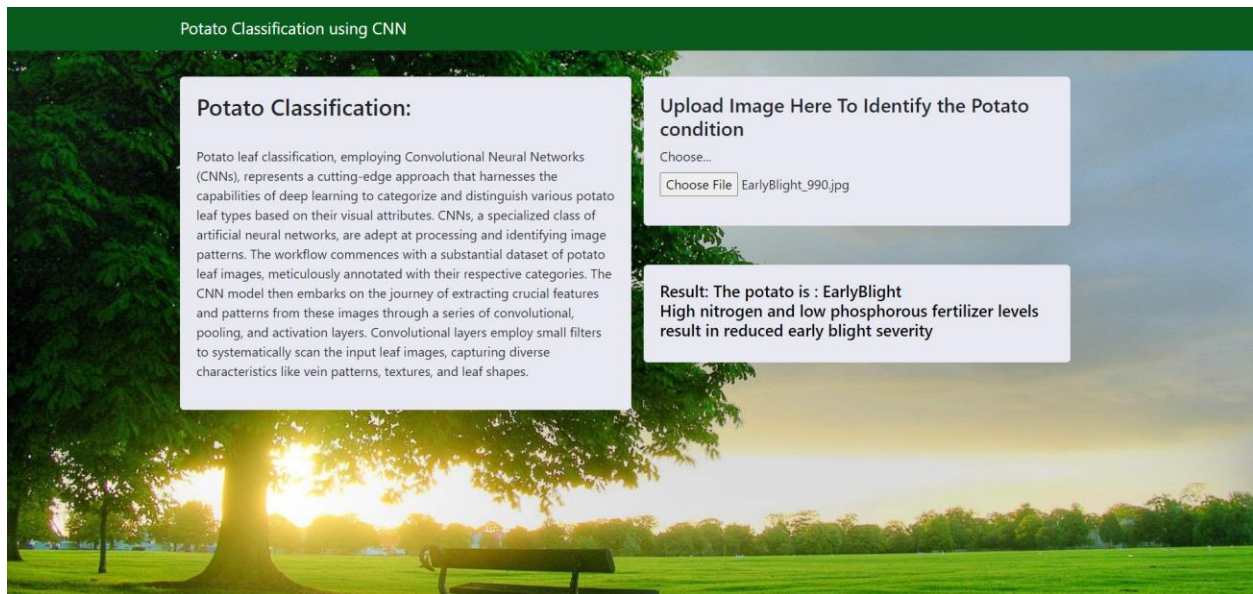
Already have an account? [Login](#)

Website



Choose the image to upload





Conclusion

The development of a potato leaf disease classification system using deep learning, particularly Convolutional Neural Networks (CNNs), offers a promising solution for agriculture. This system's ability to swiftly and accurately identify diseases in potato plants provides early detection, which can lead to increased crop yields, reduced economic losses, and enhanced food security. The user-friendly interface and feedback mechanisms make this technology accessible to a diverse user base, including non-technical users. However,

challenges related to data quality, model complexity, interpretability, and ongoing maintenance must be carefully addressed to ensure the system's effectiveness. In conclusion, this project has the potential to significantly benefit farmers and the broader agricultural industry when implemented responsibly and thoughtfully.

Future work

Expanded Crop Coverage: While the current focus is on potato diseases, the underlying deep learning framework can be extended to classify diseases in various other crops. This scalability can benefit a wide range of agricultural practices and contribute to crop health management.

AI-Driven Precision Agriculture: Incorporating this system into broader precision agriculture practices can optimize resource allocation, reducing the use of pesticides and promoting sustainable farming. The deep learning model can adapt to specific regional conditions and crop varieties.

IoT Integration: Integration with the Internet of Things (IoT) devices, such as sensors and automated irrigation systems, can enhance the system's capabilities. These devices can collect real-time data, further improving disease detection and prevention.

AI-Enhanced Recommendations: The system can evolve to not only classify diseases but also provide recommendations for disease management strategies, including the selection of appropriate pesticides, organic treatments, and preventive measures.

Mobile Apps and Accessibility: Development of dedicated mobile applications can increase accessibility, enabling farmers in remote areas to utilize the technology for disease identification and management.

