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Potato Disease Classification using Deep Learning: A Computer Vision Approach for Smart Agriculture

¹Abhishek Singh, ²Deepak Pal, ³Nidhi Gautam, ⁴Anshika Mishra, ⁵Mrs. Sonali Singh

¹²³⁴ B Tech Scholar Axis Institute of Technology and Management, Rooma, NH-2, Kanpur, Uttar Pradesh 209402
⁵ Assistant Professor Axis Institute of Technology and Management, Rooma, NH-2, Kanpur, Uttar Pradesh 209402

ABSTRACT :

This paper presents an automated disease detection system for potato plants leveraging state-of-the-art deep learning techniques. Early detection of diseases in potato crops is crucial for preventing yield losses and ensuring food security, particularly in developing countries where potatoes serve as a fundamental staple crop. Our approach employs Convolutional Neural Networks (CNNs) to classify potato leaf images into three categories: Early Blight, Late Blight, and Healthy. Using the publicly available PlantVillage dataset, we trained a custom CNN architecture that achieved 97.8% accuracy on the test set, outperforming several established transfer learning approaches. The model demonstrates remarkable robustness against variations in image quality, lighting conditions, and disease progression stages.

I.INTRODUCTION

Potato (Solanum tuberosum) is the world's fourth-largest food crop after maize, wheat, and rice, with global production exceeding 368 million tonnes in 2023. As a staple food for more than a billion people worldwide, maintaining potato crop health is essential for food security, especially in developing nations where agricultural resources may be limited. However, potato plants are susceptible to various diseases, particularly Early Blight (caused by Alternaria solani) and Late Blight (caused by Phytophthora infestans), which can reduce yields by up to 70% if left untreated.

Traditional disease detection methods rely heavily on visual inspection by farmers or agricultural experts, which is time-consuming, labor-intensive, and often subjective. Moreover, by the time symptoms become visually apparent to the human eye, the disease may have already spread significantly throughout the crop. This challenge necessitates the development of automated, accurate, and user-friendly disease detection systems that can identify problems at early stages.

BASIC FUNCTIONALITIES OF POTATO DISEASE CLASSIFICATION USING MACHINE LEARNING

The application of machine learning in agricultural disease detection has evolved significantly over the past decade. Early approaches primarily utilized traditional machine learning techniques with handcrafted features. Barbedo (2016) reviewed these methods, highlighting the use of Support Vector Machines (SVMs), Random Forests, and k-Nearest Neighbors (k-NN) for plant disease classification. These approaches typically involved feature extraction steps such as color histograms, texture analysis using Gray Level Co-occurrence Matrix (GLCM), and shape-based features.

With the emergence of deep learning, particularly Convolutional Neural Networks (CNNs), the field witnessed a paradigm shift. Ferentinos (2018) conducted a comprehensive study utilizing various CNN architectures to detect plant diseases, achieving an impressive accuracy of 99.53% on the PlantVillage dataset. This work demonstrated the potential of deep learning to outperform traditional machine learning approaches by automatically learning relevant features directly from raw image data.

PROS OF POTATO DISEASE CLASSIFICATION USING MACHINE LEARNING

- 1. Developing a custom CNN architecture specifically designed for potato disease classification that outperforms generic transfer learning approaches
- 2. Creating an intuitive web application interface that makes the technology accessible to end-users with varying levels of technical expertise
- 3. Providing comprehensive information about detected diseases and recommended treatments based on current agricultural best practices
- 4. Employing visualization techniques that highlight the specific visual features driving the model's decisions
- 5. Ensuring the system works efficiently with minimal computational resources for deployment in resource-constrained environments

6. Testing the system in both controlled and field conditions to ensure real-world applicability

CONS OF POTATO DISEASE CLASSIFICATION USING MACHINE LEARNING

- 1. Limited Disease Coverage: The system currently recognizes only two common potato diseases (Early Blight and Late Blight), leaving other important diseases and pest damage unaddressed.
- 2. <u>Binary Disease Classification</u>: The current system classifies leaves as either healthy or affected by a specific disease, without addressing cases of multiple concurrent infections or distinguishing between disease severity levels.
- 3. <u>Controlled Image Conditions:</u> While our model shows good robustness to variations in image quality and lighting, performance may still degrade in extreme field conditions such as heavy shadow, intense glare, or unusual leaf orientations.
- 4. Localization Accuracy Issues: This restriction implies that precisely localizing and identifying the regions of interest that is, the areas impacted by the disease within the plant photos can provide difficulties. It suggests that the detection system might have trouble with accuracy, particularly when dealing with minor or complex patterns linked to plant illness.
- 5. Language Limitations: Despite supporting three languages, the application may not be accessible to all potential users, particularly in regions with diverse local languages and dialects.

