Dataset Summary:

Healthy: 1123 images

Wilt_and_leaf_blight_on_lettuce: 6 images Powdery mildew on lettuce: 18 images

Bacterial: 20 images

Septoria blight on lettuce: 19 images

Viral: 15 images

Downy_mildew_on_lettuce: 30 images Shepherd purse weeds: 1106 images

Preprocessing Step:

- **Data augmentation** is performed on Wilt_and_leaf_blight_on_lettuce, Powdery_mildew_on_lettuce, Bacterial, Septoria_blight_on_lettuce, Viral, Downy mildew on lettuce to increase its number

- Less than 100*100px images are not processed and skipped
- Each image is converted to 224*224 for processing

Methodology

Classical ML Approach (Random Forest)

- 1. **Feature Extraction**: Images were processed to extract numerical features.
 - Color Histograms: Calculated for RGB channels to capture color distribution
 - Haralick Texture Features: Computed from grayscale images to represent texture properties
- 2. Data Preprocessing:
 - All extracted features were stacked into a single feature vector for each image.
 - Features were **standardized** to ensure all features contribute equally to the model.
- 3. **Model Selection**: A **Random Forest Classifier** with 100 estimators was chosen for classification.
- 4. Training and Evaluation:
 - The dataset was split into 80% training and 20% testing sets, with stratification to maintain class proportions.
 - The model was trained on the training data and evaluated on the test set.
 - Confusion Matrix, Classification Report, and Accuracy Score were generated.
 - 5-fold Cross-Validation was performed to assess model generalization.

 Feature Importance was calculated and visualized for the Random Forest model.

Deep Learning Approach (ResNet18)

- 1. **Data Preprocessing (Transforms)**: Images were transformed for input into a deep learning model.
 - o **ToTensor**: Converted images to PyTorch tensors.
 - Normalization: Pixel values were normalized using ImageNet's mean and standard deviation.

2. Dataset and Split:

- The dataset was split into 80% training and 20% validation sets.
- DataLoaders were created for batching and shuffling.
- 3. Model Architecture: A pre-trained ResNet18 model was used.
 - The final fully connected layer of ResNet18 was replaced to match the number of output classes.

4. Training Configuration:

- Loss Function: CrossEntropyLoss was used, suitable for multi-class classification.
- **Optimizer**: **Adam optimizer** with a learning rate of 0.001 was chosen.
- The model was trained for 6 epochs.

5. Training and Validation Loop:

- The model was trained in batches
- Validation was performed after each epoch to monitor performance on unseen data.
- Training and Validation Loss history was recorded.
- Confusion Matrix and Classification Report were generated from validation predictions.

Results

Classical ML Approach (Random Forest)

- Overall Accuracy: The model achieved an accuracy of 0.95 (95%).
- **Cross-validation Accuracy**: 0.9429±0.0123, indicating consistent performance across different data subsets.
- Confusion Matrix:
 - * High true positives are observed along the diagonal. For instance, Healthy (223 out of

225 actual healthy samples were correctly classified), Downy_mildew_on_lettuce (121/121), and Shepherd_purse_weeds (194/195) show excellent performance.

- Some misclassifications occurred:
 - Bacterial was sometimes misclassified as "Powdery_mildew_on_lettuce"
 (2) or "Septoria_blight_on_lettuce"
 (6).
 - Viral was frequently misclassified as "Healthy" (7).
 - "Wilt_and_leaf_blight_on_lettuce" was misclassified as "Bacterial" (1), "Powdery_mildew_on_lettuce" (2), and "Septoria_blight_on_lettuce" (1).

Classification Report:

- Healthy, Downy_mildew_on_lettuce, and Shepherd_purse_weeds classes show very high precision, recall, and f1-score (all close to 0.99 or 1.00), indicating robust performance for these classes.
- Classes like Bacterial, Powdery_mildew_on_lettuce, and
 Septoria_blight_on_lettuce have good precision/recall but slightly lower
 F1-scores compared to the best-performing classes.
- **Viral** has good recall (0.89) but slightly lower precision (0.85).
- "Wilt_and_leaf_blight_on_lettuce" shows perfect precision (1.00) and recall (1.00), but with only 25 samples.

Deep Learning Approach (ResNet18)

• Training Progress:

- Loss over Epochs: Both training and validation loss generally decreased over 6 epochs, indicating learning. Training loss decreased from around 1.2 to below 0.2, while validation loss decreased from around 0.8 to around 0.1.
- Accuracy (Approximate): The validation accuracy, approximated from validation loss, steadily increased, reaching over 0.8 by epoch 2 and fluctuating around 0.8-0.9 in later epochs.