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Complete Implementation

Importing the required libraries

```
import tensorflow as tf
from tensorflow.keras import models, layers
import matplotlib.pyplot as plt
import numpy as np
import keras
import os
from tensorflow.keras.preprocessing.image import ImageDataGenerator, load_img
```

Code for data augmentation

```
# Define directories for train, test, and validation sets
train_dir = '/kaggle/input/disease/Dataset/train'
test_dir = '/kaggle/input/disease/Dataset/test'
validation_dir = '/kaggle/input/disease/Dataset/val'

# Define parameters for image preprocessing and augmentation
batch_size = 32
target_size = (224, 224) # Change according to your image size requirements

# Create ImageDataGenerator for train set with augmentation
train_augmentation_datagen = ImageDataGenerator(
    rescale=1./255, # Normalize pixel values
    rotation_range=40,
    width_shift_range=0.2,
    height_shift_range=0.2,
    shear_range=0.2,
    zoom_range=0.2,
    horizontal_flip=True,
    fill_mode='nearest'
)

# Create ImageDataGenerator for test and validation sets (only normalization)
test_validation_datagen = ImageDataGenerator(rescale=1./255)
```

Images in the dataset

```
def plot_sample_images(directory, num_images=5):
    class_dirs = [os.path.join(directory, cls) for cls in os.listdir(directory)]

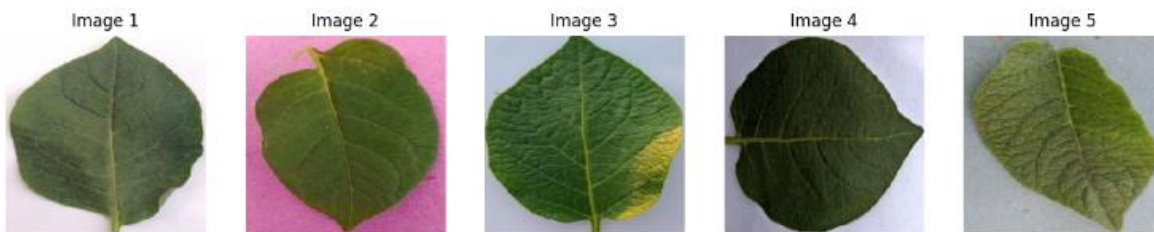
    # Plot sample images from each class directory
    for cls_dir in class_dirs:
        class_name = os.path.basename(cls_dir)
        fig, axes = plt.subplots(1, num_images, figsize=(15, 4))
        fig.suptitle(f'Sample Images from Class: {class_name}')

        # Load and plot sample images
        image_files = [os.path.join(cls_dir, file) for file in os.listdir(cls_dir)[:num_images]]
        for i, image_file in enumerate(image_files):
            image = load_img(image_file, target_size=(224, 224)) # Load and resize image
            axes[i].imshow(image)
            axes[i].set_title(f'Image {i+1}')
            axes[i].axis('off')
        plt.show()
    plot_sample_images(train_dir)
```