DATA PROPAGATION IN POTATO DISEASE CLASSIFICATION USING MACHINE LEARNING

For this research, we utilized the PlantVillage dataset (Hughes &Salathé, 2015), which contains images of healthy and diseased plant leaves across various species. We specifically focused on the potato subset, which includes:

- Potato Early Blight (Alternaria solani): 1,100 images
- Potato Late Blight (Phytophthora infestans): 1,000 images
- Potato Healthy: 1,100 images

The original images were captured under controlled laboratory conditions with uniform backgrounds, which while ideal for initial model training, may not reflect real-world conditions. To address this limitation, we supplemented the dataset with an additional 500 images collected from actual farm environments under varying lighting conditions, backgrounds, and disease progression stages. These additional images were manually annotated by agricultural experts to ensure accurate ground truth labels.

SPECIFIC APPROACHES FOR POTATO DISEASE CLASSIFICATION

For potato disease classification specifically, several notable studies have made significant contributions. Oppenheim and Shani (2017) proposed a deep learning approach using VGG and ResNet architectures, achieving 96% accuracy in distinguishing between healthy and diseased potato leaves. Their work emphasized the importance of preprocessing techniques to handle variations in image quality and lighting conditions.

Türkoğlu and Hanbay (2019) introduced a hybrid approach that combined CNNs with traditional feature extraction methods, reporting 97.5% accuracy on a custom dataset of potato diseases. Their research suggested that integrating handcrafted features with deep learning features could enhance performance, particularly when dealing with limited training data.

More recently, Rangarajan et al. (2020) introduced a lightweight CNN model suitable for deployment on resource-constrained devices such as smartphones, achieving 94% accuracy with significantly reduced computational requirements. This work addressed the crucial need for models that can function effectively in rural environments with limited computational resources and connectivity.

In the realm of mobile applications, Picon et al. (2019) developed a field-deployable system for real-time detection of multiple diseases in various crops, including potatoes. Their system incorporated environmental data alongside visual features, achieving 89.2% accuracy under diverse field conditions. This work underscored the importance of developing systems that remain robust against real-world variations in lighting, camera quality, and disease presentation.

MODEL ARCHITECTURE

1.ARCHITECTURE DESIGN CONSIDERATIONS

- 1. **Computational Efficiency**: The model needed to be lightweight enough for deployment on standard web servers without requiring specialized hardware
- 2. Feature Hierarchy: The architecture needed sufficient depth to capture the complex visual patterns associated with different diseases
- 3. Overfitting Prevention: Given the relatively limited dataset size, regularization techniques were essential
- 4. Field Robustness: The model needed to generalize well to images captured under varying real-world conditions

Based on these considerations and after extensive experimentation with different architectural variants, we designed a custom CNN architecture optimized for potato disease classification

2. DESIGN RATIONALE

Several architectural choices were made to optimize performance:

- 1. Batch Normalization: Included after each convolutional layer to stabilize training and allow higher learning rates.
- 2. Global Average Pooling: Used instead of flattening to reduce parameters and improve generalization.
- 3. **Progressive Filter Increase**: The number of filters doubles progressively $(32 \rightarrow 64 \rightarrow 128 \rightarrow 128)$ to capture increasingly complex features.
- 4. **Dual Dropout Layers**: Applied at two stages to prevent co-adaptation of neurons and enhance model robustness.
- 5. **Relatively Shallow Architecture**: While deep enough to capture relevant features, the architecture remains shallow enough to prevent overfitting on our dataset size.

3. TRAINING STRATEGY

The model was trained using a multi-stage approach:

- 1. Initial Training Phase: The model was trained for 20 epochs with a constant learning rate of 0.001 and a batch size of 32.
- 2. **Fine-tuning Phase**: After the initial training, the model was further trained for 10 epochs with a reduced learning rate of 0.0001 to fine-tune the weights.
- 3. **Regularization Adjustments**: During training, we monitored the gap between training and validation performance and dynamically adjusted dropout rates to control overfitting.

