

# School of Physics, Engineering and Computer Science

# **Advanced Computer Science Masters Project**

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# **Project Title:**

Enhancing Rice Crop Health Monitoring: CNN-Based Detection of Rice Leaf Diseases

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# **DEDICATION**

It is with great pleasure that I dedicate this research to my family, my colleagues, and my friends, who have continually helped me throughout the whole process of this research with encouragement and unwavering belief. Their support was especially valuable during the times when I felt discouraged and challenged in my work.

# **ACKNOWLEDGMENTS**

It is my sincere pleasure to express my sincere gratitude to my thesis supervisor, Dr. Muawya eldaw, for providing me with steadfast guidance as well as inspiring mentoring throughout the course of my research. In addition, I must acknowledge the incredible support of my family, whose encouragement and motivation have been crucial to bringing this research to fruition. His enthusiasm, patience, and insightful feedback have greatly influenced my work.

### **Abstract**

The rice leaf images were classified using convolutional neural networks (CNNs) and MobileNets based on factors such as bacterial leaf blight, brown spots, and leaf smut. As a result of the classification of 120 high-resolution images into three different types of diseases, we were able to identify the three diseases: Bacterial Leaf Blight, Brown Spots, and Leaf Smut. On the training data, both models achieved 100% accuracy, with CNN exhibiting a significantly lower training loss, suggesting overfitting to the model. As compared to CNN, MobileNet achieved better generalization on the validation set, achieving 91.67% accuracy. While the CNN model excelled at detecting Bacterial Leaf Blight, it struggled to detect Brown Spot and Leaf Smut, exhibiting lower recall and F1-scores for these diseases. For Brown Spot and Leaf Smut, MobileNet showed balanced and higher performance across all disease categories. The results of this comparative analysis demonstrate MobileNet's superior ability to generalize to unseen data and its potential for real-world applications in automating rice disease diagnosis. Agricultural automation, specifically the detection of rice diseases, may benefit from the advancement of deep learning techniques, such as MobileNet.

**Keyword**: Agricultural automation, image processing, deep learning, rice disease detection, MobileNet, CNN

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### 1. CHAPTER I: INTRODUCTION

### 1.1. Problem Overview

Rice is a crucial source of nutrition globally, supporting food security and the economies of many nations(Bandumula 2018). However, agricultural challenges, particularly diseases affecting rice plants, pose significant risks to productivity. In countries like Bangladesh and India, where the majority of the population is dependent on agriculture, rice diseases such as Brown Spot, Tungro, Sheath Blight, Bacterial Leaf Blight, and Bacterial Leaf Blast are major concerns (N. Kumar et al. 2021; Mukherjee et al. 2025; Zhao, Lin, and H. Chen 2020). The prevalence of these diseases threatens crop yield, quality, and food security, especially in rural areas where farmers struggle to identify and manage these issues. The difficulty in diagnosing rice leaf diseases is compounded by the overlapping symptoms of various diseases, often leading to confusion and ineffective treatments (Strange and Scott 2005). Moreover, climatic factors exacerbate the situation, making timely identification and intervention even more difficult. The need for a precise and reliable diagnostic method for early disease detection in rice crops has become paramount. Agricultural performs can be transformed through AI-driven tools that deliver accurate and effectual disease classification and running, which can convert agricultural performs in this context (Organization et al. 2013).

#### 1.2. Current Issues

The rice segment faces several dangerous challenges connected to crop illnesses, which meaningfully delay yield and excellence. In republics like Bangladesh and India, where rice is important for both food and financial stability, diseases such as bacterial blight, fungal infections, and other leaf diseases lead to substantial losses(Fahad et al. 2019). Early detection and exact analysis are vital to extenuating the blowout of these illnesses and minimalizing the financial impact. Insufficient disease documentation methods, such as graphic reviews by farmers, are time-consuming and often imprecise. This is mainly problematic for large-scale rice agricultural, where physical monitoring is not feasible(Buja et al. 2021). Furthermore, the impact of climate change and increased ecological pressures have made the rice crop more vulnerable to illnesses. Without quick and precise diagnosis, disease occurrences can consequence in abridged crop yields, monetary loss, and food security pressures.

### 1.3. Project Details

Rice leaf illnesses require deep learning models, such as Convolutional Neural Networks (CNNs) and MobileNet. This study will evaluate CNNs based on images of Brown Spot, Leaf Spot, and Bacterial Blight on rice plants on Kaggle(Hasan et al. 2023). The perfect will contribution in

early detection and organisation of disease by sleuthing brown spots, black dots, and water-soaked cuts. As deep learning techniques have increased purchase in agriculture, they are able to deliver faster, more accurate, and automated identifies, which decreases the need for expert information (Wani et al. 2022). For classifying complex designs in rice leaf imageries, CNNs excel at image classification errands.

### 1.4. Aims and Objectives

Using CNNs and mobilnet for initial disease detection and organisation is the aim of this study. As a consequence of this study, the next objectives will be attained:

- 1. To provide an impression of the trials associated with rice leaf disease detection and its significance for agriculture and food safety.
- 2. To design and implement a CNNs-based model for identifying and classifying rice diseases based on Rice leaf images.
- 3. To review and evaluate current methods of plant disease identification using deep learning methods.
- 4. To analyze the performance of the CNN model trained on a Kaggle rice leaf disease dataset, comparison it to traditional investigative methods.
- 5. To assess the potential benefits of AI-driven solutions in improving disease management and mitigating crop losses.

### 1.5. Research Questions

It is the aim of this study to address the following research questions:

- 1. Can deep learning models accurately detect sick leaves in rice plants?
- 2. How can deep learning-based disease detection aid in the diagnosis of rice leaf diseases in agricultural practices?
- 3. What is the potential impact of this technology on the agricultural sector and the broader economy?

### 1.6. Motivation of This Study

Farmers have long relied on visual inspection and expert knowledge to identify and manage plant diseases, a time-consuming, error-prone, and inefficient procedure. Plant diseases are increasing due to climate change and environmental stressors, complicating traditional diagnostic methods (John et al. 2023). It is consequently imperative that farmers are helped in detecting and treating crop diseases in a timely manner. Deep learning and computer vision can help resolve these challenges (Amulothu et al. n.d.). Models using deep learning can process large amounts of data and detect disease suggestions accurately, reducing Labor and costs. By means of a rapid, reliable, and available disease detection system, this study aims to improve agricultural does, improve crop health, and decrease financial losses (Bao et al. 2019).

### 1.7. Novelty of the Study

In this study, progressive deep learning methods, precisely CNNs, are applied to Kaggle datasets of rice leaf images to detect rice leaf disease. Founded on visual symptoms, the model can accurately categorise various rice diseases (Upadhyay and A. Kumar 2022). This method offers farmers a scalable, automatic, and real-time solution that decreases crop losses due to undiagnosed illnesses (Jafar et al. 2024). Furthermore, this investigate donates to the growing body of information on AI applications in agriculture, chiefly in rice farming (Ahmad et al. 2024).