Sample Images from Class: EarlyBlight



Sample Images from Class: Healthy

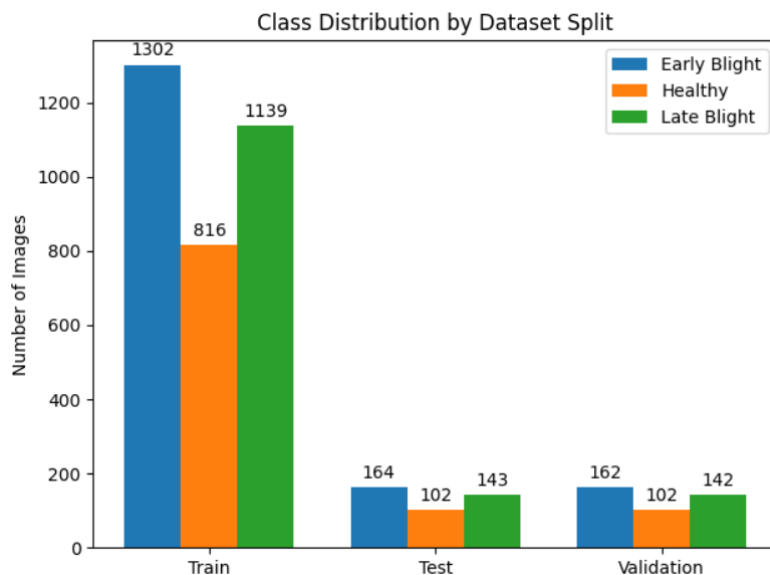


Sample Images from Class: LateBlight



Split ratio between train, test and validation

```
import os
import numpy as np
import matplotlib.pyplot as plt
def count_images_in_classes(directory):
    num_earlyblight = len(os.listdir(os.path.join(directory, 'EarlyBlight')))
    num_healthy = len(os.listdir(os.path.join(directory, 'Healthy')))
    num_lateblight = len(os.listdir(os.path.join(directory, 'LateBlight')))
    return num_earlyblight, num_healthy, num_lateblight
# Define paths to your directories (replace with your actual paths)
train_dir = '/kaggle/input/disease/Dataset/train'
test_dir = '/kaggle/input/disease/Dataset/test'
validation_dir = '/kaggle/input/disease/Dataset/val'
# Count images in train, test, and validation sets
train_earlyblight, train_healthy, train_lateblight = count_images_in_classes(train_dir)
test_earlyblight, test_healthy, test_lateblight = count_images_in_classes(test_dir)
validation_earlyblight, validation_healthy, validation_lateblight = count_images_in_classes(validation_dir)
# Create bar plot for class distribution
labels = ['Train', 'Test', 'Validation']
earlyblight_counts = [train_earlyblight, test_earlyblight, validation_earlyblight]
healthy_counts = [train_healthy, test_healthy, validation_healthy]
lateblight_counts = [train_lateblight, test_lateblight, validation_lateblight]
x = np.arange(len(labels))
width = 0.25
fig, ax = plt.subplots()
rects1 = ax.bar(x - width, earlyblight_counts, width, label='Early Blight')
rects2 = ax.bar(x, healthy_counts, width, label='Healthy')
rects3 = ax.bar(x + width, lateblight_counts, width, label='Late Blight')
ax.set_ylabel('Number of Images')
ax.set_title('Class Distribution by Dataset Split')
ax.set_xticks(x)
ax.set_xticklabels(labels)
ax.legend()
def autolabel(rects):
    for rect in rects:
        height = rect.get_height()
        ax.annotate('{}'.format(height),
                    xy=(rect.get_x() + rect.get_width() / 2, height),
                    xytext=(0, 3),
                    textcoords="offset points",
                    ha='center', va='bottom')
autolabel(rects1)
autolabel(rects2)
autolabel(rects3)
fig.tight_layout()
plt.show()
```