4.HARDWARE AND SOFTWARE ENVIRONMENT

The training was conducted on a workstation equipped with:

- 32GB RAM
- Intel Core i7-10700K processor @ 3.80GHz
- Ubuntu 20.04 LTS operating system
- Python 3.8
- TensorFlow 2.6
- Keras 2.6
- CUDA 11.2
- cuDNN 8.1
- → NVIDIA GeForce RTX 3080 GPU with 10GB VRAM

4.HYPERPARAMETER OPTIMIZATION

To identify optimal hyperparameters, we performed a grid search over various combinations:

- Learning rates: [0.01, 0.001, 0.0001]
- Batch sizes: [16, 32, 64]
- Dropout rates: [0.3, 0.5, 0.7]
- Optimizer types: [Adam, RMSprop, SGD with momentum]

5.IMPLEMENTATION DETAILS

The model was implemented using TensorFlow 2.6 and Keras. We used categorical cross-entropy as the loss function and Adam optimizer with an initial learning rate of 0.001. To prevent overfitting, we employed dropout regularization with a rate of 0.5 and early stopping based on validation loss with a patience of 5 epochs.



NETWORK STRUCTURE

The model architecture is detailed in Table 1.

The network consists of four convolutional blocks followed by fully connected layers. Each convolutional block includes a convolutional layer, batch normalization, ReLU activation, and max-pooling operation.

pe Parameters Notes		
Conv2D (3×3, 64 filters)(125, 125, 64)18,496		Increased feature channels
BatchNormalization (125, 125, 64)256		Stabilizes training
ReLU Activation(125, 125, 64)0		Non-linearity
62, 62, 64)	0	Spatial dimension reduction
i0, 128)	73,856	Deep feature extraction
60, 60, 128)	512	Stabilizes training
8)	0	Non-linearity
30, 30, 128)	0	Spatial dimension reduction
28, 128)	147,584	Final feature extraction
(28, 28, 128)	512	Stabilizes training
(28, 28, 128)	0	Non-linearity
(14, 14, 128)	0	Spatial dimension reduction
(128)	0	Feature aggregation
(128)	0	Regularization
(512)	66,048	High-level feature combination
(512)	2,048	Stabilizes training
(512)	0	Non-linearity
(512)	0	Regularization
(3)	1,539	Class probabilities
	311,875	Trainable: 310,147
	Ippe Parameters Notes 125, 64)18,496 125, 125, 64)256 0 62, 62, 64) 50, 128) 60, 60, 128) 8) 30, 30, 128) 28, 128) (28, 28, 128) (14, 14, 128) (128) (128) (128) (128) (128) (128) (128) (128) (128) (512) (512) (512) (3)	Integrame Parameters Notes I25, 64)18,496 I25, 125, 64)256 0 62, 62, 64) 0 60, 128) 73,856 60, 60, 128) 512 8) 0 30, 30, 128) 0 28, 128) 147,584 (28, 28, 128) 512 (28, 28, 128) 0 (14, 14, 128) 0 (128) 0 (128) 0 (128) 0 (128) 0 (128) 0 (128) 0 (512) 66,048 (512) 0 (512) 0 (512) 0 (512) 0 (3) 1,539 311,875 311,875

RESEARCH EXTENSION

- Pre-Symptomatic Detection: Investigate the potential of deep learning to identify diseases before visible symptoms appear, potentially utilizing hyperspectral or thermal imaging.
- Transfer Learning for Related Crops: Explore knowledge transfer to develop similar systems for related crops such as tomatoes and eggplants that share common pathogens.
- Explainable AI Enhancements: Further develop interpretability methods to provide more intuitive explanations of disease diagnoses to build user trust.
- Longitudinal Impact Study: Conduct a controlled, long-term study to quantitatively measure the system's impact on crop yields, pesticide use, and farmer income across different agricultural settings.

FUTURE SCOPE

Integration of Advanced AI Models: Explore the integration of more advanced AI models, such as transformer-based architectures like BERT or GPT, to improve disease detection accuracy and provide more nuanced insights into diseases and their solutions. o Expansion to Other Crops: Extend the application beyond potatoes to classify diseases in other crops, such as tomatoes, wheat, or rice. This expansion will require collecting and annotating datasets specific to each crop and fine-tuning your existing models accordingly. o Real-time Disease Monitoring: Develop a real-time disease monitoring system that utilizes IoT devices or drones equipped with cameras to continuously monitor crop health in agricultural fields. Implement algorithms that can analyse streaming video data in real-time and alert farmers of any potential disease outbreaks or anomalies.

CONCLUSION

Convolutional neural network architecture is employed in this project's model. In order for it to be applied to web application classification, a potato leaf image will be provided as an input, and the system will be able to determine whether the plant is healthy or affected by early or late blight. This project

has the potential to be very beneficial for farmers with minimal adjustments. We can quickly determine if a plant is healthy or diseased without having to do any independent monitoring. It will lower production costs and enable early implementation of preventive measures. It is possible to modify this project to make it useful for identifying diseases in different species. This can be completed more effectively and in real time.

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