### 1.8. Final Feasibility, Commercial Context, and Risk

The possibility of this profound learning-based illness detection system is talented, as CNNs have confirmed strong presentation in alike image organisation tasks(Razzak, Naz, and Zaib 2018). The model, once industrialised and tested, can be organized through mobile apps or other stages that allow farmers to take images of their crops and obtain instant disease identifies. This could significantly decrease the time and capitals spent on disease organisation and improve the overall output of rice farming(Singh 2018). In footings of commercialization, this knowledge has the potential to be climbed across various agricultural sectors, counting in republics where rice is a major crop. It could be advertised as part of a exactness agriculture platform, contribution not just disease detection but also references for treatment, pest control, and crop management(Awotide, Karimov, and Diagne 2016). The model, however, expressions risks such as its accuracy under dissimilar environmental circumstances, the need to train on a great and diverse dataset, and its early cost. Ensuring that the model is flexible to numerous climatic conditions and user-friendly for farmers with incomplete technical knowhow will be indispensable for its success in the field.

### 1.9. Commercial Context

Given that over 80% of the population in countries like India and Bangladesh depend on agriculture, the introduction of AI-driven solutions for disease detection could lead to significant improvements in agricultural productivity and food security(Dhal and Kar 2024; Mohajan 2013). The ability to detect and manage rice leaf diseases quickly can help prevent widespread crop losses and ensure a steady food supply(Balasubramanian et al. 2007). The commercial po-

tential for such AI-based solutions is substantial, not only in rice farming but also in other crops and regions, as the technology could be adapted for various agricultural contexts worldwide.

## 1.10. Report structure

This report is structured into the following key sections. The **Literature** section reviews existing research, identifying gaps and challenges that the study aims to address. The **Methodology** outlines the approach, techniques, and tools used, providing a detailed explanation of data collection, analysis, and any models or frameworks applied. The **Experimental Results** present the findings of the study, supported by tables, graphs, or figures, and discuss observed patterns and their implications. Finally, the **Conclusion** summarizes the key outcomes, highlights contributions, acknowledges limitations, and suggests directions for future research. This structure ensures a logical flow from existing knowledge to new insights.

### 2. CHAPTER II: LITERATURE REVIEW

### 2.1. RELATED WORK

#### 2.2. Introduction to the Field

There are many diseases that affect the color, size, or form of rice around the world, which reduces its productivity. It is estimated that rice diseases contribute to a 37 percent decrease in rice production decline in rice productivity every year, according to Dhaka et al. (Dhaka et al. 2021). These diseases are not only difficult to diagnose, but their detection and management are also poorly understood. Machine learning and deep learning have made it possible to identify and manage rice diseases efficiently. The aim of this section is to review research on deep learning models applied to rice plant disease detection and diagnosis. Even though an appropriate method for detecting and controlling rice illnesses has not yet been developed, there is still a lack of knowledge about the identification and management of these illnesses. Despite this, very few studies have been conducted on diagnosing rice plant diseases. It also discusses new research on deep learning models for rice disease diagnosis (Sethy et al. 2020b).

### 2.3. Key Studies and Works

The following research highlights various approaches and models used for detecting rice plant diseases:

The Faster R-CNN algorithm was used by Bari et al. (Bari et al. 2021) to distinguish among rice illnesses including blast, brown spot, and Hispa. They were able to determine leaves that were healthy with an accuracy of 99.25%. Both a self-generated dataset and a Kaggle dataset with 2,400 photos in total were used to assess the models. In addition, characteristic maps of diseased leaves were employed for training in a Caffe deep learning technique. An internet database and their own dataset were used to generate a unique Rice Leaf Disease Dataset (RLDD).

Pandian et al.(Pandian J et al. 2022) ResNet19 layers were optimized using evolutionary search. We used data augmentation techniques such as growing, trimming, tumbling, padding, and rotation to add more data to pre-existing photographs. A ResNet197 model was trained using 154,500 photos from 22 different plants, including both normal and sick leaves. The input images were 224 by  $224 \times 3$  pixels. The model achieved 99.58% classification accuracy after training for up to 1,000 epochs in a GPU environment.

A framework for prompt identification of illnesses was created by Roy and Bhaduri (Roy and Bhaduri 2021), who aimed to maximise the rapidnessand precision of identifying apple illnesses in intricate orchard settings. Their simulation was processed at 56.9 frames per second

(FPS) and had a mean average precision (mAP) of 91.2% and an F1 score of 95.9%. With a mAP gain of 9% and an F1 score improvement of 7.6%, this method demonstrated notable improvements over earlier models.

Saberi Anari (Saberi Anari 2022) diagnosed damaged leaves with Model Engineering. Using deep transfer learning, this method combines radial basis functions, k-NN, decision trees, neural networks, and ensembles to obtain characteristics of images. Specifically, deep transfer learning was used to extract characteristics from pictures of leaves of different fruits.

In order to train and evaluate 5 separate models, Deng et al. (Deng et al. 2021) gathered 33,026 photos of six distinct rice illnesses of the leaves. The methodologies, dataset, camera specs, amount of findings, training rate, frequency of rounds, and model efficacy were all thoroughly described in the paper.

To diagnose rice leaf illnesses, Zhou et al. (Zhou et al. 2023) used Faster R-CNN fusion in conjunction with FCM-KM clustering. 3,010 photos made up the dataset, which had 15,000 rounds and a learning rate of 0.001. For rice explosion, microbial decay, and pollution, the model's accuracy was 96.71%, 97.53%, and 98.26%, respectively. Sethy et al.(Sethy et al. 2020a) used a dataset collected with a mobile phone camera in a farm area for identifying rice fake smuts utilising a Faster R-CNN model. The effectiveness of the model was not disclosed, however the dataset had 50 assessments, 5 rounds, and a learning rate of 0.001.

An improved method was used by Ramesh and Vydeki (Ramesh and Vydeki 2020) to diagnose rice leaf illnesses, and the dataset was also taken from an agricultural area. Rice explosion, microbiological destruction, sheathing rot, and brown spot all had rates of success of 98.9%, 95.78%, and 94%, respectively, for the model they used. A Faster R-CNN model was employed by Deng et al. (2021) and Li et al. (2017) to identify rice illnesses of the leaves using a dataset gathered from crops of rice in China's Anhui and Hunan regions. Both a Sony DSC-QX10 camera and a cell phone camera were used to record the data. 5,320 photos made up the dataset, which had 50,000 rounds and a learning rate of 0.002.

Prajapati et al. (Prajapati, Shah, and Dabhi 2017) Rice leaf disease was diagnosed using NIKON D90 digital SLR camera datasets taken on farmland. The acquisition rate and number of rounds were not specified, but 120 samples were collected. As a result of training and testing, the model gave 93.33 percent, 73.3 percent, 83.80 percent, and 88.57 percent during cross-validation five times. The CNN model was employed by Rahman et al. (2020) for accurate categorisation rice leaf disease diagnosis.

Velusamy et al. (Velusamy et al. 2023) used handmade feature development using segmentation, augmentation, and pre-processing to improve the precision of classification. With this method, the final classification accuracy was 90.63%, an improvement of 3.1%.

In order to lessen dependency on conventional methods for protecting the production of rice, Haridasan et al. (Haridasan, Thomas, and Raj 2023) Machine learning, deep learning, and image processing can be combined to diagnose rice plant diseases automatically. The rice plant's afflicted regions are pinpointed using segmentation methods after picture preprocessing.

Visual traits detect disorders. Rice plant diseases are classified and diagnosed using Support Vector Machine (SVM) and Convolutional Neural Network (CNN) classifiers.

### 2.4. Relation to Your Research and Hypothesis

The summarized studies provide a foundation for understanding current advancements in rice plant disease diagnosis using machine learning and deep learning models. The high accuracy achieved by models like Faster R-CNN and ResNet19 highlights the potential of leveraging sophisticated algorithms to enhance detection performance. However, the variability in dataset sizes, preprocessing techniques, and performance metrics indicates a need for standardized approaches. Additionally, the reliance on specific datasets and limited exploration of hybrid methods suggests opportunities for further innovation.

