

# Revolutionizing Crop Disease Management Fine-Tuned Integrated Convolutional Attention Capsule Autoencoder for Automated Paddy Leaf Disease Identification

**Gangumolu Harsha Vardhan1\* , Vasa Siva Subramanyam<sup>2</sup> , Shaik Jabina Farha<sup>3</sup> , Kalkurthi Joyce Jerlen 4**

*1\* Jr. Assistant Professor, Department of Information Technology, Vishnu Institute of Technology, Bhimavaram, Andhra Pradesh, India. 2,3,4Vishnu Institute of Technology, Bhimavaram, Andhra Pradesh,India.*

*Email: <sup>2</sup> [vsivasub2003@gmail.com](mailto:vsivasub2003@gmail.com)[,](mailto:sivavaraprasad599@gmail.com)  3 [shaikjabinafarha08@gmail.com,](mailto:shaikjabinafarha08@gmail.com,)  4 [Joycejerlen1234@gmail.com](mailto:Joycejerlen1234@gmail.com)  Corresponding Email:1\*[harshavardhan5905@gmail.com](mailto:harshavardhan5905@gmail.com)*

**Received:** 29 March 2024 **Accepted:** 17 June 2024 **Published:** 01 August 2024

*Abstract: Crop diseases are a major threat to food security and agricultural productivity. Early and accurate detection of crop diseases is essential for effective disease management and prevention. However, conventional methods of crop disease identification are timeconsuming, labor-intensive, and require expert knowledge. Therefore, there is a need for developing automated and reliable methods of crop disease identification using advanced technologies such as artificial intelligence (AI). In this paper, we propose a novel AI-based method for automated paddy leaf disease identification using fine-tuned integrated convolutional attention capsule autoencoder (FICACA). FICACA is a deep learning model that combines the advantages of convolutional neural networks (CNNs), attention mechanisms, capsule networks, and autoencoders to extract and encode discriminative features from paddy leaf images. FICACA can identify 10 common paddy leaf diseases with high accuracy and efficiency. We evaluate the performance of FICACA on a large-scale dataset of paddy leaf images collected from different regions and seasons. We compare FICACA with several state-of-the-art methods and demonstrate its superiority in terms of accuracy, robustness, and generalization. We also conduct ablation studies to analyze the contribution of each component of FICACA. Our results show that FICACA can revolutionize crop disease management by providing a fast and accurate solution for paddy leaf disease identification.*

**International Journal of Agriculture and Animal Production ISSN: 2799-0907** Vol: 04, No. 05, Aug-Sept 2024 <http://journal.hmjournals.com/index.php/IJAAP> **DOI:** <https://doi.org/10.55529/ijaap.45.19.27>



#### *Keywords: Convolutional Neural Networks (CNN's), Attention Mechanisms, Capsule Networks, and Autoencoders.*

#### **1. INTRODUCTION**

The global agricultural landscape faces a perpetual challenge in safeguarding food security and ensuring sustainable agricultural productivity. Central to this challenge is the constant threat posed by crop diseases, which can devastate yields and compromise the availability of nutritious food for a growing global population. Paddy rice, a staple crop feeding billions, is no exception to this threat. The timely and precise identification of crop diseases is a pivotal step in mitigating their impact and ensuring food security.

Historically, the identification of paddy leaf diseases has been a labor-intensive and timeconsuming process, reliant on human expertise. Expert agronomists and pathologists often need to visually inspect countless leaves for symptoms, making it impractical for large-scale agricultural operations. Moreover, the accuracy of such identifications is subject to the proficiency of the individuals involved, which introduces an inherent margin of error.

In the quest for more efficient and accurate disease identification, artificial intelligence (AI) has emerged as a promising avenue. Deep learning in particular has shown tremendous promise in image-based disease diagnosis using machine learning approaches. These methods can process vast amounts of visual data rapidly and consistently, reducing the dependence on human expertise.

In this context, our research presents a groundbreaking approach to automated paddy leaf disease identification. We introduce the Fine-Tuned Integrated Convolutional Attention Capsule Autoencoder (FICACA), a deep learning model that harnesses the combined power of convolutional neural networks (CNNs), attention mechanisms, capsule networks, and autoencoders. FICACA is engineered to extract and encode highly discriminative features from paddy leaf images, enabling it to identify common paddy leaf diseases with a high degree of accuracy and efficiency.

This paper explores the development, training, and evaluation of the FICACA model. We delve into the specifics of our methodology, including the dataset used for training and evaluation, the architecture of FICACA, and the techniques employed for fine-tuning. We conduct a thorough performance evaluation, comparing FICACA against several state-of-the-art methods, and assessing its robustness and generalization capabilities across diverse datasets collected from various geographical regions and seasons.

Moreover, this paper presents ablation studies to dissect the contributions of each component within the FICACA framework, shedding light on the importance of the integrated design. The results not only showcase the superior performance of FICACA but also underline its transformative potential in revolutionizing crop disease management.

By providing a fast and accurate solution for paddy leaf disease identification, FICACA offers a means to enhance the efficiency and effectiveness of disease control measures in paddy rice cultivation. This research contributes to the broader effort to ensure food security and sustainable agriculture in the face of mounting global challenges. Furthermore, it underscores the pivotal role of AI and deep learning technologies in addressing critical issues at the intersection of agriculture and technology.

**International Journal of Agriculture and Animal Production ISSN: 2799-0907** Vol: 04, No. 05, Aug-Sept 2024 <http://journal.hmjournals.com/index.php/IJAAP> **DOI:** <https://doi.org/10.55529/ijaap.45.19.27>



## **2. RELATED WORKS**

Since crop diseases have a direct impact on crop output and quality, managing them is an essential part of agriculture. Conventional techniques for diagnosing and treating diseases need a lot of time and work. However, new developments in deep learning and machine learning methods have completely changed the management of crop diseases by making it possible to identify crop illnesses automatically and accurately.

One approach to automated crop disease identification is the use of autoencoder-based cluster ensembles. Geddes et al. [1] introduced an autoencoder-based cluster ensemble architecture for single-cell RNA-seq data processing. The proposed methodology can be applied to crop disease identification even though this work focuses on single-cell RNA-seq data. Clustering techniques have the potential to increase speed by up to 100% on the original datasets. This indicates that the proposed framework might enable a more accurate identification of the illness.

An second tactic is to use deep learning models to identify agricultural illnesses. Hu et al. developed an agricultural IoT system that uses deep learning and IoT technology to accurately identify crop illnesses [2]. They proposed the multidimensional feature compensation residual neural network (MDFC-ResNet) model for more accurate disease detection. Farmers may quickly identify crop issues and receive diagnostic results using this method, which enables timely disease treatment.

The diagnosis of crop diseases has also made extensive use of convolutional neural networks (CNNs). Bharathi and Sonai [3] identified crop leaf diseases using a CNN-based autoencoder. They discovered that the convolution filter size selection can affect how accurately diseases are identified.

Computer vision techniques, including CNNs, have been applied to various agricultural tasks, including crop disease identification. Kamath et al. [4] discussed the use of computer vision applications for crop disease identification and discrimination of crops and weeds. They proposed a multiple classifier system approach for paddy crop and weed discrimination. This highlights the potential of computer vision techniques in automating various agricultural tasks, including disease detection in crops.

Attention mechanisms have also been applied to crop disease identification. Yang et al. [6] studied the use of transfer learning models and attention mechanisms for crop classification identification. They found that transfer learning models can improve the performance of lowperformance classification models. This suggests that attention mechanisms can enhance the accuracy of crop disease identification by focusing on the most important characteristics.

Other deep learning models, including stacked denoising autoencoders, have been applied to the diagnosis of health conditions in a variety of domains in addition to CNNs. A multi-domain indicator-based optimal stacked denoising autoencoder was proposed by Yan et al. [5] for the purpose of identifying the health state of rolling bearings. They were able to identify health issues with great accuracy, suggesting that stacked denoising autoencoders could be useful for precisely identifying diseases in crops.

Identification of crop diseases has also been investigated through collaborative learning networks. A dual-branch collaborative learning network was presented by Zhang et al. [7] for the purpose of identifying crop diseases. They achieved great accuracy in recognizing specific



crop illnesses by embedding migration learning and localization algorithms for crop disease photos. This emphasizes how crucial group learning is to raising disease diagnosis accuracy.

A lightweight CNN model called GrapeNet was presented by Lin et al. [8] to diagnose illnesses of grape leaves. The model showed good recognition accuracy across multiple datasets, indicating the potential of lightweight CNN models for crop disease identification. Gao et al. [8] reported a dual-branch, effective, channel attention-based crop disease detection model that demonstrated good identification accuracy across a range of datasets. These studies show how different CNN architectures and attention mechanisms perform admirably when it comes to accurately diagnosing crop diseases.

Crop pest and disease management is another area where knowledge graphs have been applied. Knowledge graphs for crop pest and disease data management were covered by Liu et al. [9]. In the area of crop diseases and pests, they examined and categorized the principal knowledge graph technology approaches and procedures. Better disease detection and management are made possible by knowledge graphs, which offer an adaptable and effective way to manage crop disease data.

A technique for plant disease identification utilizing explainable 3D deep learning on hyperspectral images is proposed by Nagasubramanian et al. [10]. This approach combines deep learning techniques with hyperspectral imaging to accurately identify plant diseases. The explainable nature of the model allows for better understanding and interpretation of the disease identification process.

The publication by Orchi et al. [11] provides a comprehensive overview of the use of artificial intelligence and the Internet of Things (IoT) for crop disease diagnosis. In order to enable automated disease diagnosis in crops, the survey covers a variety of methods and technologies—such as deep learning and machine learning—that can be connected with Internet of Things devices.

Kamath et al. [12] present a multiple classifier system approach for paddy crop and weed discrimination. This approach combines multiple classifiers to accurately distinguish between paddy crops and weeds. The use of multiple classifiers improves the accuracy of the discrimination process.

Loti et al.'s [13] comprehensive analysis of machine learning and deep learning approaches aims to discover illnesses and pests in Chile. They look into the use of artificial intelligence methods, like convolutional neural networks, to accurately identify pests and diseases in chili plants.

A deep learning model for diagnosing cassava illness based on mobile devices is put forth by Ramcharan et al. [14]. Using photos taken using a mobile smartphone, this model applies deep learning techniques to identify and diagnose illnesses in cassava plants. Disease diagnosis in the field may now be done quickly and conveniently thanks to the mobile-based method.

Liu et al. [15] describe a generative adversarial network (GAN) based data augmentation method for grape leaf disease diagnosis. This method generates synthetic images using GANs that may be added to the training set of models to detect grape leaf disease. The improved data improves the accuracy of disease identification.



Li et al. [16] use convolutional neural networks (CNNs) to recognize crop pests in natural scenes. They show how well CNNs perform the task of precisely identifying and recognizing agricultural pests from photos taken in their natural habitats.

In summary, revolutionizing crop disease management requires the integration of various machine learning and deep learning techniques. Autoencoder-based cluster ensembles, deep learning models such as CNNs and stacked denoising autoencoders, attention mechanisms, collaborative learning networks, and knowledge graphs have all been explored for automated crop disease identification. These techniques have shown promising results in improving the accuracy and efficiency of crop disease management.

## **3. METHODOLOGY**

An explanation of the paddy leaf image dataset: The paddy leaf picture collection utilized in this work was gathered from various paddy production regions and seasons in Malaysia, Thailand, Indonesia, India, and Indonesia. One of the ten most prevalent paddy leaf diseases bacterial blight, brown spot, false smut, leaf blast, leaf scald, neck blast, red stripe, sheath blight, sheath rot, and tungro—is identified on 10,000 photos of paddy leaves in the collection. The JPEG format is used to store the images, which are 256 by 256 pixels in resolution.

Detailed explanation of the Fine-Tuned Integrated Convolutional Attention Capsule Autoencoder (FICACA) model: FICACA is a deep learning model that combines the advantages of convolutional neural networks (CNNs), attention mechanisms, capsule networks, and autoencoders to extract and encode discriminative features from paddy leaf images. FICACA consists of four main components: a feature extractor, an attention module, a capsule encoder, and a capsule decoder. The feature extractor is a CNN that applies multiple convolutional and pooling layers to the input image and produces a feature map. The attention module is a self-attention mechanism that learns to focus on the most relevant regions of the feature map and generates an attention map. The capsule encoder is a capsule network that transforms the attention map into a set of capsules, each representing a high-level feature of the image. The capsule decoder is an autoencoder that reconstructs the input image from the capsules and minimizes the reconstruction error. The output of FICACA is the class label of the input image, which is determined by the length of the capsules.

Training and fine-tuning procedures: Using a split of 80% training and 20% validation sets from the paddy leaf picture dataset, FICACA was trained and adjusted. Pre-training and finetuning were the two phases of the training procedure. Using just the feature extractor and the capsule decoder components, FICACA was trained as an autoencoder during the pre-training phase. Reconstruction error between the input and output images was to be as small as possible. FICACA was trained as a classifier employing all four components during the fine-tuning phase. Reducing the classification loss between the output label and the ground truth label was the goal. To balance the two goals, the reconstruction error was multiplied by the classification loss. .. Grid search and cross-validation methods were used to fine-tune the FICACA hyperparameters. The following were the ideal values: dropout rate  $= 0.2$ , regularization parameter =  $0.01$ ; learning rate =  $0.001$ , batch size = 32, number of epochs = 50, number of capsules  $= 16$ .



Data augmentation techniques applied: Data augmentation techniques were applied to increase the diversity and robustness of the training data. The techniques included random cropping, flipping, rotating, scaling, shifting, shearing, blurring, adding noise, changing brightness, contrast, saturation, and hue. Each image in the training set was randomly augmented with one or more techniques with a probability of 0.5.



Figure 1. The proposed architecture

Three criteria were employed to analyze FICACA on the test set of paddy leaf images in order to measure its performance: accuracy, precision, and recall. The percentage of correctly categorized photos among all photographs is known as accuracy. The percentage of accurately categorized photos among all images anticipated to be positive for a particular class is known as precision. The percentage of correctly categorized images among all images that are genuinely positive for a particular class is known as recall.

#### **4. RESULTS AND DISCUSSION**

In this section, we present the empirical findings and performance metrics obtained through rigorous experimentation and evaluation. The focus of this section is to objectively report the outcomes of our research, shedding light on how the Fine-Tuned Integrated Convolutional Attention Capsule Autoencoder (FICACA) model performs in the task of automated paddy leaf disease identification. We compare FICACA against baseline models and other relevant methods to assess its effectiveness and superiority. The results unveiled in this section provide a comprehensive view of the model's capabilities, its accuracy, precision, recall, F1-score, and overall performance, bringing us closer to understanding its potential for revolutionizing crop disease management. We critically analyze the outcomes of our experiments, examining why the FICACA model performed as it did and what the implications are for the field of paddy leaf



disease identification and crop disease management as a whole. We also explore the contributions of each component within the FICACA framework, as elucidated in our ablation studies. Furthermore, we discuss the practical implications of our work for agriculture, emphasizing the potential for real-world applications and addressing the challenges and limitations encountered during our research.

Table 1 presents the performance metrics of the Fine-Tuned Integrated Convolutional Attention Capsule Autoencoder (FICACA) model compared to baseline models and other methods for paddy leaf disease identification. Accuracy, precision, recall, and F1-score are reported for each method.



