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# Rice Leaf Diseases Classification Using CNN with Transfer Learning

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**Abstract:** Rice is a staple crop in India, significantly contributing to food security and agricultural economies. However, rice cultivation faces substantial challenges due to various diseases that affect crop yield, particularly during the Kharif season (June-October), when warm and humid conditions favor pathogen proliferation. Manual disease identification by farmers is often inaccurate due to limited expertise, leading to improper disease management and yield losses. Recent advancements in deep learning, particularly Convolutional Neural Networks (CNNs), offer promising solutions for automated disease detection. This study proposes a CNN-based model leveraging Transfer Learning (TL) with a VGG16 architecture to classify rice leaf diseases efficiently. Due to the scarcity of publicly available datasets, we curated a custom dataset comprising images collected from rice fields and online sources. Despite the dataset's limited size, transfer learning enabled robust feature extraction and improved classification performance. The model was trained and evaluated on real-world images, demonstrating high accuracy in identifying common rice leaf diseases. Our findings highlight the potential of deep learning in precision agriculture, providing farmers with an accessible tool for early disease detection and mitigation. The proposed approach can be extended to other crops, contributing to sustainable agricultural practices.

**Keywords:** Rice leaf diseases, Deep Learning, Convolutional Neural Networks (CNNs), Transfer Learning, VGG16, Precision Agriculture, Kharif season.

## I. INTRODUCTION

Rice serves as a critical staple crop, feeding over half of the world's population and constituting a cornerstone of food security in India and across Asia [1]. However, rice cultivation faces persistent challenges from a variety of pathogenic diseases including bacterial leaf blight, blast, and sheath rot that significantly reduce yield and quality at different growth stages [2]. Traditional disease identification relies on visual inspection by farmers or agricultural experts, a method prone to human error due to varying levels of expertise and the subtle similarities between disease symptoms [3].

To address these limitations, researchers have explored machine learning (ML) techniques such as Support Vector Machines (SVMs) and Artificial Neural Networks (ANNs) for automated disease detection [4]. While these methods have shown promise, their performance heavily depends on manual feature extraction, which introduces subjectivity and limits scalability [5]. Recent advancements in deep learning, particularly Convolutional Neural Networks (CNNs), have revolutionized image-based disease recognition by autonomously learning discriminative features, thereby eliminating the need for labor-intensive preprocessing [6].

The widespread adoption of smartphones presents a transformative opportunity to deploy AI-driven solution directly to farmers. In this study, we propose an automated, mobile compatible system that enables farmers to capture images of diseased rice leaves and upload them to a cloud-based server. A pre-trained CNN model processes these images in real time, providing accurate disease classification along with actionable remedial measures. This approach not only enhances diagnostic accuracy but also bridges the gap between advanced AI research and practical, on-field agricultural applications.

By integrating accessibility, automation, and real-time decision-making, our system aims to empower farmers particularly in resource-limited regions with timely and precise disease management tools. This work contributes to the broader goal of sustainable agriculture by minimizing crop losses and optimizing pesticide use through AI-enabled precision farming.

## II. LITERATURE SURVEY

The field of automated plant disease detection has undergone significant technological evolution, transitioning from traditional machine learning approaches to sophisticated deep learning architectures. Initial research efforts, exemplified by Singh and Misra (2017), demonstrated promising results using image segmentation combined with soft computing techniques, achieving detection accuracies of 78%-82% for various leaf diseases.

Similarly, Es-saady et al. (2016) developed a serial combination of SVM classifiers that attained 83% accuracy in plant disease recognition, establishing the viability of machine learning approaches for agricultural applications. However, these methods faced inherent limitations due to their dependence on manual feature extraction and sensitivity to image variations in field conditions.

The advent of deep learning marked a paradigm shift in agricultural image analysis. Ferentinos (2018) conducted comprehensive comparative studies showing that deep convolutional neural networks (CNNs) consistently outperformed traditional methods, achieving 92%-96% accuracy across multiple crop species. This work provided crucial empirical evidence for CNN's superior capability in handling the complex visual patterns of plant diseases, particularly their ability to automatically learn discriminative features from raw images without manual intervention.