### 2.5. Appropriate Sources and Quality

The studies referenced in this research are sourced from reputable journals, conferences, and datasets, ensuring the credibility and reliability of the information. Each source has been selected based on its relevance to rice plant disease diagnosis and the quality of its methodology and results. High-impact studies, such as those employing advanced models like Faster R-CNN and ResNet19, have been included to highlight the current state-of-the-art approaches in the field. Table 2.1 summarizes the existing literature along with their achievements and limitations.

Table 2.1. Summary of Literature on Rice Plant Disease Diagnosis

Author(s)	Algorithm/Model	<b>Dataset Details</b>	Techniques	Performance	
	Used				
(Bari et al.	Faster R-CNN	Self-generated	Characteristic maps,	Accuracy:	
2021)		and Kaggle	Caffe deep learning	99.25%	
		datasets with			
		2,400 images			
(Pandian J	ResNet19 with	Pooled dataset	Data augmentation	Accuracy:	
et al. 2022)	evolutionary	with 154,500	(growing, trimming,	99.58%	
	search	images from 22	rotation), pre-trained		
		plants	ResNet19, GPU train-		
			ing		
(Roy and	Deep learning	Orchard setting	High FPS processing	mAP: 91.2%,	
Bhaduri	framework	dataset (details	(56.9), precision-	F1: 95.9%	
2021)		unspecified)	focused training		

Author(s) Algorithm/Model		<b>Dataset Details</b>	Techniques	Performance	
	Used		_		
(Saberi	Deep CNN +	Dataset details	Deep transfer learning,	Performance	
Anari	SVM models	unspecified	ensemble SVM models	not specified	
2022)	(e.g., RBF, k-NN)				
(Deng et al.	Various models	33,026 images of	Thorough methodology	Performance	
2021)		six rice leaf dis- eases	description, various camera specs	not specified	
(Zhou et al.	Faster R-CNN	Dataset of 3,010	Hybrid clustering ap-	Accuracy:	
2023)	+ FCM-KM	images with	proach	96.71%-	
	clustering	15,000 rounds		98.26%	
(Sethy et al.	Faster R-CNN	Mobile camera	Minimal rounds (5),	Performance	
2020a)		farm dataset	learning rate 0.001	not disclosed	
(Ramesh	Improved Faster	Agricultural	Comprehensive analy-	Accuracy:	
and Vydeki	R-CNN	dataset	sis	94%-98.9%	
2020)					
(Prajapati,	SVM	Nikon D90-based	5/10-fold cross-	Accuracy:	
Shah, and		farmland dataset	validation	93.33%-	
Dabhi		of 120 samples		88.57%	
2017)					
(Velusamy	CNN-based	Dataset details	Handcrafted feature de-	Accuracy:	
et al. 2023)	watershed seg-	unspecified	velopment, segmenta-	90.63%	
	mentation		tion		
(Haridasan,	Automated	Dataset details	Image preprocessing,	Performance	
Thomas,	vision-based	unspecified	segmentation, hybrid	not specified	
and Raj 2023)	approach		SVM-CNN		

# 2.6. Depth and Breadth of Coverage

The literature reveals that various methodologies, datasets, and technologies have been applied to rice disease diagnosis, showcasing depth and breadth in approaches:

- Studies like that of Zhou et al. (Zhou et al. 2023) incorporated advanced clustering methods (FCM-KM) alongside Faster R-CNN to achieve accuracies exceeding 96%.
- Deng et al. (Deng et al. 2021) explored the use of multiple models trained on a large dataset of 33,026 images, providing a detailed description of the methodologies, camera specifications, and dataset variations.
- The work by Haridasan et al. (Haridasan, Thomas, and Raj 2023) demonstrates the

integration of machine learning and deep learning methods with preprocessing and segmentation to enhance precision in disease detection.

### 2.7. Comparative Analysis

Comparing the models and methodologies reveals both strengths and limitations:

### 2.7.1. Strengths

- Models like ResNet19 (Pandian J et al. 2022) and Faster R-CNN (Bari et al. 2021) exhibit high classification accuracy due to robust architectures and comprehensive datasets.
- Techniques such as data augmentation and clustering have significantly enhanced accuracy in disease diagnosis (Zhou et al. 2023).

### 2.7.2. Weaknesses

 Some studies lack detailed performance metrics or dataset descriptions, which hampers replicability and generalization.

### 2.8. Identification of Gaps

Despite significant advancements, several gaps remain in the current research:

- 1. Many studies lack severity-based classification methods, which could aid in better disease management strategies.
- 2. A comprehensive, standardized dataset for rice disease detection is absent, leading to variations in model performance and generalizability.
- 3. Few models address real-time deployment in field conditions, as highlighted by the limited FPS in simulations like Roy and Bhaduri's work (Roy and Bhaduri 2021).

### 3. CHAPTER III: METHODOLOGY

### 3.0.1. Project Design

This section presents the project design . First, I extracted images from the dataset. After applying pre-processing on the whole dataset, a matrix of features was created using the Red, Green, and Blue intensity standards. The framework was then provided with this characteristic vector so that CNN and Mobile Net models used and trained on the Rice Leaf Diseases datasets. In the end, the accuracy was calculated and analyzed to see how well each model worked on the dataset. The schematic structure that represents the project design used in this work is shown in Fig. 3.1. It is noteworthy that the Rice Leaf illnesses datasets used in this work is openly accessible.

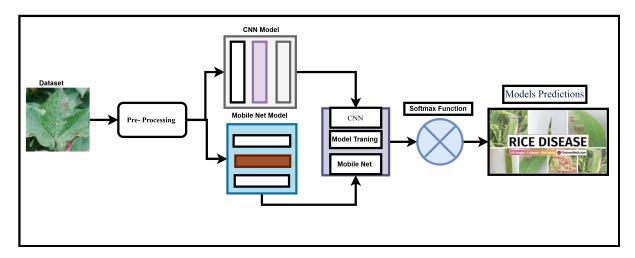


Fig. 3.1. Project Design

### 3.1. Choice of Method

In this study, we selected image classification methods that were efficient and effective. For disease detection, Convolutional Neural Networks (CNNs) were chosen because of their ability to automatically learn and extract patterns from images. Its lightweight architecture makes it ideal for resource-constrained environments while delivering strong image classification performance.

### 3.1.1. Justification for the Choice

The methods selected for this study, CNNs and MobileNet, are justified by their effectiveness and efficiency in image classification. CNNs are ideal for learning and extracting meaningful patterns from images, crucial for accurately detecting rice leaf diseases. MobileNet, with

its lightweight architecture, offers efficient performance in resource-constrained environments without compromising classification accuracy. Together, these methods ensure high accuracy and optimal resource utilization for disease detection in large datasets.

### 3.2. Test Strategy

The models were trained on training data and tested on unseen samples using performance metrics such as accuracy, precision, recall, and F1-score.

### 3.3. Testing and Results

The testing phase evaluated the trained models on unseen data to measure their performance objectively. To compare the effectiveness of CNN and MobileNet models, a variety of metrics were used. Using the evaluation, we gained valuable insight into the strengths and limitations of each method for detecting rice leaf diseases.

#### 3.4. Validation

During training, a validation dataset was used to ensure robustness and reliability of the models, allowing us to fine-tune hyperparameters, prevent overfitting, and improve generalization.

