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# Intelligent Cotton Leaf Disease Detection System using Deep Learning and Multi-Layer Validation Framework

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**Abstract:** *Crop diseases in cotton pose a significant economic threat to agriculture across the globe. Most of the diseases can cause a reduction in crop yield of up to 30–40%. For the agriculture sector to thrive, a robust system is required to diagnose and treat crop diseases efficiently. Current approaches rely on manual approaches and are time-consuming, error-prone and require the specialist expertise of agronomists. In this paper, we present a full-stack webbased system that includes deep learning classification for disease detection and employs a novel multi-layer validation system. The classification system utilises a deep Convolutional Neural Network and achieves up to 87.5% classification accuracy across ten diseases including the difficult to diagnose Fusarium Wilt, Bacterial Blight and aphids infestation. Our system employs a dual-validation system where rule-based image analysis is first used to verify if an image is of a cotton leaf, followed by AI-powered verification of the disease. Our system achieves 98% rejection accuracy for non-cotton images and 95% acceptance accuracy for real cotton leaf images. It includes a Flask-based REST API backend that utilises strict validation protocols and evaluates 16 distinct parameters from an image, including green color dominance, skin tone detection, edge density and texture uniformity. A bilingual treatment recommendation system provides on-the-spot treatment advice in both English and Hindi. The end-to-end system requires no specialist hardware, enabling specialist disease diagnosis to be carried out on a web browser, making it accessible to farmers worldwide.*

**Keywords:** *Deep learning, cotton disease detection, convolutional neural networks, precision agriculture, image validation, web-based diagnosis, Flask API, agricultural AI.*

## I. INTRODUCTION

Cotton is one of the world's most economically important fibre crop and is cultivated in 70+ countries of world with production level of more than 25 million metric tons every year. In addition to various phytopathogens, a number of pests are affecting cotton seriously and are impacting crop yield and fibre quality. Losses to major cotton-producing regions are in excess of \$3 billion per year [1].

Present approaches to identify plant diseases rely on visual identification by trained agricultural experts. These traditional methods carry several caveats, such as variability in identification, the need for specialized knowledge, time and resources required for large scale field surveys, and the lag time between surveys and decision making that allows for disease progression. There is a compelling need for automated, objective, simple to use plant disease diagnostics that function accurately and at scale.

Plant disease detection in agriculture has recently gained much attention using deep learning, and Computer Vision techniques. While such systems have achieved state-of-the-art results, current agricultural disease detectors suffer from several critical issues, for example, they have a high false detection rate on non-plant images; they are unable to differentiate between different species of plants; they are not sufficiently validated; or they are inaccessible to farmers.

This research addresses these gaps by presenting a comprehensive cotton disease detection system with the following key contributions:

- 1) Development of a novel dual-layer validation method comprising rule-based image analysis (based on 16 parameters) and AI-powered verification of cotton leaves, attaining 98% rejection rate for non-cotton samples.
- 2) Deep CNN architecture trained to recognize 10 cotton diseases, with high accuracy obtained using data augmentation techniques reaching 87.5% classification.
- 3) Full-stack web application capable of disease diagnosis in real time, developed using RESTful API architecture accessible through standard web browser interfaces.

- 4) New bilingual treatment recommendation engine populated with recommendations in English and Hindi languages for healthcare practitioners to provide appropriate care to patients.
- 5) Comprehensive validation system which compares features of objects in the image, including color distribution (HSV/YCrCb) and edge parameters such as edge density and texture, as well as shape features (morphology).

The remainder of this paper is organized as follows: Section II reviews related work in plant disease detection systems. Section III describes the proposed methodology including system architecture, validation framework, and deep learning model. Section IV presents implementation details and experimental setup. Section V discusses results and performance evaluation. Section VI concludes with future research directions.

## II. LITERATURE REVIEW

### A. Plant Disease Detection Using Deep Learning

Deep learning approaches have revolutionized agricultural disease detection in recent years. Mohanty et al. [2] developed a deep CNN trained on the PlantVillage dataset containing 54,306 images across 38 disease categories, achieving 99.35% accuracy. However, their system lacked robustness when tested on field images with complex backgrounds and varying lighting conditions.

Ferentinos [3] proposed a comparative study of CNN architectures (AlexNet, VGG, ResNet) for plant disease identification across 25 plant species, reporting best accuracy of 99.53% using VGG architecture. The study acknowledged significant performance degradation when processing images outside the training distribution, highlighting the need for robust validation mechanisms.

Amara et al. [8] proposed a deep learning-based approach that utilised CNN to achieve high-accuracy plant disease classification on leaf disease images. However, issues regarding robustness to diverse field conditions were pointed out.

Too et al. compared the performance of some of the deep learning architectures such as VGG16, ResNet and DenseNet for plant disease detection using fine-tuned models, indicating that there was significant improvement in the classification accuracy for plant disease detection using transfer learning approach [9].

### B. Cotton-Specific Disease Detection

In addition to work specifically focused on cotton, there is a growing interest in cotton diseases through image classification and plant disease diagnostics. Xie, et al. [4] recently used improved AlexNet deep learning architecture to accurately identify cotton diseases achieving 93.6% accuracy between 4 different disease classifications. Their work did however fall short in two key areas: the number of images used to train and validate the model and field validation on real world samples. The model was trained and validated using a dataset consisting of only 1,200 images.

Ramesh and Vydeki in [5] proposed cotton leaf spot diseases recognition and classification approach based on edge detection and multiclass SVM method, they get accuracy rate 94% on LBM database.

### C. Image Validation and Quality Assessment

In existing literature on image quality assessment, approaches typically focus on measures of blur, noise, and resolution. Zhang et al. in [6] approached no-reference image quality assessment using a Convolutional Neural Network (CNN), a shift from traditional full-reference and reduced-reference methods. However, to date, there are no approaches capable of plant species classification or validating leaf images.

Although Kumar et al. [7] reported 82% classification accuracy of leaf images using color histogram, they experienced high false-positives in practical implementations because the methodology failed to distinguish between different species of plants.

### D. Research Gaps

Analysis of existing literature reveals several critical gaps:

- 1) No robust validation mechanisms to filter out non-target images before disease classification.
- 2) Few trials in real-world environments, specifically addressing accessibility for farming communities.
- 3) Integrated validation using a combination of rule-based and AI approaches is not being given sufficient consideration.
- 4) Lack of fully integrated systems that perform a comprehensive diagnosis and give treatment recommendations.
- 5) Limited bilingual support for diverse agricultural regions

This research addresses these gaps through a comprehensive system design that prioritizes both accuracy and practical deployability in resource-constrained agricultural settings.

### III. METHODOLOGY

#### A. System Architecture

In this work we propose a three tiered system consisting of presentation layer (web interface), application layer (Flask API server) and data layer (trained models and validation rules) as shown in Figure 1.

The client side interface is built using HTML5, CSS3 and JavaScript allowing for optimal display on desktop and mobile browsers. Users can upload their cotton leaf images by simply dragging and dropping or selecting them from their hard drive. The client interface provides real time feedback to users during the validation and analysis processes.

This module consists of a server-side, flask-based application that has been implemented with RESTful APIs for image processing, validation and disease detection prediction. The images processed by the application undergo several layers of validation before being sent through to the deep learning model for disease classification.

#### B. Dual-Layer Validation Framework

A critical innovation in this system is the dual-layer validation framework designed to ensure only genuine cotton leaf images proceed to disease classification. This approach significantly reduces false positives and computational overhead.

1) *Layer 1: Rule-Based Validation:* The first validation layer implements 16 distinct image analysis parameters:

**Color Space Analysis:** Images are converted to multiple color spaces (HSV, YCrCb, LAB) for comprehensive color analysis. Green dominance is assessed by computing mean green channel values with threshold  $G_{mean} > 35$  to accept diseased or yellowing leaves. Green pixel percentage is calculated in HSV space using the criterion:

$$P_{green} = \frac{\sum_{i,j} M_{green}(i,j)}{W \times H} \times 100 > 12\% \quad (1)$$

where  $M_{green}(i,j)$  is a binary mask defined as:

$$\begin{aligned} &1 \text{ if } 20 \leq H(i,j) \leq 100 \\ &\text{and } S(i,j) > 10 \\ &\text{and } V(i,j) > 15 \\ &0 \text{ otherwise} \end{aligned} \quad (2)$$

**Skin Tone Detection:** To reject human images, skin detection is performed in YCrCb color space using empirically determined ranges:

$$S_{skin} = \frac{\sum_{i,j} M_{skin}(i,j)}{W \times H} \times 100 < 35 \quad (3)$$

where skin mask  $M_{skin}$  identifies pixels within ranges:  $Y \in [0, 255]$ ,  $Cr \in [133, 173]$ ,  $Cb \in [77, 127]$ .

**Texture Analysis:** Standard deviation of grayscale intensity measures texture complexity:

$$\sigma_{intensity} = \sqrt{\frac{1}{N} \sum_{i,j} (I(i,j) - \mu_I)^2} > 8 \quad (4)$$

where  $\mu_I$  is mean intensity. Low standard deviation indicates uniform backgrounds.

**Edge Density Computation:** Canny edge detection with adaptive thresholds identifies edge pixels. Edge density is calculated as:

$$D_{edge} = \frac{\sum_{i,j} E(i,j)}{W \times H} > 0.008 \quad (5)$$

where  $E(i,j)$  represents binary edge map from Canny detector with thresholds (30, 120).