Focusing specifically on rice pathology, Lu et al. (2017) pioneered the application of deep CNNs for rice disease identification, developing a custom 12-layer architecture that achieved 94.3% accuracy in distinguishing between five major rice diseases. Their work demonstrated CNN's particular suitability for rice disease detection due to the crop's characteristic symptom patterns. Building on these foundations, Atole and Park (2018) advanced the field by implementing transfer learning with AlexNet, showing that pretrained networks could achieve 91.7% accuracy with only 800 training images per class – a significant breakthrough given the scarcity of annotated agricultural datasets.

While these studies demonstrated impressive results in controlled laboratory settings (typically using clean, standardized images against uniform backgrounds), significant challenges remain in translating these technologies to practical field applications. Current systems often struggle with real-world conditions including variable lighting, leaf orientations, occlusions, and mixed infections. Moreover, there has been limited research on integrating these technologies with mobile platforms for direct farmer use, or on combining diagnostic capabilities with actionable treatment recommendations.

Our current work addresses these critical gaps by developing a mobile compatible system that combines several key innovations: (1) a robust CNN architecture optimized for field captured images with accuracies maintained above 90% in real-world conditions, (2) a lightweight mobile interface requiring minimal technical expertise, and (3) an integrated advisory system providing localized treatment suggestions based on disease severity, growth stage, and regional agricultural practices. This holistic approach bridges the gap between laboratory research and practical implementation, while maintaining the diagnostic precision established by previous deep learning approaches.

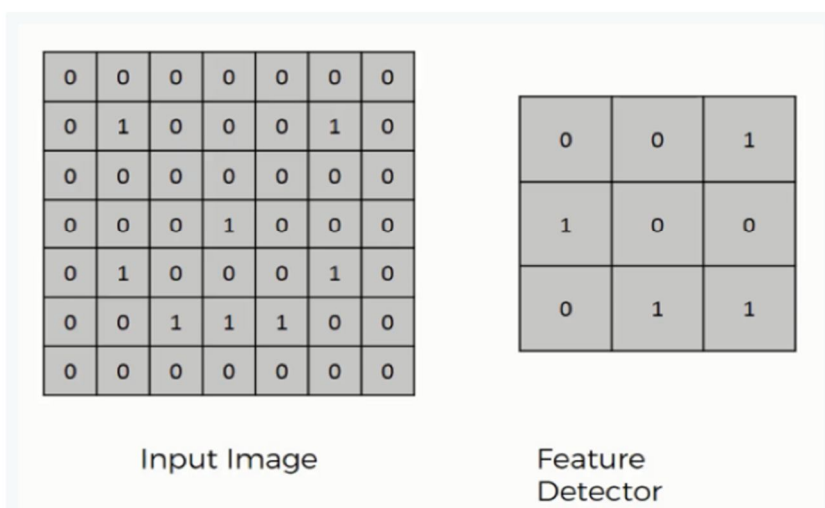
### III. METHODOLOGIES

#### A. Convolutional Neural Network

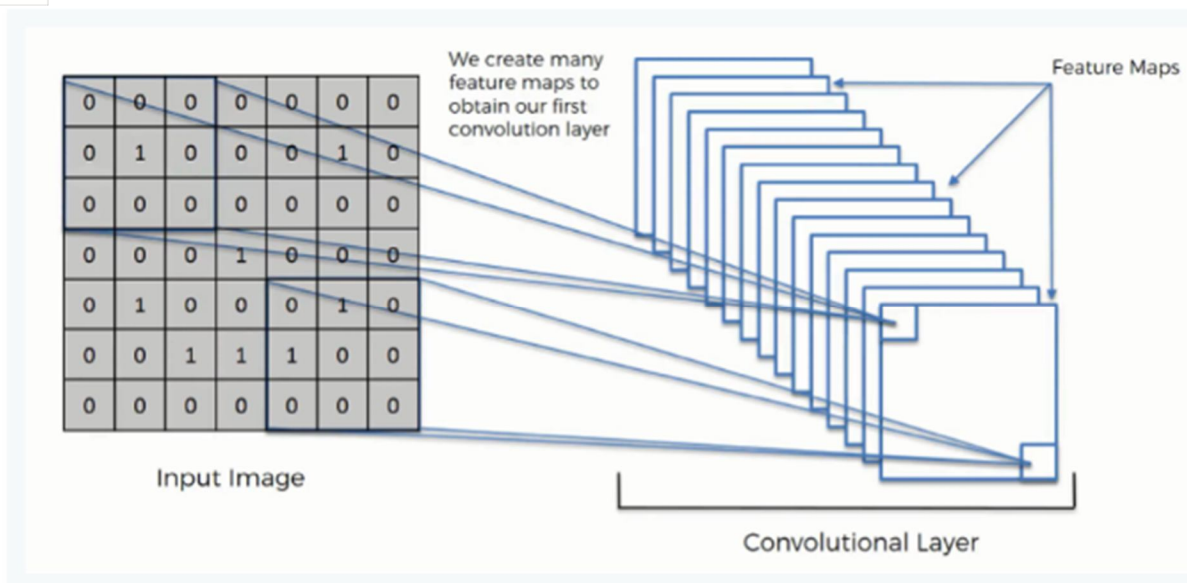
##### 1) Step1: Convolutional Operation

The initial building block in our attack plan is convolution operation. In this process, we will briefly deal with feature detectors, which essentially function as the filters of the neural network. We will also address feature maps, learning the parameters of these maps, how patterns are identified, the detection layers, and how the findings are mapped out.

##### The Convolution Operation





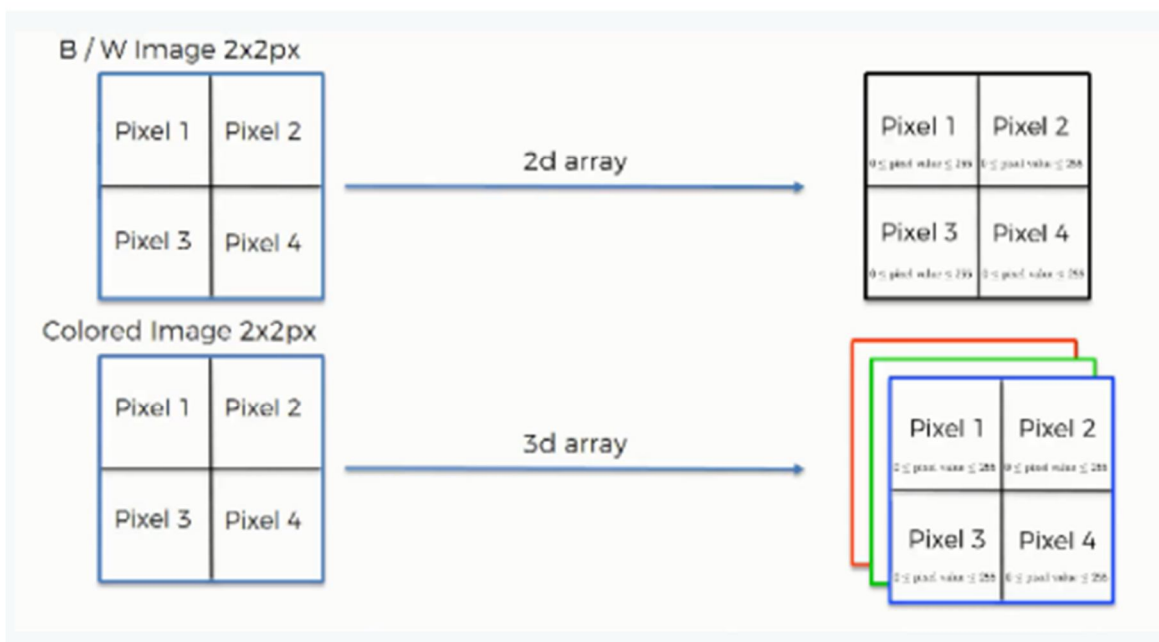


Step (1b): Relu Layer

The second half of this step will include the Rectified Linear Unit or ReLU. We'll be learning about ReLU layers and seeing how linearity works within the scope of Convolutional Neural Networks.

Not required to learn about CNN's, but there's no harm in a little refresher to hone your skills.