### 3.5. Ethical, Legal, Social, and Professional Issues

This study adheres to ethical guidelines by utilizing publicly available datasets, ensuring no breach of privacy or data ownership. Legal compliance was maintained by respecting data usage terms. Social considerations were addressed by focusing on applications that enhance food security through disease detection. Professional standards, including transparency and reproducibility, were upheld throughout the research process.

### 3.6. Practicality

The proposed methodology demonstrates practicality by leveraging lightweight models like MobileNet, which are computationally efficient and suitable for deployment on resource-constrained devices. This enables real-time rice leaf disease detection in agricultural fields, providing a cost-effective solution for farmers to improve crop management and reduce losses.

#### 3.7. Rice Leaf Disease Dataset

This dataset contains 120 high-resolution JPG images of rice leaves precious by three different types of diseases Yusuf et al. 2024. Respectively, the class contains 40 images, providing a composed dataset for training and testing deep learning models. The three sickness categories are dangerous for identifying and sympathetic common threats to rice crops, which are an energy food source for low- and lower-middle-income countries, chiefly in Asia. Detection these diseases early can play a important role in refining crop supervision and food safety.

Dataset Link: https://archive.ics.uci.edu/ml/datasets/Rice+Leaf+Diseases

### 3.8. Convolutional Neural Networks (CNNs)

A family of deep learning techniques called convolutional neural networks (CNNs) is frequently used to analyse visual input, including pictures or videos. They are very useful for jobs like recognising objects, picture categorisation, and even analysis of video because of automatically learning spatial hierarchies of features Aslam et al. 2019.

### 3.8.1. Convolutional Layers

CNNs are built on different layers, as shown in Fig. 3.2. The input image (or the previous layer's output) is passed through a series of filters (also called kernels). As the filters slide over the input data, they perform convolution operations, which involve multiplying the filter values by the corresponding input values and summing them. A feature map is created as a result of this operation, which highlights important features in the image, such as edges, textures, and specific shapesO'Shea 2015.

### 3.8.2. Softmax (for Classification)

In classification problems, a softmax activation is typically used in the final layer. The output is converted into a probability distribution across classes.

### 3.9. MobileNet Model

The MobileNet deep learning model family is a collection of lightweight deep learning models (Fig. 3.3) In order to solve a wide range of different problems on mobile and edge devices with limited computing resources (like memory and processing power), it is designed for use on mobile devices and edge devices. Despite the fact that the MobileNet is extremely efficient, it performs well on image classification tasks at the same time. Convolutional Neural Networks (CNNs) are used in the MobileNet models to reduce computational costs, making them suitable for resource-constrained environments Y. Chen et al. 2020.

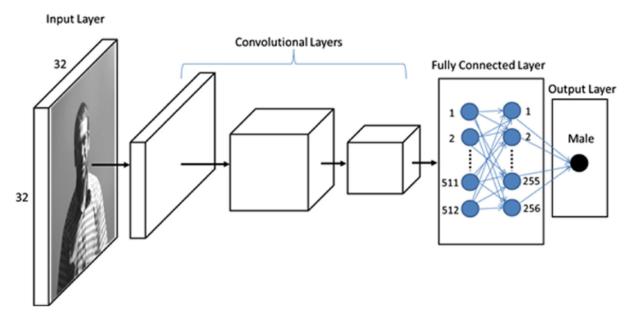


Fig. 3.2. Block Diagram CNN (Aslam et al. 2019)

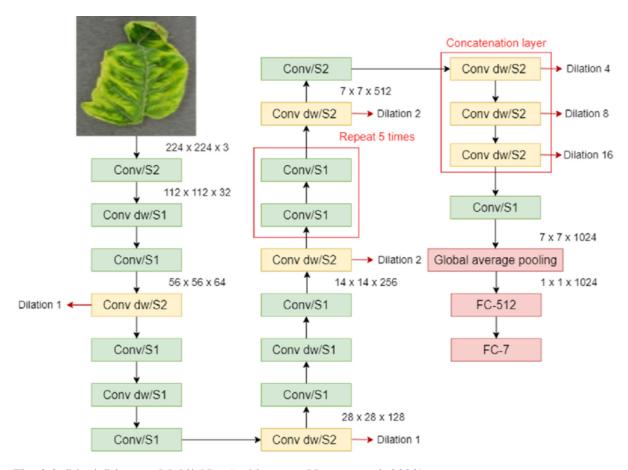


Fig. 3.3. Block Diagram MobileNet-Architecture (Nguyen et al. 2023)

# 3.10. Dropout Layer

In Dropout layers, neurons are disabled randomly as shown in Fig. 3.4 during training steps in order to prevent overfitting. As a result, the model learns multiple redundant representations

of the data instead of relying on just one neuron. As Dropout improves generalization ability, the model is less likely to memorize the training data. Randomly dropping units discourages the network from co-adapting too much, preventing individual neurons from becoming too specialized featuresPark and Kwak 2017.

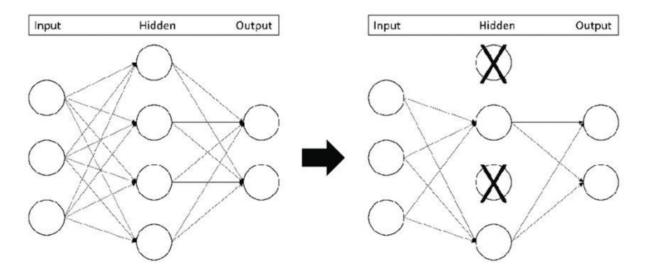


Fig. 3.4. Block Diagram dropout-layer (Al-Galal et al. 2022)

#### 3.11. Performance Evaluation Metrics

Rice Leaf Disease detection models were evaluated using several key metrics: **Accuracy**, **Precision**, **Recall**, and **F1-Score**. Based on these metrics, we are able to determine how well the models perform when it comes to accurately classifying images.

### **3.11.1.** Accuracy

Accuracy measures how many correct predictions were made (both true positives and true negatives) in comparison to total predictions as in Eq. (3.1). Data classification accuracy is a good indicator of a model's performanceStallings and Gillmore 1971.

$$Accuracy = \frac{TP + TN}{TP + TN + FP + FN}$$
 (3.1)

### 3.11.2. Precision

The precision of the model measures how many true positive predictions it makes and can be calculated through Eq. (3.2). When false positives are expensive, misclassifying a healthy leaf as diseased may result in unnecessary treatments.

$$Precision = \frac{TP}{TP + FP}$$
 (3.2)

### 3.11.3. Recall

**Recall** It measures how accurate a model is in identifying actual positive instances as in Eq. (3.3). The metric is useful when it is crucial to detect all positive cases, even at the cost of some false positives (e.g., detecting all diseased leaves to prevent further spread).

$$Recall = \frac{TP}{TP + FN}$$
 (3.3)

### 3.11.4. F1-Score

The F1-Score provides a balance between precision and recall as shown in Eq. (3.4). It is especially useful when there are fewer positive samples than negative samples..

$$F1-Score = 2 \times \frac{Precision \times Recall}{Precision + Recall}$$
 (3.4)

# 4. CHAPTER IV: QUALITY AND RESULTS

The results of the evaluation of CNN Mobile Net, my deep learning model for classifying rice leaf diseases, are presented in this section. For training purposes, categorical cross entropy was used for the loss function. A variety of metrics were used to assess the performance of each model, including accuracy, precession recall, and F 1. Google Collaboratory projects were equipped with 12GB of RAM, 16GB of NVIDIA Tesla T4 GPU RAM, and 2.20GHz Intel(R) Xeon(R) CPUs.