**Additional Validation Parameters:** Resolution check (minimum 100×100 pixels), aspect ratio validation (0.254.0), brightness assessment (50-245 in V channel), saturation verification ( $\sigma_S > 15$ ), and gradient magnitude analysis.

2) *Layer 2: AI-Based Verification:* For images passing rulebased validation, an optional secondary CNN classifier distinguishes cotton leaves from other plant species. This binary classifier employs transfer learning from VGG16 pre-trained weights, fine-tuned on cotton versus non-cotton datasets. The classifier outputs probability  $P_{cotton}$  with acceptance threshold:

$$P_{cotton} = \sigma(W^T \phi(x) + b) > 0.7 \quad (6)$$

where  $\sigma$  is sigmoid activation,  $\phi(x)$  represents learned features, and  $W, b$  are learned parameters.

### C. Disease Classification CNN Architecture

The disease classification network employs a custom CNN architecture optimized for cotton disease patterns:

Architecture Design:

- Input layer: 224×224×3 RGB images
- Convolutional block 1: 32 filters (3×3), ReLU, BatchNorm, MaxPool (2×2)
- Convolutional block 2: 64 filters (3×3), ReLU, BatchNorm, MaxPool (2×2)
- Convolutional block 3: 128 filters (3×3), ReLU, BatchNorm, MaxPool (2×2)
- Convolutional block 4: 256 filters (3×3), ReLU, BatchNorm, MaxPool (2×2)
- Fully connected layer: 512 neurons, ReLU, Dropout (0.5)
- Fully connected layer: 256 neurons, ReLU, Dropout (0.3) • Output layer: 10 neurons, Softmax activation

The network classifies images into 10 categories: Aphids, Army worm, Bacterial Blight, Cotton Boll Rot, Curl Virus, Fusarium Wilt, Green Cotton Boll, Healthy, Powdery mildew, and Target Spot.

Training Strategy: The model is trained using categorical cross-entropy loss:

$$L = -\frac{1}{N} \sum_{i=1}^N \sum_{c=1}^C y_{ic} \log(\hat{y}_{ic}) \quad (7)$$

where  $y_{ic}$  is ground truth label,  $\hat{y}_{ic}$  is predicted probability for class  $c$  of sample  $i$ .

Optimization employs Adam optimizer with learning rate  $\alpha = 0.0001$ ,  $\beta_1 = 0.9$ ,  $\beta_2 = 0.999$ . Data augmentation includes rotation ( $\pm 40$ ), width/height shifts (30%), shear (30%), zoom (30%), horizontal/vertical flips, and brightness variation (0.8-1.2).

### D. Treatment Recommendation System

The system maintains a comprehensive disease-treatment knowledge base providing bilingual recommendations (English and Hindi) for each disease category. Recommendations are structured into immediate actions, prevention strategies, and application guidelines based on agricultural best practices and phytopathology research.

For each detected disease, the system calculates severity level:

$$\begin{aligned} \text{Severity} &= \text{Low if } P_{\text{disease}} < 60\% \\ \text{Severity} &= \text{Moderate if } 60\% \leq P_{\text{disease}} < 80\% \\ \text{Severity} &= \text{High if } P_{\text{disease}} \geq 80\% \end{aligned} \quad (8)$$

where  $P_{\text{disease}}$  is the model's confidence score for the predicted disease class.

## IV. IMPLEMENTATION

### A. Dataset Preparation

The training dataset comprises 12,500 images collected from multiple sources: field photographs from cotton farms across three geographical regions (2,500 images), publicly available cotton disease datasets (8,000 images), and augmented synthetic images (2,000 images). Each disease category contains minimum 800 training images and 200 validation images. Image acquisition followed standardized protocols: natural daylight photography, varied disease stages, multiple cotton varieties, and diverse environmental conditions. Images were manually annotated by agricultural experts and verified through two-stage quality control.

### B. Backend Implementation

The Flask backend (Python 3.8+) implements modular architecture with separation of concerns:

Core Dependencies: TensorFlow 2.15.0, OpenCV 4.8.0, Pillow 10.0.0, NumPy 1.24.0, Flask 2.3.0, Flask-CORS 4.0.0. API Endpoints:

- POST /api/predict - Disease prediction with validation
- GET /api/health - System health check
- GET / - Serves frontend interface

Image Processing Pipeline: Uploaded images undergo format conversion (RGB), resolution verification, validation execution, model preprocessing (resize 224×224, normalization [0,1]), and inference with confidence thresholding (minimum 45%).

**C. Frontend Implementation**

The responsive web interface employs vanilla JavaScript for API communication, HTML5 for semantic structure, CSS3 for styling, and Toastify.js for user notifications. The interface supports drag-and-drop upload, real-time preview, progress indication, and result visualization with treatment details.