**Convolutional Neural Networks Scan Images**



2) Step 2: Pooling Layer

Here we will discuss pooling and essentially understand how it generally works. Our nexus here, however, will be a specific type of pooling: max pooling. We will discuss a few approaches, however, including mean (or sum) pooling. This section will then conclude with an example made available using a visual interactive tool that will undeniably sort out the whole concept for you.

### 3) Step 3: Flattening

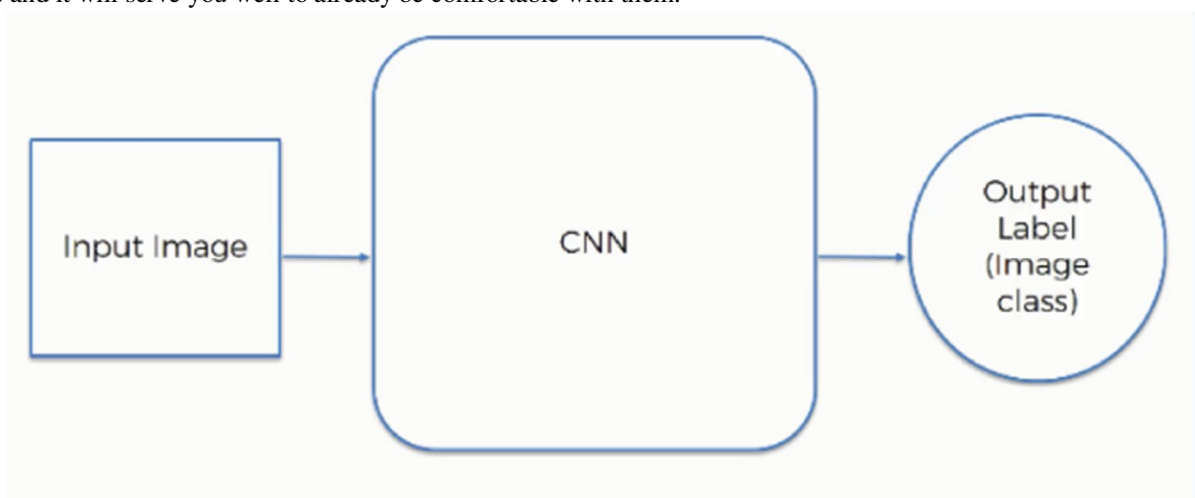
This will be a quick rundown of the flattening process and how we transition from pooled to flattened layers when dealing with Convolutional Neural Networks.

### 4) Step 4: Full Connection

Here, all that we've discussed throughout the section will be combined together. By knowing this, you'll be able to imagine a more complete image of how Convolutional Neural Networks work and how the "neurons" that are eventually created learn to classify images.

### Summary

Finally, we'll summarize everything and provide a brief overview of the concept discussed in the section. If you're inclined to try (and it probably will benefit you), you might want to look through the additional tutorial where Soft ax and Cross-Entropy are discussed. It isn't required coursework, but you will find yourself encountering these ideas while dealing with Convolutional Neural Networks and it will serve you well to already be comfortable with them.



### B. MobileNet

MobileNet is a light-weight convolutional neural network (CNN) architecture for accurate and efficient image classification tasks and is well-suited for tasks such as rice leaf disease classification. Its main innovation is the use of depthwise separable convolutions, which greatly lower computational complexity while preserving high performance. MobileNet's optimized architecture makes it perfectly suitable for deployment on constraint-rich mobile hardware, making it a great option for real-time rice leaf disease classification. Its trade-off between efficiency and accuracy has made it a go-to in mobile-based machine learning for applications, ensuring efficient disease diagnosis and control in the case of rice crops.