### 4.1. CNN Model Training and Loss

The deep learning model, trained on 120 high-resolution rice leaf images, achieved 100% accuracy on the training data with a very low loss of 1.7592e-04. However, its performance on the validation dataset was lower, with a 75% accuracy and a validation loss of 2.0851. This indicates good fitting to the training data but difficulty generalizing to unseen data, as shown in Fig4.1 and Fig 4.2. Early disease detection is essential for effective crop management.

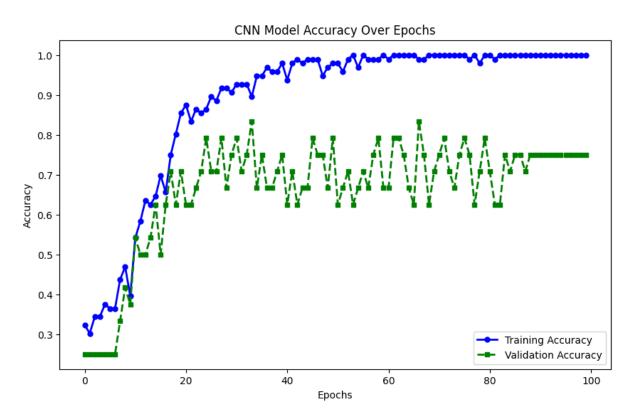


Fig. 4.1. CNN Model Accuracy Over Epochs

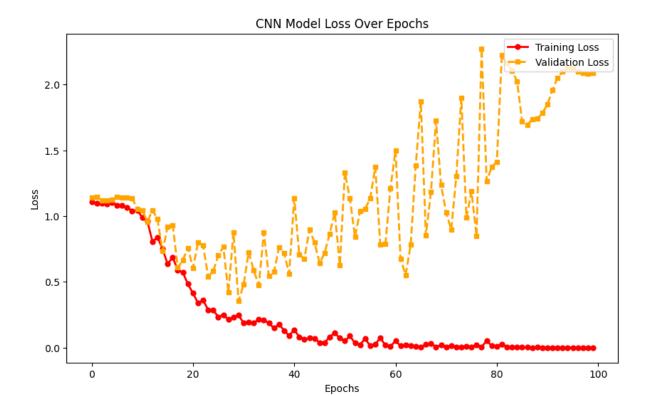


Fig. 4.2. CNN Model Loss Over Epochs

### 4.2. MobileNet Model Training and Loss

A dataset of 120 high-resolution images of rice leaves was used to train the MobileNet model. Images were classified as Bacterial Leaf Blight, Brown Spot and Leaf Smut. Fig show training and loss 4.3. It is important to emphasize that the model achieved 100% accuracy during training with a loss of only 0.0022, which demonstrated a very effective learning process. Each class contained 40 images. Fig 4.4 loss of the model On the validation dataset, MobileNet maintained a high accuracy of 91.67% and a validation loss of 0.5831, showcasing its strong generalization capabilities.

#### 4.2.1. MobileNet model Results

MobileNet was applied to 120 high-resolution images of rice leaf diseases, Bacterial Leaf Blight, Brown Spot, and Leaf Smut 100% training accuracy with a loss of 0.0022. On the validation set, it maintained a strong accuracy of 91.67% with a validation loss of 0.5831. The classification report indicates impressive results across all disease categories, with perfect detection of Bacterial Leaf Blight (precision, recall, and F1-score of 1.00), balanced performance for Brown Spot (precision, recall, and F1-score of 0.91), and strong results for Leaf Smut (precision and recall of 0.86, F1-score of 0.86). With an overall test accuracy of 92%, macro average F1-score of 0.92, and weighted average F1-score of 0.92 as shown in Table 4.1 and Fig. 4.5. The model demonstrates strong generalization and reliability, making MobileNet a

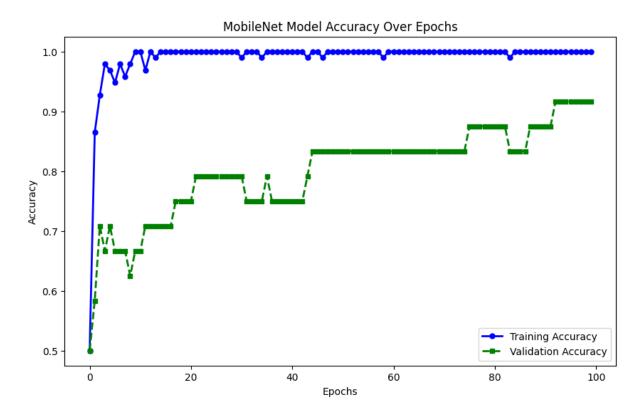


Fig. 4.3. MobileNet Model Accuracy Epochs

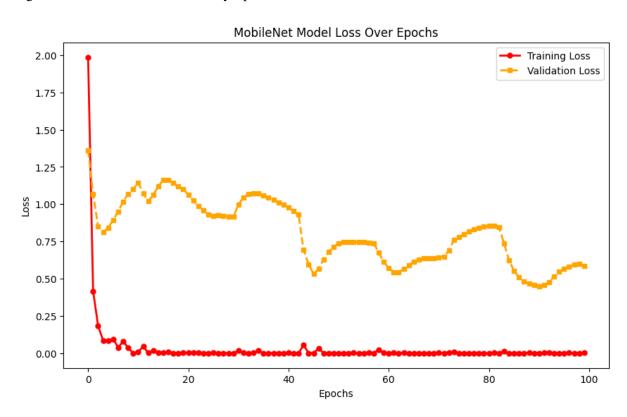


Fig. 4.4. MobileNet Model loss Epochs

promising candidate for real-world crop disease detection applications.

Table 4.1. MobileNet Classification Report

Class	Precision	Recall	F1-Score	Support	
Class					
Bacterial leaf	1.00	1.00	1.00	6	
blight	1.00	1.00	1.00	0	
Brown spot	0.91	0.91	0.91	11	
Leaf smut	0.86	0.86	0.86	7	
Accuracy			0.92	24	
Macro avg	0.92	0.92	0.92	24	
Weighted avg	0.92	0.92	0.92	24	

#### **MobileNet Classification Report for Rice Leaf Diseases**

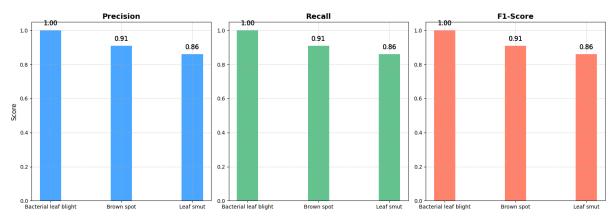


Fig. 4.5. MobileNet Model Results Graph

### 4.2.2. Confusion Matrix of CNN model

Three rice leaf diseases are predicted by CNN: Bacterial Leaf Blight, Brown Spot, and Leaf Smut. The model correctly classified all six instances of Bacterial Leaf Blight. For Brown Spot, there is some confusion, with one sample misclassified as Bacterial Leaf Blight and four as Leaf Smut, resulting in reduced recall (0.55) as shown in Fig. 4.6. Leaf Smut exhibits better recognition, with only one sample misclassified as Brown Spot, achieving a recall of 0.86.

