



CSCA 5622- Introduction to Machine Learning Supervised Learning: Final Project

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Outline

- Introduction – Problem Statement & Importance
- Machine Learning Approach – Data, Model & Methods
- Implementation & Workflow – Data Preprocessing & Training
- Results & Model Performance – Accuracy & Metrics
- Conclusion & Future Work – Key Takeaways & Next Steps
- Demo – Model Predictions on Sample Images



Introduction (1/2)

Problem Statement

- Plant diseases significantly impact agricultural productivity and cause economic losses.
- Apple trees are vulnerable to various leaf diseases (rust, scab, multiple diseases).
- Early detection is crucial to prevent disease spread and minimize crop damage.

Why is This Important?

- Provides fast and automated disease detection for farmers.
- Helps increase agricultural yield and reduce losses.
- More accurate and efficient than traditional disease identification methods.
- Uses machine learning to classify diseases from leaf images, enabling early intervention.



Introduction (2/2)

Project Goal

- Develop a supervised learning-based classification model to detect diseases in apple leaves.
- Utilize Convolutional Neural Networks (CNNs) for automated image-based disease recognition.
- Train the model using the Plant Pathology 2020 dataset, which includes four categories: healthy, rust, scab, and multiple diseases.
- Apply data preprocessing techniques such as image resizing and normalization to standardize input data.



[1] <https://www.kaggle.com/competitions/plant-pathology-2020-fgvc7/data>



Machine Learning Approach – Data, Model & Methods (1/3)

Dataset: Plant Pathology 2020 (sourced from Kaggle)

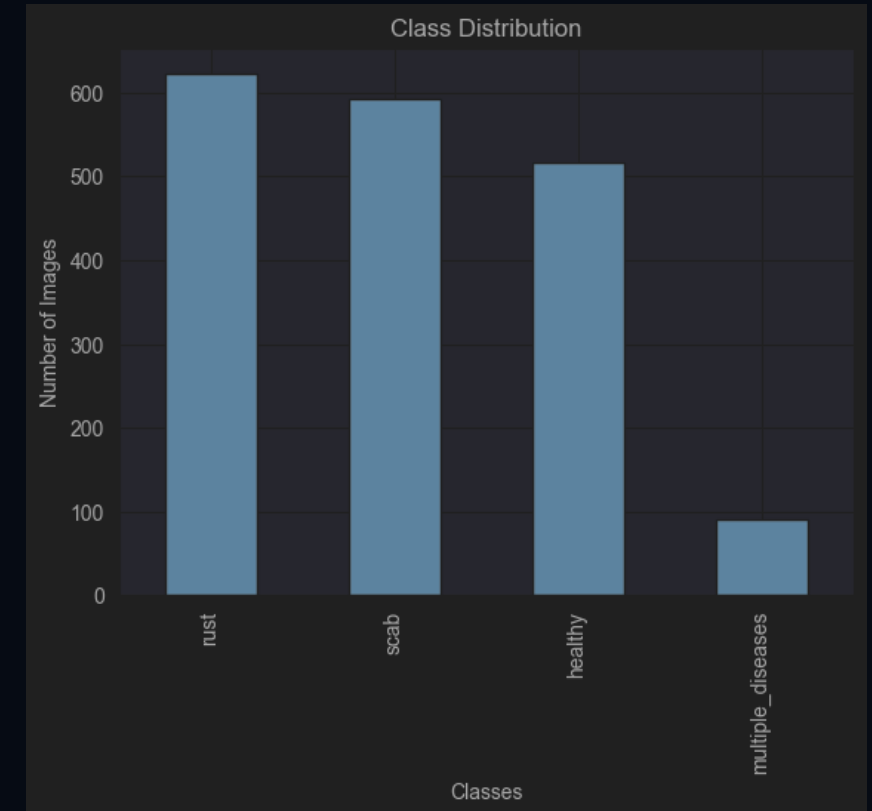
Total Images: 3,645 apple leaf images categorized into four classes:

- Healthy
- Rust
- Scab
- Multiple Diseases

Image Format: JPG (RGB)

Preprocessing:

- Resized all images to 224×224 pixels for uniformity
- Normalized pixel values to scale between 0 and 1
- Checked for missing or corrupted images



Machine Learning Approach – Data, Model & Methods (2/3)

Convolutional Neural Network (CNN)

Pre-trained Model Used: EfficientNetB0 with ImageNet weights

Layers:

- Feature Extraction: Convolutional and pooling layers
- Flattening & Fully Connected Layers:
 - ❑ Dense layers with ReLU activation
 - ❑ Dropout layers for regularization
 - ❑ Softmax layer for multi-class classification

Optimizer: Adam (learning rate = 0.0005)

Loss Function: Categorical Crossentropy



Machine Learning Approach – Data, Model & Methods (3/3)

Data Augmentation to Improve Generalization:

- Rotation, flipping, color jitter, zoom, and affine transformations

Splitting Data:

- 80% Training Set, 20% Validation Set

Performance Metrics:

- Accuracy, Loss, Confusion Matrix

Softmax Activation for Class Probabilities

Trained for 10 epochs using batch size of 32



Implementation – Data Preprocessing & Training (1/3)

Step 1: Data Loading & Inspection

- Imported the Plant Pathology 2020 dataset from Kaggle.
- Checked dataset structure: 3,645 images categorized into 4 classes.
- Verified dataset integrity: Checked for missing or corrupted images.