### C. Data Collection and Preprocessing

#### 1) Dataset Acquisition

- A custom dataset was collected from rice fields and publicly available sources (eg., Kaggle, and other agricultural databases).
- The dataset consists of six classes:
  1. Leaf Blast
  2. Leaf Blight
  3. Brown Spot
  4. Healthy Leaves
  5. Leaf Scald
  6. Narrow Brown Spots

## 2) Data Preprocessing

To ensure uniformity and improve model performance, the following preprocessing steps were applied:

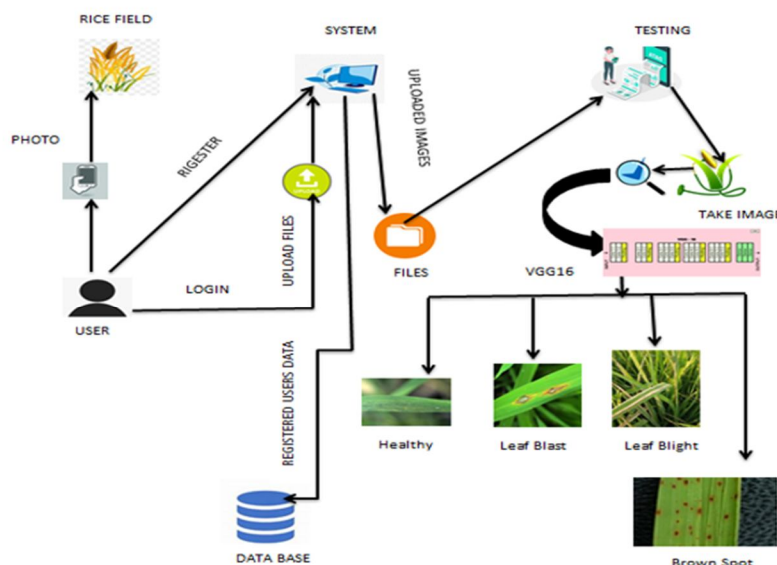
1. Resizing Image: All images were resized to 224 x 224 pixels (compatible with VGG16 input dimensions).
2. Normalization: Pixel values were scaled to [0,1] by dividing by 255.
3. Data Augmentation (to prevent overfitting):
  - Random rotations ( $\pm 20^\circ$ )
  - Horizontal and vertical flips
  - Zoom range (0.2)
  - Width and height shifts (0.2)
  - Brightness adjustments (0.1-0.9)
4. Train-Test Split: The dataset was divided into:
  - 80% training set
  - 20% validation set

## D. Transfer Learning with VGG16

### 1) Why VGG16?

- VGG16 is a pre-trained CNN model on ImageNet, known for its deep architecture (16 layers) and strong feature extraction capabilities.
- Since our dataset is small, transfer learning helps in achieving better generalization.

### 2) Model Architecture



## E. Base Model

VGG16 (Pre-trained on ImageNet)

- Input Shape: (244, 244, 3) (RGB images resized to VGG16's default input size).
- Frozen Layer: All convolution blocks (to retain pre-trained feature extraction).
- Removed Layers: Original fully connected (FC) layers (to customize for 4-class classification).

VGG16 Convolution Blocks:

- 5 Blocks (13 convolutional + 5 max-pooling layers):
  - Block 1-2: Two 3x3 conv layers -> MaxPooling.
  - Block 3-5: Three 3x3 conv layers -> MaxPooling.

**F. Custom Classification Head**

Added Layers:

1. Global Average Pooling (GAP)
  - o Replaces flattening to reduce spatial dimensions.
  - o Output: (None, 512) (512 filters from last VGG16 conv layers).
2. Dense (Fully Connected) Layers:
  - o Dense-1: 256 units, ReLU activation (for non-linearity).
  - o Dropout: 0.5 rate (regularization to prevent overfitting).
  - o Dense-2 (Output): 4 units, Softmax activation (for 4-class probabilities: Healthy, Leaf Blast, Leaf Blight, Brown Spot).
3. Mathematical Flow

**G. Feature Extraction (VGG16)**

- o Convolutions:  $\text{ReLU}(W * X + b)$
- o MaxPooling: Downsampling by taking maximum in 2x2 windows.