Confusion Matrix:

- * Excellent performance is seen for Healthy (241/242), Downy_mildew_on_lettuce (136/136), and Shepherd_purse_weeds (193/193).
 - Some misclassifications:
 - Powdery_mildew_on_lettuce had 3 samples misclassified as "Downy_mildew_on_lettuce" and 1 as "Healthy."
 - Viral had 4 misclassified as "Downy mildew on lettuce."
 - Wilt_and_leaf_blight_on_lettuce had 1 misclassified as "Shepherd_purse_weeds."

Classification Report:

- Similar to the classical approach, Healthy, Downy_mildew_on_lettuce, and Shepherd_purse_weeds show outstanding precision, recall, and f1-scores (all near 1.00).
- Bacterial, Septoria_blight_on_lettuce, and Viral also show strong performance (F1-scores > 0.90).

- Powdery_mildew_on_lettuce has a precision of 0.95 and recall of 0.92, indicating good but not perfect performance.
- Wilt_and_leaf_blight_on_lettuce shows very good precision and recall (0.96 and 1.00, respectively).

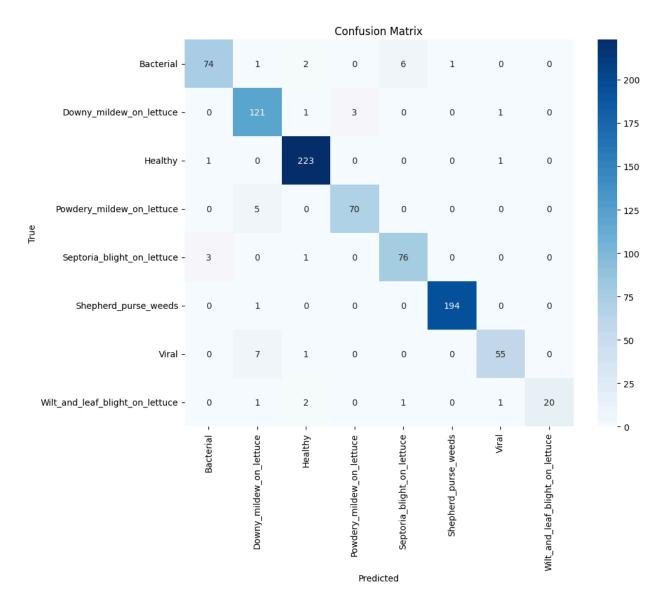
Observations

Both classical machine learning with **Random Forest** and the **Deep Learning approach with ResNet18** achieved high accuracy in classifying plant diseases.

- The Random Forest model achieved an overall accuracy of 95% with a cross-validation accuracy of 94.29±1.23%. It demonstrated strong performance across most classes, particularly for "Healthy," "Downy_mildew_on_lettuce," and "Shepherd_purse_weeds." However, some misclassifications occurred, notably "Viral" being confused with "Healthy" in some instances.
- The Deep Learning model (ResNet18) also showed excellent performance, with training and validation losses decreasing effectively over epochs. The confusion matrix indicates very high accuracy across several classes, with only a few misclassifications. It seems to handle the nuances of visual features effectively due to its pre-trained nature and convolutional capabilities.
- Comparing the confusion matrices both models show similar patterns of strong
 performance for well-represented and distinct classes. The deep learning model appears
 to have slightly fewer scattered misclassifications for some classes (e.g.,
 "Powdery_mildew_on_lettuce" and "Wilt_and_leaf_blight_on_lettuce" have fewer
 misclassifications to diverse classes in the DL approach).
- The deep learning model's training process, as visualized by the loss and approximate accuracy plots, shows a stable learning curve, suggesting good model convergence.
- The use of pre-trained models in the deep learning approach likely contributed significantly to its high performance, as these models have learned rich feature representations from large datasets like ImageNet. In contrast, the classical approach relied on handcrafted features (color histograms and Haralick textures), which, while effective, might not capture the same level of complexity as learned features from a deep network.

In conclusion, both methods are highly effective for this plant disease classification task. The Deep Learning approach with ResNet18 appears to slightly edge out the classical Random Forest model in overall robustness and handling of more complex disease patterns, likely due to its ability to learn hierarchical features directly from raw image data.

Classical ML Approach confusion matrix:



DeepLearning Approach confusion matrix:

