

Comparative Analysis of MobileNetV2 and ResNet50 for Cotton Leaf Disease Classification using Deep Learning

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Abstract - Cotton is a vital cash crop in India, especially in Maharashtra, where it contributes significantly to the agricultural economy and textile industry. However, its productivity is frequently threatened by foliar diseases such as cotton leaf curl, bacterial blight, fusarium wilt, powdery mildew, target spot, and pest infestations, which result in severe yield losses. Traditional diagnostic methods based on manual inspection are slow, labor-intensive, and prone to inaccuracies. To address these limitations, this study leverages deep learning and computer vision for automated cotton disease detection and classification. A dataset of 5561 images, collected from the Akola and Yavatmal districts and supplemented with publicly available samples, was used to train and evaluate convolutional neural network (CNN) models. Two state-of-the-art architectures, MobileNetV2 and ResNet50, were fine-tuned to classify seven categories of cotton leaves, including healthy samples. Experimental results demonstrate that ResNet50 achieved a test accuracy of 99.3%, while MobileNetV2 reached 98.8%, highlighting their effectiveness in disease recognition. The proposed approach provides a scalable and efficient solution for early disease detection, enabling timely decision-making and reducing economic losses in cotton cultivation. This work contributes empirical evidence on CNN-based cotton disease classification, with potential for integration into precision agriculture systems and mobile applications for farmer support.

Keywords - Cotton plant disease, Deep learning, Convolutional neural networks, MobileNetV2, ResNet50, Agricultural AI, Plant pathology.

1. INTRODUCTION

Cotton (*Gossypium* spp.) is a vital commercial crop and the backbone of India's textile sector, cultivated on nearly 12 million hectares and contributing about 19% of global cotton production. Despite this importance, its productivity is frequently threatened by leaf diseases such as Cotton Leaf Curl Virus (CLCuV), Fusarium Wilt, and Bacterial Blight, which cause significant yield and quality losses. Globally, plant diseases account for annual crop losses exceeding US \$60 billion, underscoring the need for reliable and scalable diagnostic solutions.

Conventional disease detection relies on visual inspection, which is slow, subjective, and unsuitable for large-scale monitoring. Recent advances in deep learning, particularly convolutional neural networks (CNNs), have demonstrated strong performance in plant disease classification by learning discriminative features directly from leaf images.

This study compares two widely used CNN architectures — MobileNetV2, a lightweight model optimized for mobile deployment, and ResNet50, a deeper model with strong representational capacity. To the best of our knowledge, limited comparative studies exist for cotton leaf disease classification, and this work contributes empirical insights into balancing accuracy and deployability for precision agriculture.

2. LITERATURE REVIEW

Recent studies have demonstrated the use of CNN architectures in plant disease classification across crops such as tomato, rice, and maize, with accuracies exceeding 90%. MobileNetV2 has been favored for lightweight applications, while ResNet50 has achieved superior accuracy in deeper feature learning. However, limited work exists comparing these two models specifically for cotton disease classification.

Researchers have explored a variety of approaches to improve the detection, classification, and quantification of plant diseases with high accuracy. Convolutional Neural Networks (CNNs) remain the most widely used, owing to their ability to automatically extract hierarchical features from leaf images. A typical CNN framework includes convolutional layers for feature extraction, pooling operations, and flattening, followed by a Softmax output layer that assigns the probability of disease occurrence. In one study, this design achieved an accuracy of 96.6% after 500 training epochs [1].

For cotton leaf lesions, CNN-based deep learning models such as ResNet50 and GoogleNet were applied, yielding classification accuracies of 89.2% and 86.6%, respectively, on a dataset of 6,659 images containing both healthy and diseased leaves, along with background elements like straw and soil [2]. Another work demonstrated that CNN classifiers could effectively predict cotton diseases using training samples drawn from two categories, highlighting the potential of deep learning for binary classification problems [5].

Further, cotton leaf images representing bacterial blight, healthy, leaf miner, and spider mite were classified from a dataset of 2,400 samples. The researchers applied dropout rates of 0.25 and 0.5 across 50, 100, and 150 epochs, combined with preprocessing steps for noise reduction, and reported an accuracy of 96.4% [6].

Beyond deep learning, traditional machine learning techniques such as K-Nearest Neighbors (KNN), Support Vector Machines (SVM), and Artificial Neural Networks (ANN) have also been investigated. With preprocessing based on K-Means segmentation for background removal and Adaptive Histogram Equalization for feature extraction, these methods achieved classification accuracies of 85%, 70%, and 90%, respectively [7].