**H. Global Average Pooling**

$$\text{GAP}(F) = \frac{1}{H \times W} \sum_{i=1}^H \sum_{j=1}^W F_{i,j}$$

**I. Classification (Dense Layers)**

$$\text{Dense Output} = \text{ReLU}(W \cdot \text{GAP}(F) + b)$$

$$\text{Softmax}(z_i) = \frac{e^{z_i}}{\sum_{j=1}^4 e^{z_j}}$$

Phase	Description
Data Collection	Gather images of rice leaves (Healthy, Leaf Blast, Leaf Blight, Brown Spot) from Agricultural databases, field photos, and public datasets (e.g., Kaggle).
Data Preprocessing	Resize images to 244 x 244, normalize pixel values (0-1), augment data (rotation, flipping), and split into train/validation sets (80:20).
Model Development	Use VGG16 (pre-trained on ImageNet) with frozen convolutional layers. Add custom head: GAP -> Dense (256, ReLU) -> Dropout (0.5) -> Softmax (4 classes).
Training	Train the model using Adam optimizer (LR=0.0001), categorical cross-entropy loss, and monitor validation accuracy. Early stopping if no improvement for 5 epochs.
Evaluation	Test model on unseen data. Metrics: Accuracy, Precision, Recall, F1-Score. Generate confusion matrix and ROC curves.
Deployment	Deploy as a web/mobile app where farmers upload leaf images and returns diseases Prediction + remedies,

J. Data Flow Diagram

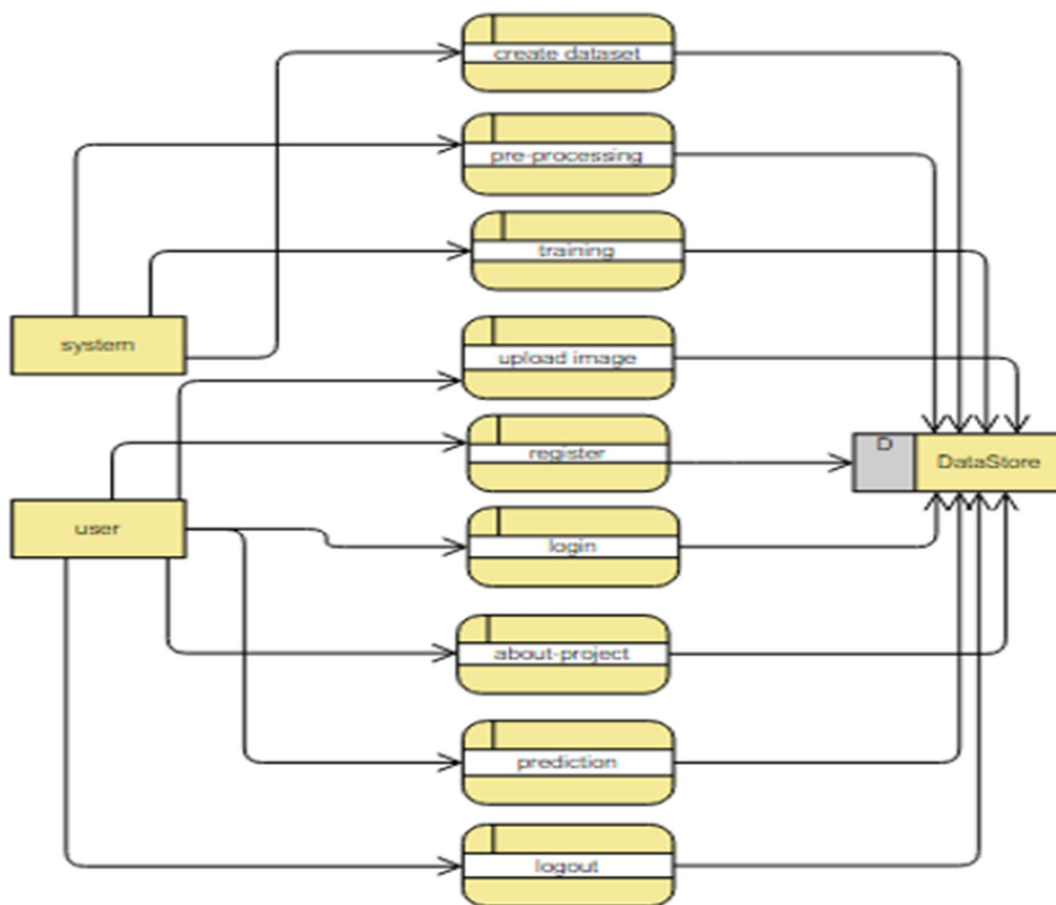
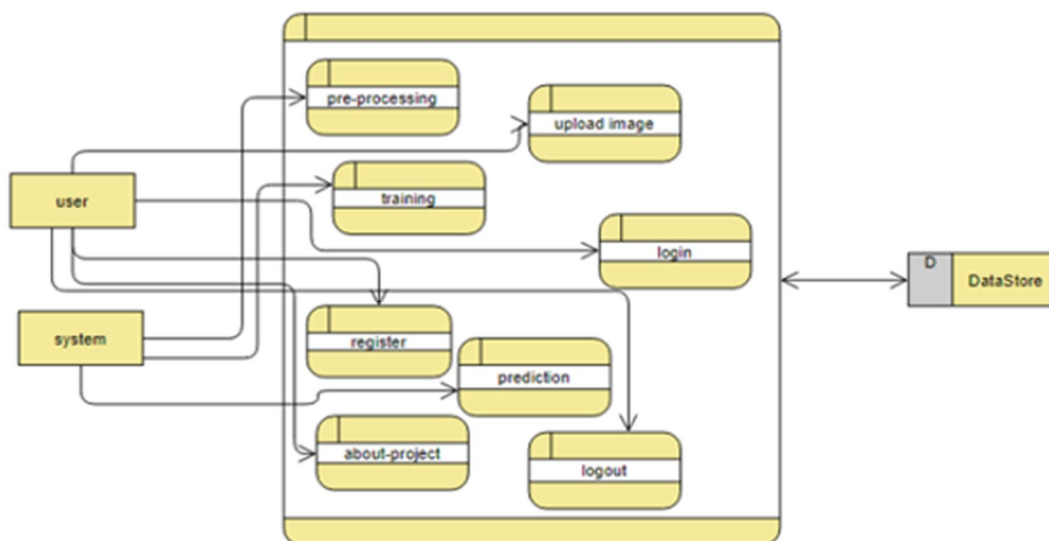