Augmented Images after image augmentation

```
original_images, _ = next(train_augmentation_datagen.flow_from_directory(
    train_dir,
    target_size=target_size,
    batch_size=9,
    class_mode='binary',
    shuffle=False
))

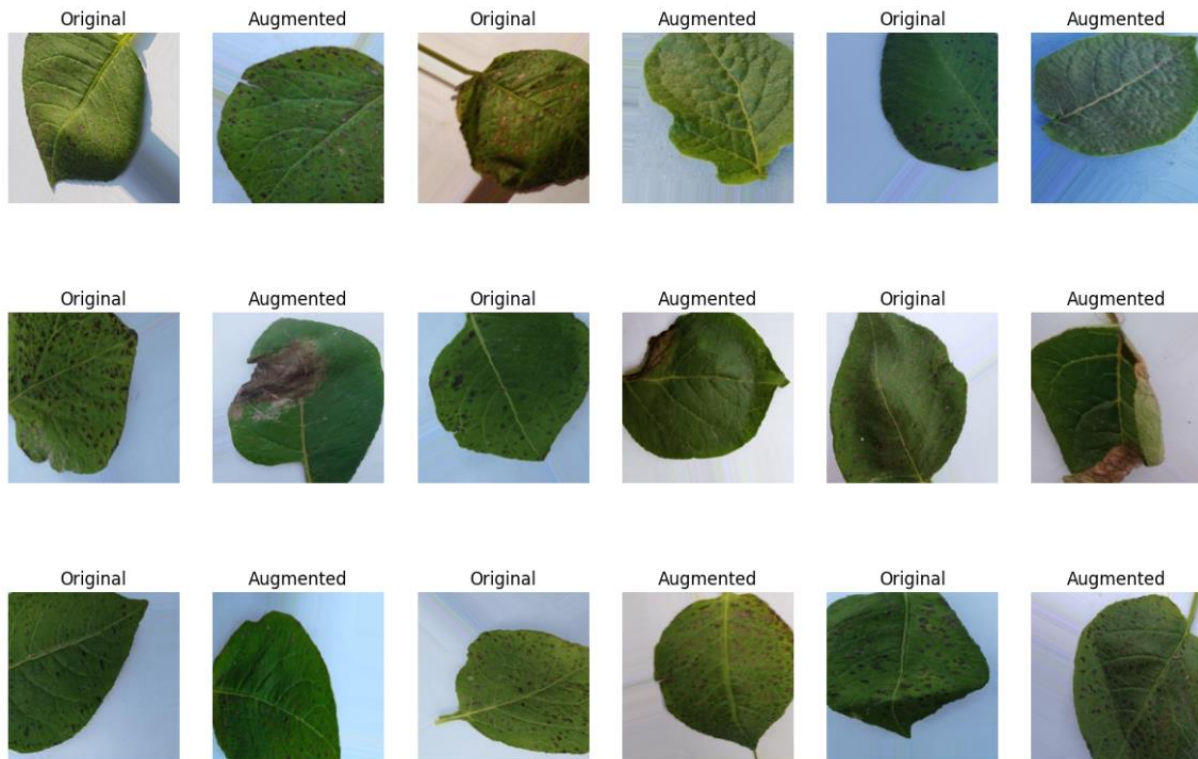
augmented_images, _ = next(train_augmentation_generator)

plt.figure(figsize=(15, 10))
for i in range(9):
    plt.subplot(3, 6, 2*i + 1)
    plt.imshow(original_images[i])
    plt.title('Original')
    plt.axis('off')

    plt.subplot(3, 6, 2*i + 2)
    plt.imshow(augmented_images[i])
    plt.title('Augmented')
    plt.axis('off')

plt.show()
```

Found 3257 images belonging to 3 classes.



ResNet50

```
from tensorflow.keras.applications import ResNet50
from tensorflow.keras.models import Model
from tensorflow.keras.layers import Flatten, Dense
from tensorflow.keras.optimizers import Adam
from keras.models import Sequential
from tensorflow.keras.layers import BatchNormalization
```

+ Code + Markdown

```
resnet_model = Sequential()
pretrained_model = ResNet50(include_top=False,
                             input_shape=(240,240,3),
                             pooling='avg',
                             weights='imagenet')
for layer in pretrained_model.layers:
    layer.trainable=False
resnet_model.add(pretrained_model)
resnet_model.add(Flatten())
resnet_model.add(BatchNormalization())
resnet_model.add(Dense(512, activation='relu'))
resnet_model.add(BatchNormalization())
resnet_model.add(Dense(4, activation='softmax'))
```

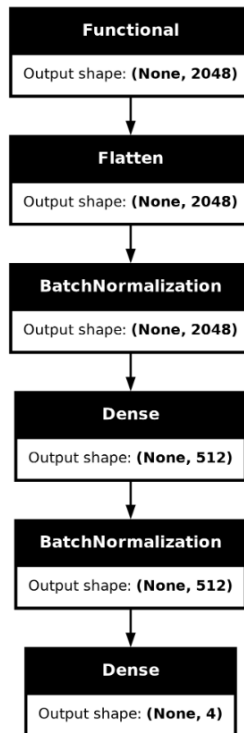
Downloading data from https://storage.googleapis.com/tensorflow/keras-applications/resnet/resnet50_weights_tf_dim_ordering_tf_kernels_notop.h5
94765736/94765736 0s 0us/step