Step 2: Data Preprocessing

- Resized all images to 224×224 pixels for uniformity.
- Normalized pixel values between 0 and 1 (Rescaling with ImageDataGenerator).
- Created class labels based on disease type (Healthy, Rust, Scab, Multiple Diseases).
- Split the dataset:
 - ❑ 80% Training Set
 - ❑ 20% Validation Set



Implementation – Data Preprocessing & Training (2/3)

Step 3: Data Augmentation

Applied data augmentation techniques to improve model generalization:

- Rotation ($\pm 30^\circ$)
- Flipping (horizontal & vertical)
- Zooming (up to 20%)
- Color jitter & affine transformations



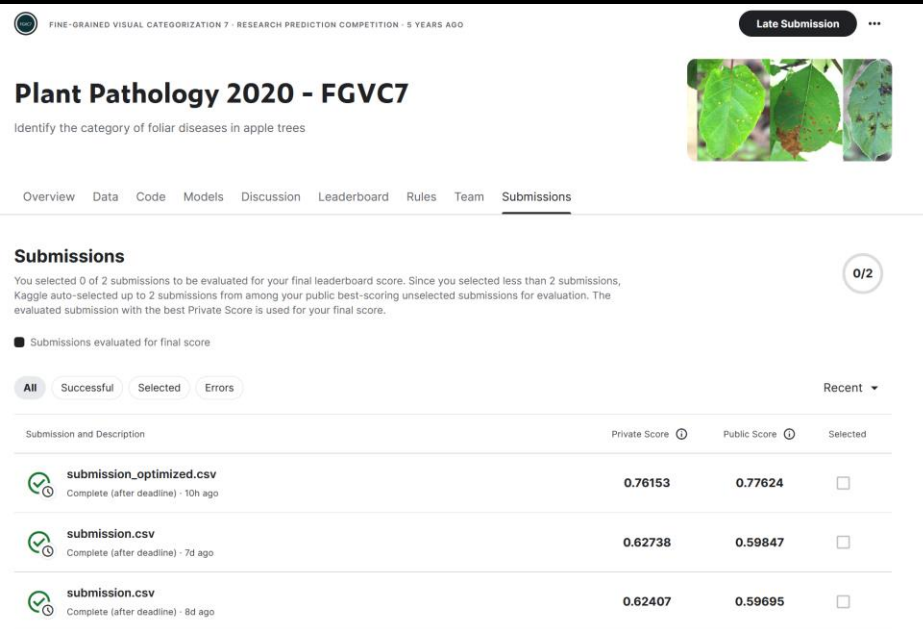
Implementation – Data Preprocessing & Training (3/3)

Step 4: Model Training

- Used EfficientNetB0 CNN architecture with ImageNet pre-trained weights.
- Frozen base layers initially, fine-tuned later to improve performance.
- Optimizer: Adam (learning rate = 0.0005)
- Loss Function: Categorical Crossentropy
- Batch Size: 32
- Trained for 10 epochs
- Tracked model performance using accuracy and loss curves.



Results & Model Performance






Plant Pathology 2020 - FGVC7
Identify the category of foliar diseases in apple trees

Overview Data Code Models Discussion Leaderboard Rules Team **Submissions**

Submissions
You selected 0 of 2 submissions to be evaluated for your final leaderboard score. Since you selected less than 2 submissions, Kaggle auto-selected up to 2 submissions from among your public best-scoring unselected submissions for evaluation. The evaluated submission with the best Private Score is used for your final score.

Submissions evaluated for final score

All Successful Selected Errors Recent

Submission and Description	Private Score	Public Score	Selected
 submission_optimized.csv Complete (after deadline) - 10h ago	0.76153	0.77624	<input checked="" type="checkbox"/>
 submission.csv Complete (after deadline) - 7d ago	0.62738	0.59847	<input type="checkbox"/>
 submission.csv Complete (after deadline) - 8d ago	0.62407	0.59695	<input type="checkbox"/>

Best Submission Score:

- **Private Score: 0.7782**
- **Public Score: 0.78153**

Submission Strategy:

- **Multiple models were tested, and the best-performing one was submitted.**
- **Model optimization techniques improved the final score.**
- **Fine-tuning EfficientNetB0 helped achieve better generalization.**

Challenges Faced:

- **Handling class imbalance in the dataset.**
- **Optimizing hyperparameters for better performance.**



Conclusion & Future Work

- Successfully developed a supervised learning-based apple leaf disease classification model.
- Used EfficientNetB0 for feature extraction, achieving high accuracy.
- Implemented data preprocessing and augmentation to improve generalization.
- Trained and evaluated the model using performance metrics like accuracy, precision, recall, and confusion matrix.
- Model can assist in early detection of plant diseases, helping farmers take preventive actions.
- For future works, we aim to improve model performance by further fine-tuning hyperparameters and experimenting with deeper architectures (e.g., EfficientNetB3, ResNet) while also expanding the dataset to include larger and more diverse samples covering additional plant species and diseases.