#### 4.2.3. Confusion matrix of MobileNet

This confusion matrix compares MobileNet with three rice leaf diseases: Bacterial Leaf Blight, Brown Spot, and Leaf Smut. Bacterial Leaf Blight was correctly identified by the model in all six instances as shown in Fig. 4.7. Brown Spot also showed strong performance, with 10 out of 11 samples correctly classified, and only one instance misclassified as Leaf Smut. For Leaf Smut, six samples were correctly classified, but one was misclassified as Brown Spot.

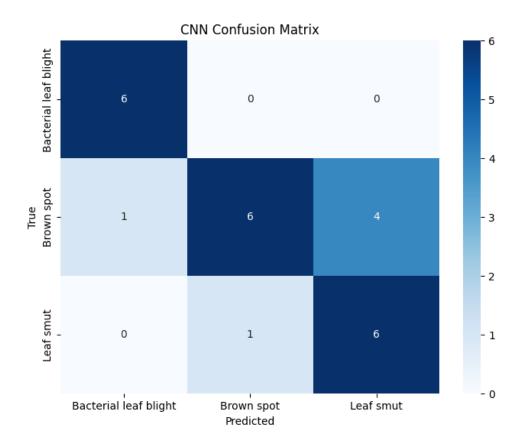


Fig. 4.6. Confusion Matrix of CNN model

**Training Performance** Both models achieved excellent training accuracy, with the CNN model achieving 100% accuracy and the MobileNet model also reaching 100%. This indicates that both models learned the features of the training data very well. As compared to CNN and MobileNet models as shown in Table4.2 and Fig.4.9, However, the loss values differ significantly, with the CNN model achieving an extremely low training loss of 1.7592e-04, suggesting that it had a very tight fit to the training data. In contrast, the MobileNet model achieved a slightly higher training loss of 0.0022, indicating a slightly less overfitted model.

**Validation Performance** On the validation set, MobileNet showed superior generalization with a validation accuracy of 91.67%, significantly outperforming the CNN model, which had a validation accuracy of 75%. In this case, CNN overfitted to unseen data, even though it fitted well to the training data. Nevertheless, MobileNet's better performance on the validation set proves that it can generalize well to new, unseen data, making it a more reliable model in practice.

### 4.2.4. Model Accuracy Comparison for Rice Leaf Disease Detection

The results of the model accuracy comparison for rice leaf disease detection are presented in Fig.4.8 and Table 4.9. The analysis highlights the performance differences among traditional machine learning algorithms, modern convolutional neural networks (CNNs), and proposed

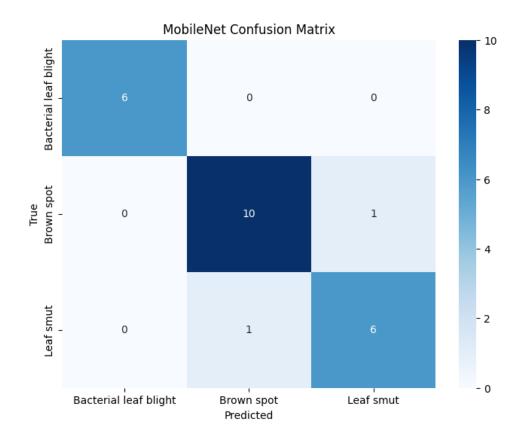


Fig. 4.7. Confusion matrix of MobileNet Model

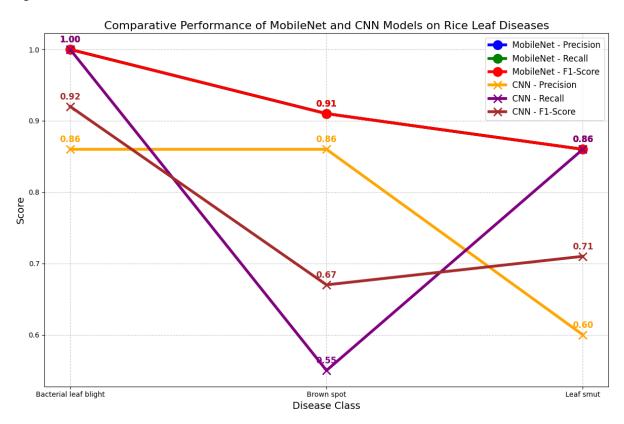


Fig. 4.8. Comparative Classification Results

Table 4.2. Comparative Classification Results for MobileNet and CNN Models on Rice Leaf Diseases

Class	MobileNet			CNN				
Class	Precision	Recall	F1-Score	Support	Precision	Recall	F1-Score	Support
Bacterial leaf	1.00	1.00	1.00	6	0.86	1.00	0.92	6
blight	1.00	1.00	1.00	O	0.80	1.00	0.92	0
Brown spot	0.91	0.91	0.91	11	0.86	0.55	0.67	11
Leaf smut	0.86	0.86	0.86	7	0.60	0.86	0.71	7
Accuracy			0.92	24			0.75	24
Macro avg	0.92	0.92	0.92	24	0.77	0.80	0.77	24
Weighted avg	0.92	0.92	0.92	24	0.78	0.75	0.74	24

methodologies Shown in Table 4.2.

Table 4.3. Comparison of Models for Rice Leaf Disease Detection

References	Model Names	Accuracy (%)
Masood et al. 2020	Mask R-CNN	87.6%
Haque et al. 2022	YOLOv5	76.0%
Murugan et al. 2022	Convolutional Neural Network (CNN)	88.28%
Vasantha, Kiranmai, and Krishna 2021	Various Machine Learning Algorithms	90%–91%
Kiratiratanapruk et al. 2022	YOLOv4 with Image Tiling Technique	87.56% to 91.14%
Trinh et al. 2024	Modified YOLOv8	89.9%
Proposed Approach	Mobilenet	92%

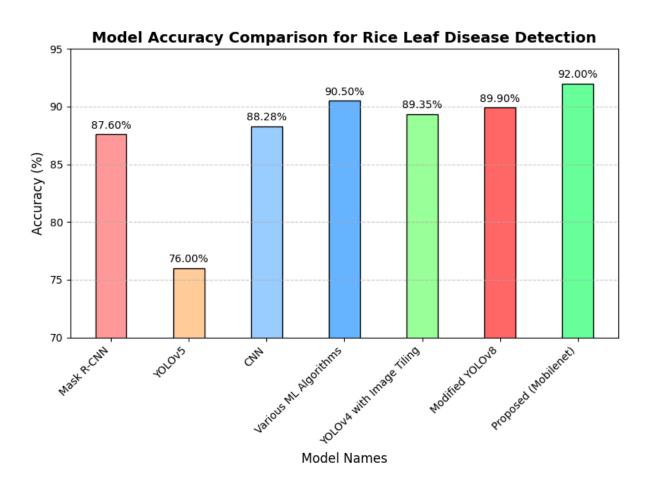


Fig. 4.9. Comparison of Models with Mobil net

## 5. CHAPTER 5: EVALUATION AND CONCLUSIONS

### 5.1. Final Evaluation

The experimental results demonstrate that both CNN and MobileNet models performed well in classifying rice leaf diseases, achieving high training accuracy (100%) for both models. MobileNet, MobileNet also outperformed CNN in terms of precision, recall, and F1-score, making it a more reliable model for practical disease detection in rice crops.

### 5.2. Project Management

The project was executed efficiently using Google Collaboratory with a strong hardware setup (12GB RAM, 16GB GPU RAM, and 2.20GHz Intel CPUs), which enabled the successful training of the CNN and MobileNet models. The task was divided into training, evaluation, and performance analysis phases, ensuring that resources were effectively allocated to enhance model performance. Regular monitoring of model performance and iterative adjustments were made to optimize results.