Training Accuracy

```
history = resnet_model.fit(training_data,
                            steps_per_epoch=len(training_data),
                            epochs=20,
                            validation_data=validation_data,
                            validation_steps=len(validation_data))
```

```
Epoch 1/20
WARNING: All log messages before absl:InitializeLog() is called are written to STDERR
I0000 00:00:17.173730164.597369 124 device_compiler.h:106] Compiled cluster using XLA! This line is logged at most once for the lifetime of the process.
W0000 00:00:17.173730164.646521 124 graph_launch.cc:671] fallback to op-by-op mode because mnsnet node breaks graph update
50/50 ----- 0s 392ms/step - accuracy: 0.0519 - loss: 0.4916
W0000 00:00:17.173730184.000069 124 graph_launch.cc:671] fallback to op-by-op mode because mnsnet node breaks graph update
W0000 00:00:17.173730189.073345 126 graph_launch.cc:671] fallback to op-by-op mode because mnsnet node breaks graph update
50/50 ----- 0s 773ms/step - accuracy: 0.0531 - loss: 0.4880 - val_accuracy: 0.9492 - val_loss: 0.1593
Epoch 2/20
50/50 ----- 0s 1ms/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
Epoch 3/20
W0000 00:00:17.173730202.638199 124 graph_launch.cc:671] fallback to op-by-op mode because mnsnet node breaks graph update
repeat() function when building your dataset.
self.gen.throw(typ, value, traceback)
50/50 ----- 13s 245ms/step - accuracy: 0.9757 - loss: 0.0783 - val_accuracy: 0.9492 - val_loss: 0.1595
Epoch 4/20
50/50 ----- 0s 389us/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
Epoch 5/20
50/50 ----- 13s 252ms/step - accuracy: 0.9909 - loss: 0.0341 - val_accuracy: 0.9639 - val_loss: 0.1195
Epoch 6/20
50/50 ----- 0s 316us/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
Epoch 7/20
50/50 ----- 13s 258ms/step - accuracy: 0.9961 - loss: 0.0162 - val_accuracy: 0.9639 - val_loss: 0.1055
Epoch 8/20
50/50 ----- 0s 540us/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
Epoch 9/20
50/50 ----- 13s 257ms/step - accuracy: 0.9976 - loss: 0.0112 - val_accuracy: 0.9623 - val_loss: 0.0973
Epoch 10/20
50/50 ----- 0s 321us/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
Epoch 11/20
50/50 ----- 13s 250ms/step - accuracy: 0.9945 - loss: 0.0129 - val_accuracy: 0.9609 - val_loss: 0.0903
Epoch 12/20
50/50 ----- 0s 630us/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
Epoch 13/20
50/50 ----- 13s 247ms/step - accuracy: 0.9982 - loss: 0.0067 - val_accuracy: 0.9721 - val_loss: 0.0870
Epoch 14/20
50/50 ----- 0s 309us/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
Epoch 15/20
50/50 ----- 13s 240ms/step - accuracy: 0.9950 - loss: 0.0165 - val_accuracy: 0.9689 - val_loss: 0.0939
Epoch 16/20
50/50 ----- 0s 309us/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
Epoch 17/20
50/50 ----- 13s 249ms/step - accuracy: 0.9974 - loss: 0.0081 - val_accuracy: 0.9689 - val_loss: 0.0977
Epoch 18/20
50/50 ----- 0s 438us/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
Epoch 19/20
50/50 ----- 13s 253ms/step - accuracy: 0.9994 - loss: 0.0052 - val_accuracy: 0.9689 - val_loss: 0.0970
Epoch 20/20
50/50 ----- 0s 308us/step - accuracy: 0.0000e+00 - loss: 0.0000e+00 - val_accuracy: 0.0000e+00 - val_loss: 0.0000e+00
```