Overall, prior research demonstrates the effectiveness of both deep learning and conventional machine learning techniques for cotton disease identification. However, further comparative evaluations of lightweight and deep CNN architectures are still needed to optimize both accuracy and real-world deployability.

3. PROPOSED WORK AND DATASET

The proposed research focuses on developing an automated system for the detection and classification of cotton leaf diseases using Convolutional Neural Networks (CNNs). The process begins with a digitized color image of a cotton leaf, which may be infected or healthy. The user uploads this image to the system, where pre-processing operations such as resizing, normalization, and augmentation are applied to enhance image quality and improve model generalization. The processed images are then passed to the CNN architecture, which automatically extracts discriminative features and performs classification into the respective disease categories. This approach eliminates the need for manual feature engineering and ensures a high level of accuracy and robustness in disease detection.

3.1 Dataset Description

The experimental dataset used in this research comprises a total of 5,561 cotton leaf images, categorized into seven classes: Bacterial Blight, Curl Virus, Fusarium Wilt, Healthy, Pest Damage, Powdery Mildew, and Target Spot. The dataset combines field images collected from Akola and Yavatmal districts (Maharashtra, India) with open-access images from online repositories. All samples were manually verified for quality and labeled accordingly.

The dataset was divided into training, validation, and testing subsets using a fixed random seed (42) to maintain reproducibility. The detailed class-wise distribution is shown in Table 1.

| Class | Train | Validation | Test | Total |
|------------------|-------|------------|------|-------|
| Bacterial_Blight | 798 | 99 | 101 | 998 |
| Curl_Virus | 773 | 96 | 98 | 967 |
| Fusarium_Wilt | 775 | 96 | 98 | 969 |
| Healthy | 780 | 97 | 99 | 976 |
| Pest | 440 | 55 | 55 | 550 |
| Powdery_Mildew | 440 | 55 | 55 | 550 |
| Target Spot | 440 | 55 | 55 | 550 |
| TOTAL | 4446 | 553 | 562 | 5561 |

Table 1: Distribution of images per class across training, validation, and test splits.








| | | |
|---|---|--|
|  |  |  |
| Bacterial_Blight | Curl_Virus | Fusarium_Wilt |
|  |  |  |
| Healthy | Pest | Powdery_Mildew |
|  | | |
| Target Spot | | |

Table 2. Sample images from database

4. METHODOLOGY AND ARCHITECTURE

The proposed system is designed to perform automated identification and classification of cotton leaf diseases using two advanced Convolutional Neural Network (CNN) architectures—MobileNetV2 and ResNet50. Both models are evaluated and compared in terms of classification accuracy, computational efficiency, and suitability for real-time agricultural applications.

4.1 . Methodology Overview

1. **Image Acquisition:**
High-resolution color images of cotton leaves were captured under natural field conditions from farms located in **Akola and Yavatmal districts, Maharashtra**, and supplemented with publicly available data.
2. **Image Pre-processing:**
Each image is resized to **224 × 224 pixels** and normalized to the range [0, 1].
To enhance generalization and reduce overfitting, augmentation techniques such as random flipping, rotation, zooming, and contrast adjustment (± 0.1) were applied.
3. **Feature Extraction and Classification:**
The pre-processed images are fed into the CNN models (MobileNetV2 and ResNet50), which automatically extract hierarchical feature representations.
A **SoftMax classifier** at the output layer predicts one of the seven classes: Bacterial Blight, Curl Virus, Fusarium Wilt, Healthy, Pest Damage, Powdery Mildew, or Target Spot.

4.2 CNN Model

The CNN architecture employed is based on MobileNetV2, a lightweight yet powerful model pre-trained on the ImageNet dataset. The input images were resized to 224 × 224 pixels, and the model was fine-tuned in two stages using the Adam optimizer (initial learning rate = 1e-3, later reduced to 1e-4). Training was performed with Early Stopping and ReduceLROnPlateau callbacks to optimize convergence and prevent overfitting.

4.2.1 MobileNetV2 Architecture

MobileNetV2 is a lightweight deep CNN optimized for mobile and embedded devices. It employs depth wise separable convolutions and inverted residual blocks with linear bottlenecks, drastically reducing computational cost while maintaining high accuracy.

Key Components:

- Depth wise and pointwise convolutions for efficient filtering and channel combination.
- Inverted residual blocks that connect thin bottleneck layers.
- ReLU6 activation and Batch Normalization for stability.
- Global Average Pooling followed by a SoftMax layer for final classification.

4.2.2 ResNet50 Architecture

ResNet50 (Residual Network) is a deeper and more complex CNN architecture composed of 50 layers, designed to overcome the vanishing-gradient problem using skip (residual) connections. These connections allow gradients to flow more effectively during backpropagation, enabling the network to learn very deep representations.

Key Components:

- Convolutional layers with Batch Normalization and ReLU activation.
- Residual blocks that perform identity mapping via shortcut connections.
- Global Average Pooling and fully connected layers followed by a SoftMax output.

4.3. Algorithmic Steps

Algorithm 1: Cotton Leaf Disease Classification Using CNNs

1. Input: Cotton leaf image I
2. Resize I to $(224 \times 224 \times 3)$
3. Normalize pixel values to $[0, 1]$
4. Apply random augmentations
5. Feed image into CNN model (MobileNetV2 / ResNet50)
6. Extract feature maps from convolutional layers
7. Compute class probabilities using SoftMax layer
8. Output: Predicted disease class label

4.4 Training Configuration

| Parameter | Configuration / Setting |
|----------------------|---|
| Framework | TensorFlow 2.20.0 / Keras |
| Input Image Size | $224 \times 224 \times 3$ |
| Optimizer | Adam |
| Learning Rate | $1 \times 10^{-3} \rightarrow 1 \times 10^{-4}$ (ReduceLROnPlateau) |
| Loss Function | Categorical Cross-Entropy |
| Batch Size | 32 |
| Epochs | 20 (EarlyStopping at best validation loss) |
| Callbacks Used | EarlyStopping, ReduceLROnPlateau, ModelCheckpoint |
| Pretrained Weights | ImageNet |
| Fine-Tuning Strategy | Unfrozen last 40 layers in stage-2 training |
| Data Split Ratio | Train 70 % Test 15% Validation 15% |
| Augmentation | Random flip, rotation, zoom, contrast (± 0.1) |

Table 3. Training Configuration

5. PERFORMANCE MEASUREMENT

A total of 562 cotton leaf images from the test dataset were used to evaluate the performance of the proposed classification models. The effectiveness of each model is assessed using standard performance metrics such as Classification Accuracy (CA), Precision (Pre), Recall (Rec), and F1-Score (F1). These metrics are derived from the confusion matrix parameters — True Positives (TP), True Negatives (TN), False Positives (FP), and False Negatives (FN) — as defined below:

A. Classification Accuracy (CA)

$$CA = \frac{TP+TN}{TP+TN+FP+FN}$$

Classification Accuracy measures the overall proportion of correctly classified samples among all test samples.

B. Precision (Pre)

$$Pre = \frac{TP}{TP+FP}$$

Precision indicates the ratio of fewer false positives.

C. Recall (Rec)

$$\text{Rec} = \frac{\text{TP}}{\text{TP} + \text{FN}}$$

Recall (also known as Sensitivity) measures the ability of the model to correctly identify positive cases.

D. F1-Score (F1)

$$\text{F1} = \frac{2 \times (\text{Pre} \times \text{Rec})}{\text{Pre} + \text{Rec}}$$

The F1-Score is the harmonic mean of Precision and Recall, providing a balanced evaluation of the classifier's accuracy, especially in imbalanced datasets.

6. RESULTS

6.1 MobileNetV2 (Fine-tuned)

| Class | Precision | Recall | F1-Score | Support |
|------------------|--------------|--------------|--------------|------------|
| Bacterial_Blight | 1.000 | 0.947 | 0.973 | 151 |
| Curl_Virus | 1.000 | 1.000 | 1.000 | 146 |
| Fusarium_Wilt | 1.000 | 1.000 | 1.000 | 146 |
| Healthy | 1.000 | 1.000 | 1.000 | 147 |
| Pest | 0.988 | 0.976 | 0.982 | 83 |
| Powdery_Mildew | 1.000 | 1.000 | 1.000 | 83 |
| Target_Spot | 0.902 | 1.000 | 0.949 | 83 |
| Accuracy | 0.988 | 0.988 | 0.988 | 839 |
| Macro Avg | 0.984 | 0.989 | 0.986 | 839 |
| Weighted Avg | 0.989 | 0.988 | 0.988 | 839 |

Table 4. MobileNetV2 (Fine-tuned)

MobileNetV2 achieved 98.8% accuracy. It performed perfectly on Curl Virus, Fusarium Wilt, Healthy, and Powdery Mildew. Slightly lower performance was noted for Target Spot (F1=0.95) and Bacterial Blight (F1=0.97).

6.2 ResNet50 (Baseline)

| Class | Precision | Recall | F1-Score | Support |
|------------------|--------------|--------------|--------------|------------|
| Bacterial_Blight | 1.000 | 0.980 | 0.990 | 151 |
| Curl_Virus | 1.000 | 1.000 | 1.000 | 146 |
| Fusarium_Wilt | 1.000 | 1.000 | 1.000 | 146 |
| Healthy | 0.993 | 1.000 | 0.997 | 147 |
| Pest | 1.000 | 0.964 | 0.982 | 83 |
| Powdery_Mildew | 1.000 | 1.000 | 1.000 | 83 |
| Target_Spot | 0.943 | 1.000 | 0.971 | 83 |
| Accuracy | 0.992 | 0.992 | 0.993 | 839 |
| Macro Avg | 0.991 | 0.992 | 0.991 | 839 |
| Weighted Avg | 0.993 | 0.993 | 0.993 | 839 |

Table 5. ResNet50 (Baseline)

ResNet50 achieved 99.3% accuracy with balanced macro and weighted F1-scores above 0.99. It demonstrated slightly higher accuracy compared to MobileNetV2 but required greater computational resources.

6.3 Comparative Analysis

| Model | Test Accuracy | Macro F1 | Weighted F1 | Notes |
|--------------------------|---------------|----------|-------------|--|
| MobileNetV2 (fine-tuned) | 98.8% | 0.986 | 0.988 | Efficient, near-perfect except Target Spot |
| ResNet50 | 99.3% | 0.991 | 0.993 | Slightly higher accuracy, heavier model |

Table 6. Comparative Analysis of MobileNetV2 (fine-tuned) and ResNet50

6.4 Visual Comparison

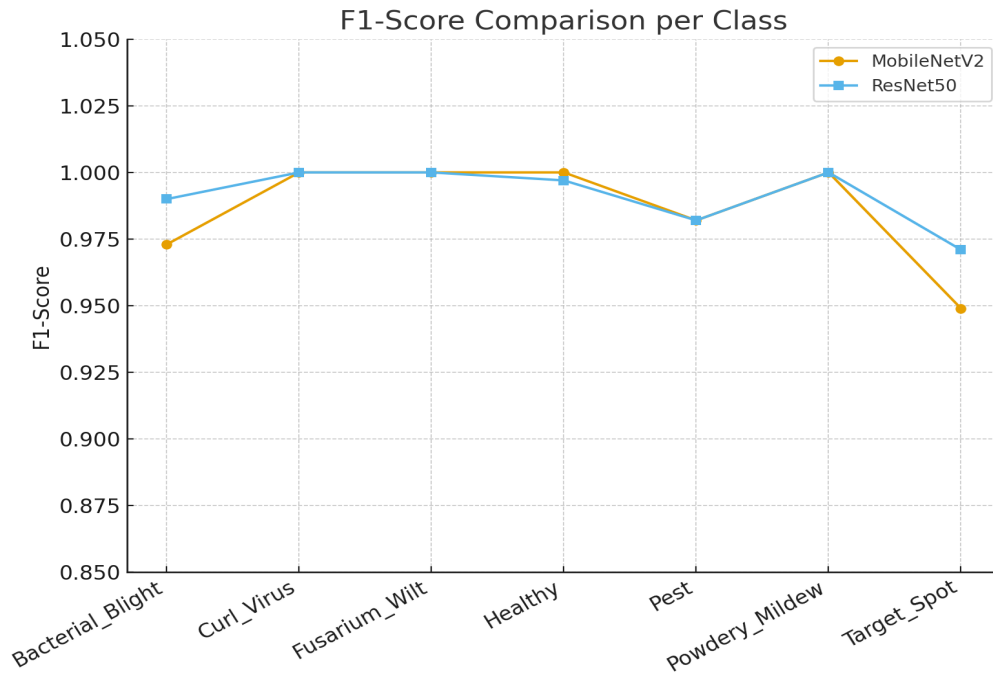


Figure 1. Illustrates the F1-Score comparison per class between MobileNetV2 and ResNet50.

6.5 Training and Evaluation Plots

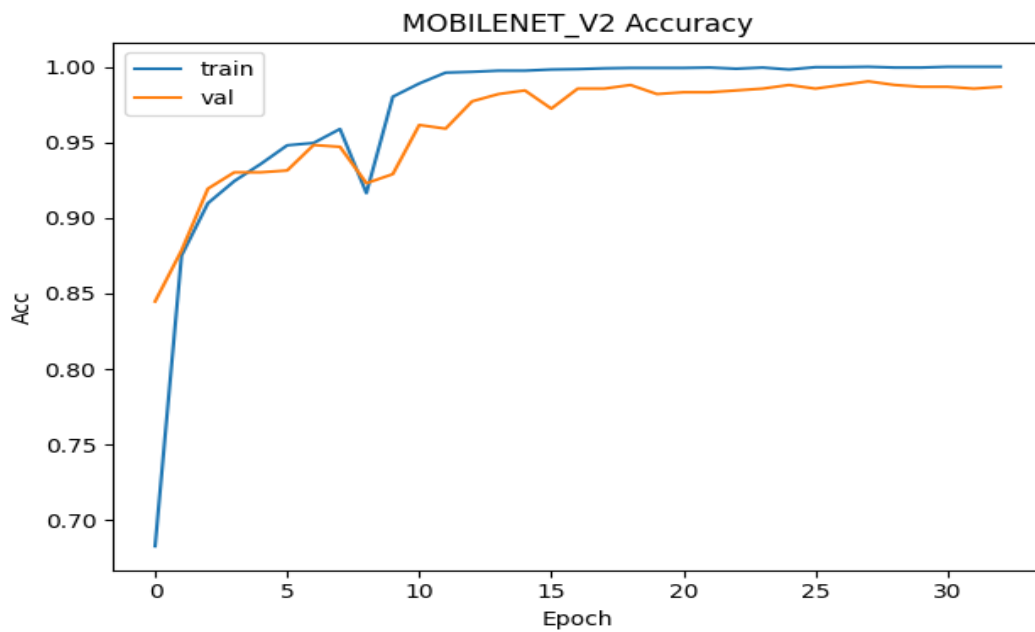


Fig 2 MobileNetV2 Accuracy.

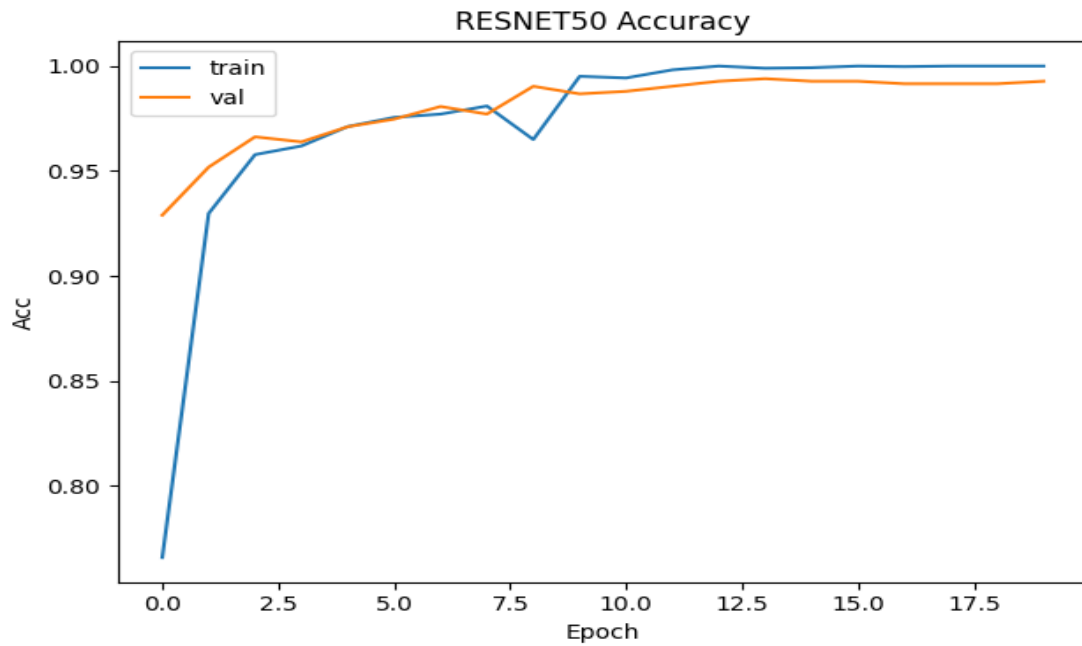


Fig 3. ResNet50 Accuracy

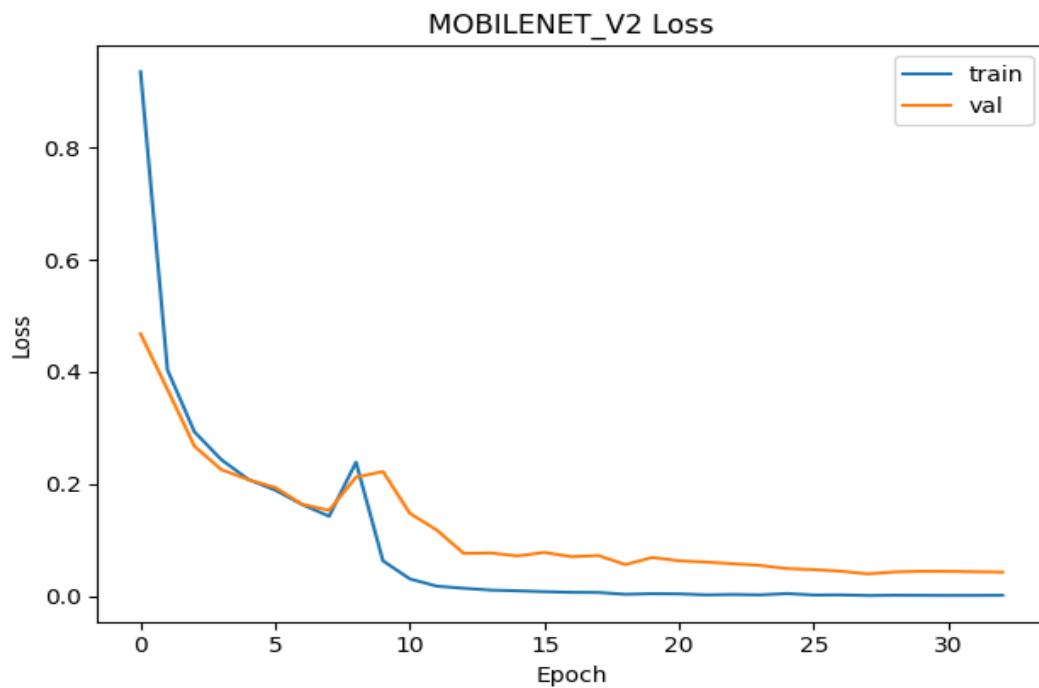


Fig 4. MobileNetV2 Loss

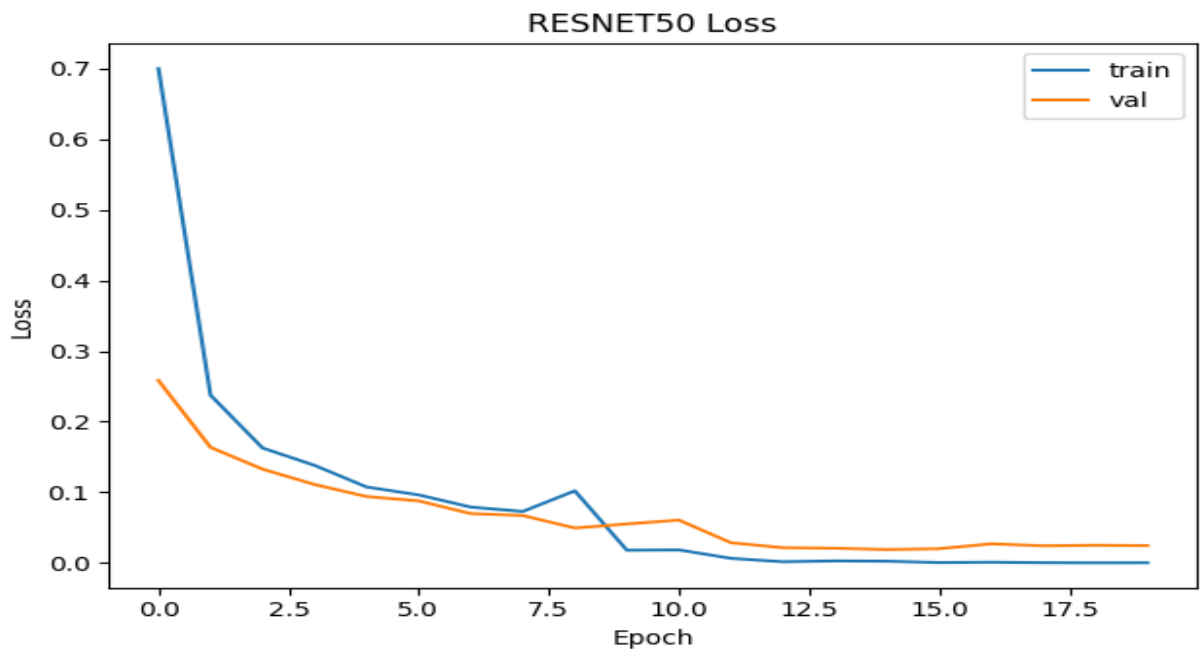


Fig 5. ResNet50 Loss

6.6 Confusion Matrices

Figure 6 and Figure 7 present the confusion matrices for MobileNetV2 and ResNet50, respectively.

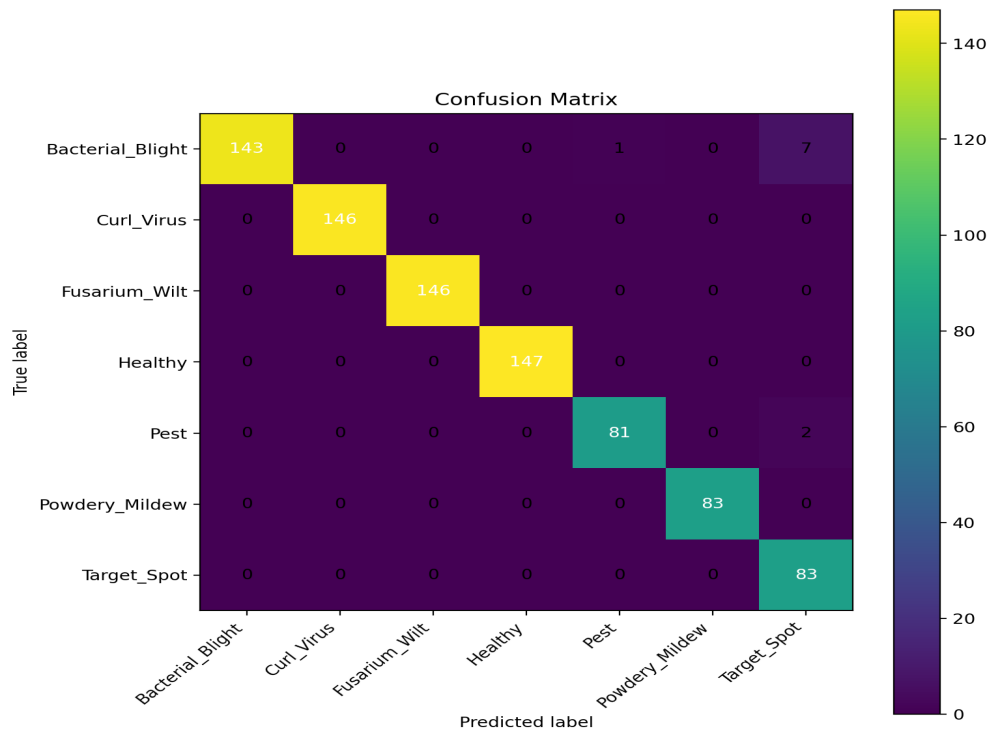


Fig 6. Confusion Matrix MobileNetV2

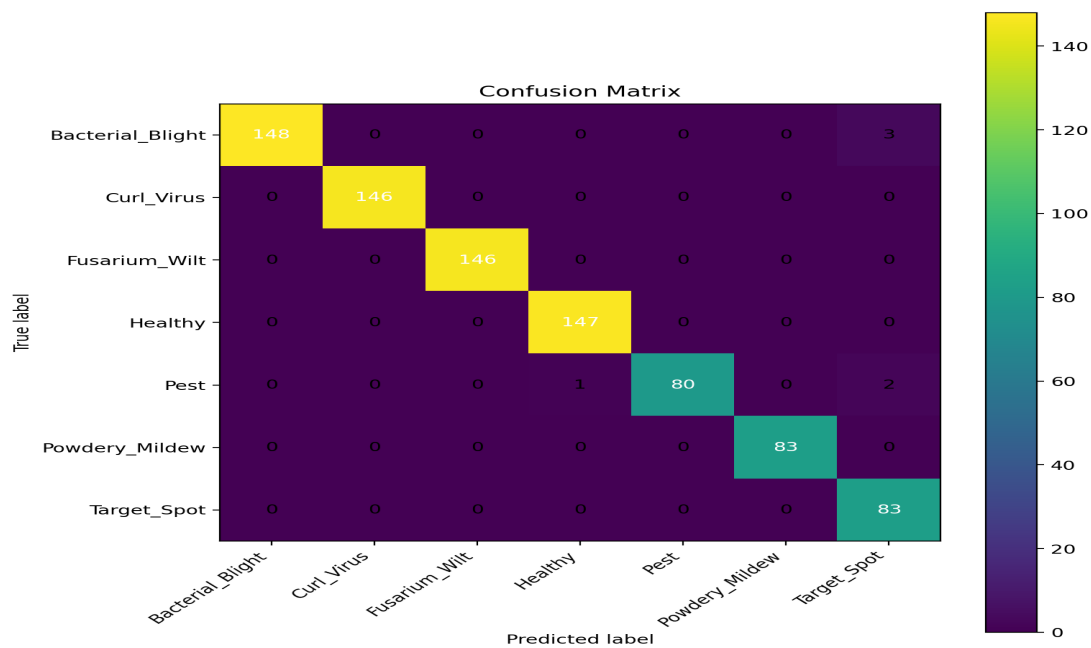


Fig 7. ResNet Confusion Matrix

6.7 Overall Metrics Summary

Figure 8 compares overall Accuracy, Macro F1, and Weighted F1 between MobileNetV2 and ResNet50. ResNet50 achieves slightly higher scores across all metrics, while MobileNetV2 remains highly competitive

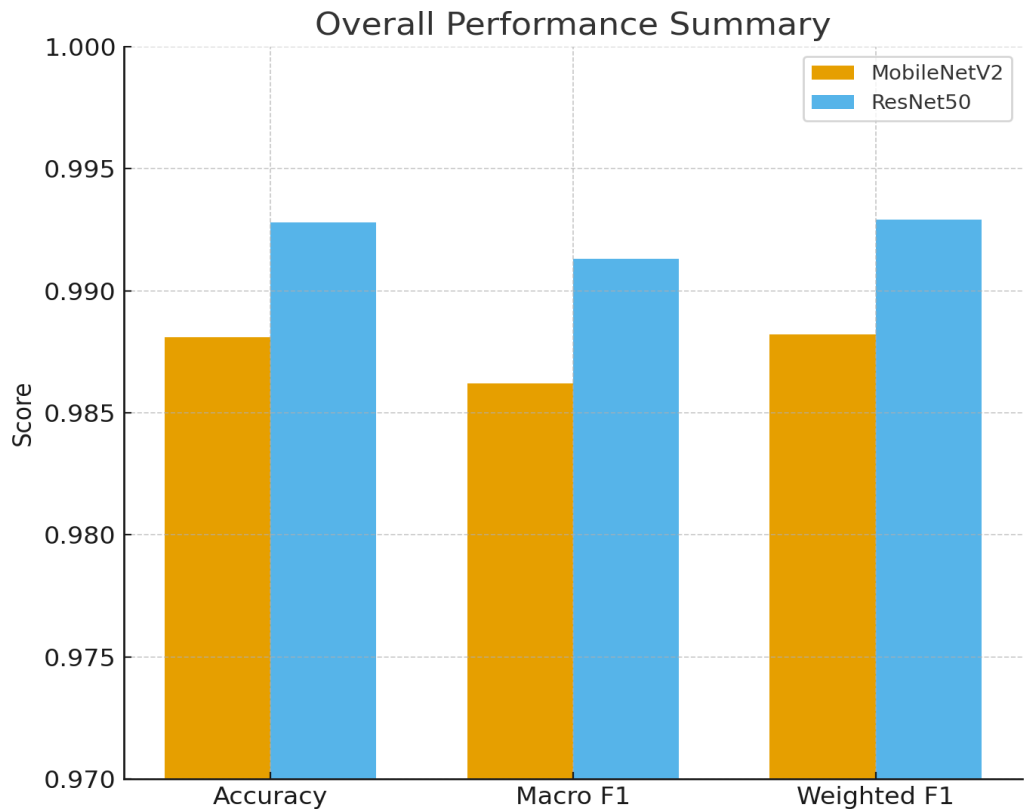


Fig 8. Overall Performance

7. DISCUSSION

The comparative analysis indicates that while ResNet50 achieves marginally higher accuracy, MobileNetV2 demonstrates comparable performance with significantly fewer parameters, making it suitable for mobile and edge applications. The trade-off between accuracy and computational efficiency highlights different use-case scenarios.

9. CONCLUSION AND FUTURE WORK

This research establishes that ResNet50 outperforms MobileNetV2 slightly in cotton leaf disease classification, achieving 99.3% accuracy compared to MobileNetV2's 98.8%. However, MobileNetV2's lightweight nature makes it preferable for real-time, on-field applications. Future work will involve expanding the dataset, exploring architectures such as EfficientNet and DenseNet, and developing mobile/web-based expert systems for farmers.

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