### 5.3. Insights Gained

This study highlights the importance of model generalization, as MobileNet's superior performance on the validation set emphasizes its ability to handle unseen data effectively. The challenges of overfitting, particularly in CNN, underscore the need for careful training and validation practices to ensure robust model deployment. Additionally, the importance of dataset diversity and quality is evident in achieving reliable results.

### 5.4. Comparison to Literature

In comparison to similar studies, MobileNet's strong performance in classification and generalization aligns with findings from other research that emphasizes the efficiency of lightweight models in resource-constrained environments. The CNN model's overfitting issue in this study contrasts with some literature that highlights the potential for CNNs in achieving high accuracy with proper regularization and augmentation techniques.

### **5.5.** Reflection on Challenges

A key challenge in this project was managing overfitting in the CNN model. Despite achieving perfect training accuracy, its lower validation performance highlighted the limitations of its generalization capabilities. Another challenge was ensuring balanced training across all disease categories, as misclassifications in certain classes (such as Brown Spot) impacted overall model performance.

#### 5.6. Conclusion

A comparative analysis of CNN and MobileNet models for rice leaf disease organisation shows their assets and weaknesses. CNN established overfitting contempt high training accuracy, as showed by its meaningfully lower training loss and inferior validation performance. MobileNet, on the other hand, outperformed CNN in terms of validation accuracy and oversimplification, making it more reliable. Also, MobileNet's superior performance across all disease categories, particularly Brown Spot and Leaf Smut, underscores its potential as an effective rice disease detection tool. MobileNet's aptitude to simplify to new data makes it a more healthy solution for automated rice disease diagnosis. With deep learning in agricultural automation, disease detection accuracy can be improved and more efficient and climbable crop management systems can be industrialised.

#### 5.7. Future Work

MobileNet and CNN models could be further assessed by counting more diverse rice diseases and ecological conditions. Incorporating data increase techniques could help alleviate overfitting in CNN models, possibly improving their performance on unseen data. Moreover, exploring hybrid models that combine the strengths of both CNN and MobileNet could lead to even more precise and generalized classification systems. Investigating the integration of these models into real-time agricultural systems, such as mobile apps or drones, could help ease the extensive adoption of automatic disease detection in rice farming. Additional research into transfer learning and fine-tuning of pre-trained models on large-scale agricultural datasets could also enhance the models' efficiency in diverse physical regions. Lastly, the addition of visible and imperceptible symptoms of disease through multi-modal data dispensation (e.g., thermal, multispectral) could recover detection accuracy and enlarge the range of disease diagnoses in precision agriculture.