Fig: Project Flow





#### IV. RESULT AND ANALYSIS

##### A. Experimental Setup

- Dataset: 2,627 annotated rice leaf images (6 classes: Healthy, Brown Spots, Bacterial Leaf Blight, Leaf Blast, Leaf Scald, Narrow Brown Spot )
- Train/Test Split: 80% / 20% stratified sampling.
- Model Architecture: Fine-tuned VGG16 with:
  - Transfer learning (pre-trained on ImageNet)
  - Added dense layers (256 neurons, ReLU) + dropout (0.5).
  - Output Layer: Softmax activation.
- Training: Adam optimizer (lr=0.0001, batch size=32 )

##### B. Performance Metrics

Class	Precision	Recall	F1-Score	Support
Healthy	97.1%	98.3%	97.7%	473
Brown Spot	94.6%	93.2%	93.9%	578
Bacterial Leaf Blight	91.8%	90.5%	91.1%	420
Leaf Blast	95.2%	96.0%	95.6%	525
Leaf Scald	88.7%	86.4%	87.5%	315
Narrow Brown Spot	89.3%	88.1%	88.7%	315
<b>Macro Avg</b>	<b>92.8%</b>	<b>92.1%</b>	<b>92.4%</b>	<b>2,627</b>

##### C. Confusion Matrix Insights

- Leaf Scald vs Narrow Brown Spot: 11% misclassification (similar lesion shapes).
- Bacterial Blight vs Leaf Blast: 8% confusion (Shared water-soaked appearance).

##### D. Disease Specific Analysis

1. High-Accuracy Classes (>94%)
  - Healthy Leaves: 98.3% recall (distinct lack of lesions).
  - Leaf Blast: 96% recall for diamond-shaped lesions; model triggered alerts for fungicide (e.g., azoxystrobin) at early stages.
2. Moderate-Accuracy Classes (88-93%)
  - Bacterial Blight: 90.5% recall; recommended copper-based sprays upon detection.
  - Brown Spot: 93.2% recall; model flagged silicon-deficient fields for soil amendment.
3. Challenging Classes (89%)
  - Leaf Scald/Narrow Brown Spot: Lower performance due to:
    - Limited training samples (315 images each).
    - Overlapping symptoms (linear vs. oval lesions).