**D. Deployment Configuration**

The system operates on lightweight infrastructure: Flask development server (production deployment recommended with Gunicorn/uWSGI), CPU-based inference (1-2 seconds per image), memory footprint 500MB including loaded models, and horizontal scaling capability through load balancing.

**V. RESULTS AND DISCUSSION**

**A. Validation Framework Performance**

The dual-layer validation framework was evaluated on 5,000 test images comprising 2,500 cotton leaf images and 2,500 non-cotton images (humans, objects, other plants, landscapes). **Layer 1 (Rule-Based) Performance:**

- True Positive Rate (cotton accepted): 96.8%
- True Negative Rate (non-cotton rejected): 97.2%
- False Positive Rate: 2.8%
- False Negative Rate: 3.2%
- Processing time: 0.15-0.25 seconds per image

The system successfully rejected 98.5% of human photographs through skin detection, 96.8% of objects through green pixel analysis, and 95.2% of other plants through combined validation parameters.

Common Rejection Reasons: Insufficient green color (42% of rejections), low edge density (28%), skin tone detection (18%), extreme aspect ratios (7%), low resolution (5%).

**B. Disease Classification Results**

The CNN classifier achieved the following performance metrics on 2,000-image test set:

Overall Metrics:

- Overall Accuracy: 87.5%
- Macro-averaged Precision: 86.3%
- Macro-averaged Recall: 85.8%
- Macro-averaged F1-Score: 86.0%
- Inference time: 1.2 seconds (CPU), 0.3 seconds (GPU)

Per-Class Performance: Healthy leaves achieved highest classification accuracy (92%) due to distinct visual characteristics.

Fusarium Wilt showed excellent performance (90% F1-score) attributed to characteristic vascular discoloration patterns.

**C. Confusion Matrix Analysis**

Common misclassifications occurred between visually similar disease categories. Army Worm was occasionally confused with Aphids (8% cases) due to similar damage patterns. Target Spot showed 6% confusion with Powdery Mildew in early disease stages.

TABLE I

DISEASE CLASSIFICATION PERFORMANCE

Disease	Precision	Recall	F1
Fusarium Wilt	0.92	0.89	0.90
Bacterial Blight	0.88	0.91	0.89
Aphids	0.85	0.87	0.86
Curl Virus	0.90	0.85	0.87
Powdery Mildew	0.84	0.86	0.85
Army Worm	0.82	0.80	0.81
Target Spot	0.87	0.84	0.85
Boll Rot	0.83	0.88	0.85
Green Boll	0.89	0.86	0.87
Healthy	0.93	0.92	0.92

TABLE II  
COMPARISON WITH RELATED WORK

System	Accuracy	Validation	Real-time
Xie et al. [4]	93.6%	No	No
Ramesh [5]	94.0%	No	No
Our System	87.5%	Yes (98%)	Yes

#### D. Comparative Analysis

While our disease classification accuracy (87.5%) is lower than some specialized systems, the complete solution including robust validation, real-time processing, and deployment readiness provides significant practical advantages.

#### E. System Usability Evaluation

Field testing with 50 farmers across two agricultural regions yielded positive feedback:

- 94% found the interface intuitive and easy to use
- 88% reported bilingual recommendations as highly valuable
- Average diagnosis time: 3.5 seconds from upload to results
- 92% expressed willingness to use the system regularly

#### F. Limitations and Challenges

Current limitations include: reduced accuracy for early stage disease symptoms, challenges with overlapping infections (multiple diseases), dependency on image quality and lighting conditions, and requirement for internet connectivity.

The system occasionally struggles with severely diseased leaves showing extensive necrosis, where green color percentage falls below validation thresholds, resulting in false rejections.

## VI. CONCLUSION

This research presents a comprehensive cotton disease detection system that addresses critical gaps in agricultural AI applications through innovative dual-layer validation, robust deep learning classification, and accessible web-based deployment. The system achieves 98% accuracy in filtering noncotton images while maintaining 87.5% disease classification accuracy across 10 disease categories. Key contributions include the novel multi-parameter validation framework combining rule-based and AI approaches, production-ready full-stack implementation with RESTful API architecture, bilingual treatment recommendation system, and demonstrated real-world usability in farming communities. Future research directions include: integration of temporal disease progression tracking through multiple image uploads, expansion to additional cotton varieties and geographical regions, development of mobile applications for offline operation, incorporation of environmental data (temperature, humidity) for enhanced prediction, and implementation of federated learning for continuous model improvement while preserving farmer privacy.

The system demonstrates that practical, accessible AI solutions for agriculture require careful attention to validation, deployment infrastructure, and end-user needs beyond pure classification accuracy. This holistic approach provides a blueprint for developing impactful agricultural AI systems.

## VII. ACKNOWLEDGMENT

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