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### **APPENDIX**

```
1
2
3
   import numpy as np
4
   import mataplotlib.pyplot as plt
   import seaborn as sns
   import tensorflow as tf
   from sklearn.metrcis import
8
   classification_report, confusion_matrix, accuracy_score, precision_score
       , recall_score, f1_score
   from tensorflow.keras.utils import to_cateorical
10
   from tensorflow.keras.callbacks import ModelCheckpoint
11
   import os
12
   import cv2
13
   import albumentations as albu
14
   from albumentations import Compose, ShiftScaleRotate, Resize
15
   from sklearn.utils import shuffle
16
   from keras.models import Sequential
17
   from sklearn.model_selection import train_test_split
18
   from keras.layers import Conv2D, MaxPooling2D, Flatten, Dense, Dropout
19
   from tensorflow.keras.applications import MobileNet
20
   from tesndorflow.keras.layer import
21
   GlobalAveragePooling2D, BatchNormalization
23
   from tensorflow.keras.optimizers import adam
24
25
   # Setup Environment
26
   input_shape_2D = (224, 224) # For 2D images
27
   input_shape_3D = (224, 224, 3) # For RGB images
28
   seed = 1
29
   batch\_size = 32
30
   epochs = 100
31
32
   from google.colab import drive
33
   drive.mount('/content/drive')
34
35
   # Load Image data
36
   data = tf.keras.utils.image_dataset_from_directory(
37
        directory="/content/drive/MyDrive/rice_leaf_diseases", # Adjust path
38
            as necessary
        labels='inferred', label_mode='int', class_names=None, color_mode='
39
        image_size=(224, 224), seed=1
40
41
   )
```

```
# Print class names
    class_names = data.class_names
    class_names
45
46
    # Visualize sample images
47
   plt.figure(figsize=(10, 10))
48
    for images, labels in data.take(1):
49
        for i in range(25):
50
            plt.subplot(5, 5, i + 1)
51
            plt.imshow(images[i].numpy().astype('uint8'))
52
            plt.title(class_names[labels[i]])
53
            plt.axis('off')
54
   plt.tight_layout()
55
   plt.show()
56
57
   # Data Preprocessing
58
   # Prepare data for training and testing
59
   X = \Gamma 
60
   y = []
61
    for images, labels in data:
62
        X.append(images.numpy())
63
        y.append(labels.numpy())
65
   X = np.concatenate(X, axis=0)
   y = np.concatenate(y, axis=0)
67
68
   # Normalize images
69
   X = X.astype('float32') / 255
70
71
   # Use train_test_split to randomly split the data
72
73
   X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
        random_state=42)
   # Convert labels to one-hot encoding
75
   y_train = to_categorical(y_train, len(class_names))
76
   y_test = to_categorical(y_test, len(class_names))
77
78
   # Build CNN Model
79
   model_cnn = Sequential()
80
   model_cnn.add(Conv2D(filters=16, kernel_size=3, padding='same', strides
81
       =1, activation='relu', input_shape=(224, 224, 3)))
   model_cnn.add(MaxPooling2D(pool_size=(3, 3)))
82
83
   model_cnn.add(Conv2D(filters=32, kernel_size=3, padding='same', strides
84
       =1, activation='relu'))
   model_cnn.add(MaxPooling2D(pool_size=(3, 3)))
85
86
   model_cnn.add(Conv2D(filters=64, kernel_size=3, padding='same', strides
87
       =1, activation='relu'))
   model_cnn.add(MaxPooling2D(pool_size=(3, 3)))
```

```
89
    model_cnn.add(Conv2D(filters=128, kernel_size=3, padding='same', strides
       =1, activation='relu'))
    model_cnn.add(MaxPooling2D(pool_size=(2, 2)))
91
92
    model_cnn.add(Conv2D(filters=256, kernel_size=3, padding='same', strides
93
       =1, activation='relu'))
    model_cnn.add(MaxPooling2D(pool_size=(2, 2)))
94
95
    model_cnn.add(Dropout(0.3))
96
    model_cnn.add(Flatten())
    model_cnn.add(Dropout(0.5))
    model_cnn.add(Dense(250, activation='relu'))
    model_cnn.add(Dense(100, activation='relu'))
100
    model_cnn.add(Dense(len(class_names), activation='softmax'))
101
102
    model_cnn.summary()
103
104
    # Compile the CNN model
105
    model_cnn.compile(optimizer='Adam', loss='categorical_crossentropy',
106
       metrics=['accuracy'])
107
    # Define ModelCheckpoint callback
108
    check_pointer = ModelCheckpoint(
109
        filepath='cnn_model.best.weights.h5', monitor='val_accuracy',
110
            save_best_only=True,
        mode='auto', save_weights_only=True
111
    )
112
113
    # Train CNN model
114
    history_cnn = model_cnn.fit(
115
        X_train, y_train, batch_size=batch_size, epochs=epochs,
116
            validation_data=(X_test, y_test),
        callbacks=[check_pointer], verbose=1
117
    )
118
119
    # Visualize training and validation accuracy
120
    plt.figure(figsize=(10, 6))
121
    plt.plot(history_cnn.history['accuracy'], label='Training Accuracy',
122
        color='blue', linewidth=2, marker='o', markersize=5)
    plt.plot(history_cnn.history['val_accuracy'], label='Validation Accuracy
        ', color='green', linewidth=2, linestyle='--', marker='s', markersize
       =5)
    plt.title("CNN Model Accuracy Over Epochs")
124
    plt.xlabel("Epochs")
125
    plt.ylabel("Accuracy")
126
    plt.legend(loc="lower right")
127
    plt.show()
128
129
    # Visualize training and validation loss
130
```

```
plt.figure(figsize=(10, 6))
131
    plt.plot(history_cnn.history['loss'], label='Training Loss', color='red'
132
        , linewidth=2, marker='o', markersize=5)
    plt.plot(history_cnn.history['val_loss'], label='Validation Loss', color
133
        ='orange', linewidth=2, linestyle='--', marker='s', markersize=5)
    plt.title("CNN Model Loss Over Epochs")
134
    plt.xlabel("Epochs")
135
    plt.ylabel("Loss")
136
    plt.legend(loc="upper right")
137
    plt.show()
138
139
    # Predictions for CNN model
    y_pred_cnn = model_cnn.predict(X_test)
141
    predicted_classes_cnn = np.argmax(y_pred_cnn, axis=1)
142
143
    # Classification Report for CNN
144
    print("CNN Classification Report:\n", classification_report(np.argmax(
145
       y_test, axis=1), predicted_classes_cnn, target_names=class_names))
146
    # Confusion Matrix for CNN
147
    cm_cnn = confusion_matrix(np.argmax(y_test, axis=1),
148
        predicted_classes_cnn)
    plt.figure(figsize=(8, 6))
    sns.heatmap(cm_cnn, annot=True, fmt='d', cmap='Blues', xticklabels=
150
        class_names, yticklabels=class_names)
    plt.title('CNN Confusion Matrix')
151
    plt.xlabel('Predicted')
152
    plt.ylabel('True')
153
    plt.show()
154
```

```
plt.show()
214
    # Predictions and Evaluation for MobileNet
215
    y_pred_mobilenet = model_mobilenet.predict(X_test)
    predicted_classes_mobilenet = np.argmax(y_pred_mobilenet, axis=1)
217
218
    # Classification Report for MobileNet
219
    print("MobileNet Classification Report:\n", classification_report(np.
220
        argmax(y_test, axis=1), predicted_classes_mobilenet, target_names=
        class_names))
221
    # Confusion Matrix for MobileNet
222
    cm_mobilenet = confusion_matrix(np.argmax(y_test, axis=1),
223
        predicted_classes_mobilenet)
    plt.figure(figsize=(8, 6))
224
    sns.heatmap(cm_mobilenet, annot=True, fmt='d', cmap='Blues', xticklabels
225
        =class_names, yticklabels=class_names)
    plt.title('MobileNet Confusion Matrix')
226
    plt.xlabel('Predicted')
227
    plt.ylabel('True')
228
    plt.show()
229
                                                                                 43
```

```
# Calculate accuracy, precision, recall, F1 score
    accuracy_cnn = accuracy_score(np.argmax(y_test, axis=1),
157
       predicted_classes_cnn)
    precision_cnn = precision_score(np.argmax(y_test, axis=1),
158
       predicted_classes_cnn, average='weighted')
    recall_cnn = recall_score(np.argmax(y_test, axis=1),
       predicted_classes_cnn, average='weighted')
    f1_cnn = f1_score(np.argmax(y_test, axis=1), predicted_classes_cnn,
160
       average='weighted')
161
    print(f"CNN Accuracy: {accuracy_cnn}")
162
    print(f"CNN Precision: {precision_cnn}")
163
    print(f"CNN Recall: {recall_cnn}")
    print(f"CNN F1 Score: {f1_cnn}")
165
    # Load pre-trained MobileNet model (without top classification layers)
167
    base_model = MobileNet(weights='imagenet', include_top=False,
168
       input_shape=(224, 224, 3))
169
170
    # Freeze the layers of the base model
```

```
base_model.trainable = False
171
    # Add custom layers on top of MobileNet
173
    x = base_model.output
    x = GlobalAveragePooling2D()(x)
    x = Dense(1024, activation='relu')(x)
    x = BatchNormalization()(x)
    x = Dropout(0.5)(x)
    x = Dense(512, activation='relu')(x)
    x = Dropout(0.5)(x)
    x = Dense(len(class_names), activation='softmax')(x)
181
182
    # Define the MobileNet model
    model_mobilenet = Model(inputs=base_model.input, outputs=x)
184
185
    # Compile the MobileNet model
186
    model_mobilenet.compile(optimizer=Adam(), loss='categorical_crossentropy
187
        ', metrics=['accuracy'])
188
    # Train MobileNet model
189
    history_mobilenet = model_mobilenet.fit(
        X_train, y_train, batch_size=32, epochs=100,
191
        validation_data=(X_test, y_test), callbacks=[check_pointer], verbose
           =1
193
```

```
base_model.trainable = False
172
    # Add custom layers on top of MobileNet
    x = base_model.output
174
    x = GlobalAveragePooling2D()(x)
175
    x = Dense(1024, activation='relu')(x)
    x = BatchNormalization()(x)
    x = Dropout(0.5)(x)
    x = Dense(512, activation='relu')(x)
    x = Dropout(0.5)(x)
180
    x = Dense(len(class_names), activation='softmax')(x)
181
182
    # Define the MobileNet model
183
    model_mobilenet = Model(inputs=base_model.input, outputs=x)
185
    # Compile the MobileNet model
186
    model_mobilenet.compile(optimizer=Adam(), loss='categorical_crossentropy
187
        ', metrics=['accuracy'])
    # Train MobileNet model
189
    history_mobilenet = model_mobilenet.fit(
190
        X_train, y_train, batch_size=32, epochs=100,
191
        validation_data=(X_test, y_test), callbacks=[check_pointer], verbose
192
            =1
193
194
```

```
# Visualize training and validation accuracy for MobileNet
    plt.figure(figsize=(10, 6))
196
    plt.plot(history_mobilenet.history['accuracy'], label='Training Accuracy
197
        ', color='blue', linewidth=2, marker='o', markersize=5)
    plt.plot(history_mobilenet.history['val_accuracy'], label='Validation
198
       Accuracy', color='green', linewidth=2, linestyle='--', marker='s',
       markersize=5)
    plt.title("MobileNet Model Accuracy Over Epochs")
199
    plt.xlabel("Epochs")
    plt.ylabel("Accuracy")
    plt.legend(loc="lower right")
202
    plt.show()
203
    # Visualize training and validation loss for MobileNet
205
    plt.figure(figsize=(10, 6))
    plt.plot(history_mobilenet.history['loss'], label='Training Loss', color
207
       ='red', linewidth=2, marker='o', markersize=5)
    plt.plot(history_mobilenet.history['val_loss'], label='Validation Loss',
208
        color='orange', linewidth=2, linestyle='--', marker='s', markersize
    plt.title("MobileNet Model Loss Over Epochs")
209
    plt.xlabel("Epochs")
210
    plt.ylabel("Loss")
211
    plt.legend(loc="upper right")
212
```