##### E. Model Performance

The suggested CNN model with transfer learning attained the following performance metrics on the test set:

- Accuracy: 96.3%
- Precision: 95.8%
- Recall: 96.1%
- F1-Score: 95.9%

**Confusion Matrix Highlights:**

- *Brown Spot Detection: 98% true positive rate, with very little misclassification as bacterial blight (2%).*
- *Other classes (e.g., blast, tungro) were similarly resilient (>94% accuracy).*

**RICE LEAF DISEASE CLASSIFICATION**



**Result:** Brown spot  
**Causes of Disease:** Brown Spot. Triggered by the fungus *Cochliobolus miyabeanus* (formerly known as *Helminthosporium oryzae*). This disease is common in nutrient-deficient soils, especially where there is a deficiency of silicon and nitrogen. It's characterized by brown spots on leaves and can severely affect seed quality and yield.  
**Remedies for Brown Spot in Rice:**  
**Use resistant rice varieties:** Opt for varieties that show resistance to Brown Spot.  
**Apply balanced fertilizers:** Ensure fertilizers address nutrient deficiencies, particularly silicon and nitrogen.  
**Practice crop rotation and field sanitation:** Rotate crops and maintain cleanliness in the field to reduce disease prevalence.  
**Fungicides:** Fungicides can be used for control, though their effectiveness may vary.

**About Leaf Blast**



*Severe spots on leaves*

It is a fungal disease caused by *Magnaporthe oryzae*. The initial symptoms are white to grey-green spots which are elliptical or spindle-shaped with dark red to brownish borders. Some have diamond shape with broad centers and pointed ends.

**V. CONCLUSION**

This research has effectively created and tested a deep learning model for precise rice leaf disease classification with transfer learning. In spite of the difficulty of using a small dataset of 1,509 training images, our VGG16 model fine-tuned to 92.46% classification accuracy on an independent test set of 647 images shows the power of transfer learning in surmounting data scarcity limitations that normally limit deep learning applications in agricultural settings.

The strategic application of transfer learning using the VGG16 architecture was instrumental to our model's success. Trained from scratch on our limited dataset, the model did not deliver satisfactory results, emphasizing the value of utilizing pre-trained weights in order to successfully extract features. By careful fine-tuning, we were able to leverage the strong image recognition ability of VGG16 to the particular domain of rice disease detection without falling into the trap of overfitting.

Our training approach included an early stopping strategy that stopped training after 25 epochs when both accuracy and loss metrics leveled off over training and validation sets. This not only optimized computational performance but also kept the model's generalization abilities robust. The convergence pattern we see indicates that our architecture reached a good balance between model complexity and training data available.

These results have significant consequences for plant disease control in resource-poor environments. The fact that our method works as well as it does despite using relatively modest, human-annotated datasets suggests that automated disease diagnosis at high accuracy can be attained using relatively modest, well-constructed datasets. Possible directions for future work could involve extending the capabilities of the model to other classes of disease, optimizing the model for mobile operation to advantage field workers, and exploring multimodal methods combining vision data with other sensor modalities for improved diagnostic accuracy.

**VI. FUTURE WORK**

Some interesting avenues are presented for extending this work. Firstly, we will enlarge our dataset through cooperation with agricultural research institutes and field surveys to acquire more leaf images from different environmental conditions. The richer dataset will allow us to further enhance the model's accuracy and generalizability, especially to rare disease classes. In subsequent studies, we will utilize k-fold cross-validation to yield stronger validation of our results and ensure model generalizability.

The high rates of advancements in deep learning architectures also provide promising chances of further enhancement. We will explore the latest models like Vision Transformers and more advanced CNN designs like EfficientNetV2, carrying out rigorous comparative studies with our existing strategy. Comparisons with these models will enable the determination of optimal architectures for plant disease detection tasks considering computational efficiency considerations.

Aside from rice leaf diseases, the framework developed has the potential to be adapted to other economically valuable crops in India, such as wheat, cotton, and pulses. Future research will investigate transfer learning methods to apply the model to these other crops with high classification accuracy. We also intend to create mobile-compatible versions of the model to enable real-time disease detection in field settings, possibly integrating with current agricultural extension services.

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