Layers added to the pretrained model

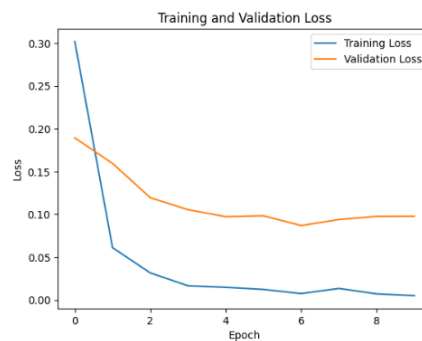


Plotting the Training and Validation Accuracy and Training and Validation Loss

```
import matplotlib.pyplot as plt

# Plot every other accuracy for training and validation
plt.plot(history.history['accuracy'][:,::2], label='Training Accuracy')
plt.plot(history.history['val_accuracy'][:,::2], label='Validation Accuracy')
plt.xlabel('Epoch')
plt.ylabel('Accuracy')
plt.title('Training and Validation Accuracy')
plt.legend()
plt.show()

# Plot every other loss for training and validation
plt.plot(history.history['loss'][:,::2], label='Training Loss')
plt.plot(history.history['val_loss'][:,::2], label='Validation Loss')
plt.xlabel('Epoch')
plt.ylabel('Loss')
plt.title('Training and Validation Loss')
plt.legend()
plt.show()
```



Testing Accuracy

```
resnet_model.evaluate(validation_data)
```

```
9/9 ----- 2s 201ms/step - accuracy: 0.9800 - loss: 0.0622  
[0.10186779499053955, 0.9688524603843689]
```

Manually testing the images

```
from tensorflow.keras.preprocessing import image  
import numpy as np  
  
# Load and preprocess the image  
img = image.load_img('/kaggle/input/potato/Dataset/EarlyBlight/EarlyBlight_1.jpg', target_size=(240, 240))  
img_array = image.img_to_array(img)  
img_array = np.expand_dims(img_array, axis=0) # Add a batch dimension  
  
# Make predictions  
predictions = resnet_model.predict(img_array) # Assuming 'model' is your trained VGG19 model  
  
# Get the class with the highest probability  
predicted_class_index = np.argmax(predictions)  
  
# Define your class labels  
class_labels = ['EarlyBlight', 'Healthy', 'LateBlight'] # Update with your actual class labels  
  
# Get the predicted class label  
predicted_class = class_labels[predicted_class_index]  
  
print("Predicted class:", predicted_class)
```

```
1/1 ----- 6s 6s/step  
Predicted class: EarlyBlight
```

Performance Metrics

```
import numpy as np  
from sklearn.metrics import accuracy_score, precision_score, recall_score, f1_score, roc_auc_score, confusion_matrix, accuracy_score  
from tensorflow.keras.preprocessing.image import ImageDataGenerator  
image_size = (240, 240)  
batch_size = 70  
validation_generator = ImageDataGenerator().flow_from_directory(  
    '/content/Dataset/val',  
    target_size=image_size,  
    batch_size=batch_size,  
    shuffle=False  
)  
  
y_pred_prob = resnet_model.predict(validation_generator)  
y_pred = np.argmax(y_pred_prob, axis=1)  
y_true = validation_generator.classes  
precision = precision_score(y_true, y_pred, average='weighted')  
print("Precision:", precision)  
recall = recall_score(y_true, y_pred, average='weighted')  
print("Recall:", recall)  
f1 = f1_score(y_true, y_pred, average='weighted')  
print("F1 Score:", f1)  
acc=accuracy_score(y_true,y_pred)  
print("Accuracy:",acc)  
conf_matrix = confusion_matrix(y_true, y_pred)  
print("\nConfusion Matrix:")  
print(conf_matrix)  
import matplotlib.pyplot as plt  
import seaborn as sns  
plt.figure(figsize=(10, 8))  
sns.heatmap(conf_matrix, annot=True, fmt="d", cmap="Blues", cbar=False)  
plt.title("Confusion Matrix")  
plt.xlabel("Predicted Label")  
plt.ylabel("True Label")  
plt.show()
```

Found 815 images belonging to 3 classes.

12/12 [=====] - 4s 318ms/step

Precision: 0.9443536402437807

Recall: 0.943558282208589

F1 Score: 0.9433615305057043

Accuracy: 0.943558282208589

Confusion Matrix:

```
[[322   2   2]
 [  9 189   6]
 [  9  18 258]]
```