Table 1: Performance Metrics of FICACA Model on Paddy Leaf Disease Identification

When it comes to identifying paddy leaf diseases, the Fine-Tuned Integrated Convolutional Attention Capsule Autoencoder (FICACA) performs the best across the board in terms of all relevant criteria. Outperforming traditional Convolutional Neural Networks (CNN), Recurrent Neural Networks (RNN), and Long Short-Term Memory (LSTM) models, FICACA obtains the highest accuracy, precision, recall, and F1-Score. Notably, fewer false positive predictions are implied by FICACA's improved precision, which is important for managing disease. These findings highlight the potential of FICACA to transform crop disease management by offering reliable, efficient, and accurate identification of paddy leaf diseases, ultimately leading to increased agricultural output and food security.

## **5. CONCLUSION**

We have introduced a new method in this work that uses the Fine-Tuned Integrated Convolutional Attention Capsule Autoencoder (FICACA) model to automatically identify paddy leaf diseases. Our thorough testing and assessment has produced results that show FICACA outperforms traditional techniques like Convolutional Neural Networks (CNN), Recurrent Neural Networks (RNN), and Long Short-Term Memory (LSTM) models in key performance metrics, such as accuracy, precision, recall, and F1-Score. FICACA's superior precision and recall rates, in particular, highlight its potential for accurate and comprehensive disease detection—a pivotal factor in crop disease management. These findings underscore the transformative potential of FICACA in revolutionizing agricultural practices by providing a robust and efficient solution for paddy leaf disease identification. As we look toward the future, the integration of advanced technologies like FICACA represents a promising step forward in ensuring global food security and sustainable agricultural productivity in the face of mounting challenges. Further research and practical implementations of FICACA hold the key to more resilient and productive agricultural systems.



In the realm of automated paddy leaf disease identification and agricultural technology, future enhancements hold substantial promise. Fine-tuning and hyperparameter optimization can further elevate the accuracy and efficiency of models like the Fine-Tuned Integrated Convolutional Attention Capsule Autoencoder (FICACA). Exploring transfer learning for expedited training and multi-spectral imaging for a more comprehensive perspective on crop health could enhance disease detection capabilities. Real-time field applications, including integration with drones and smart farming systems, stand to revolutionize on-site disease monitoring. Moreover, ensuring resource-efficient, lightweight model variants would expand the model's reach to resource-constrained agricultural settings. Collaboration with international agricultural institutions, interdisciplinary research with domain experts, and user-friendly interfaces for farmers will further facilitate adoption. Additionally, research into disease forecasting and environmental impact assessment underscores the broader potential of AIbased approaches in sustainable agriculture. These future directions collectively represent a dynamic path forward in addressing the global challenge of crop disease management and enhancing food security.

### **6. REFERENCES**

- 1. T. Geddes, T. Kim, L. Nan, J. Burchfield, J. Yang, D. Taoet al., "Autoencoder-based cluster ensembles for single-cell rna-seq data analysis", BMC Bioinformatics, vol. 20, no. S19, 2019. https://doi.org/10.1186/s12859-019-3179-5
- 2. W. Hu, J. Fan, Y. Du, B. Li, N. Xiong, & E. Bekkering, "Mdfc–resnet: an agricultural iot system to accurately recognize crop diseases", Ieee Access, vol. 8, p. 115287-115298, 2020. https://doi.org/10.1109/access.2020.3001237
- 3. I. Bharathi and V. Sonai, "Image-based crop leaf disease identification using convolution encoder networks",, 2022. https://doi.org/10.5772/intechopen.106989
- 4. R. Kamath, M. Balachandra, & S. Prabhu, "Paddy crop and weed discrimination: a multiple classifier system approach", International Journal of Agronomy, vol. 2020, p. 1- 14, 2020. https://doi.org/10.1155/2020/6474536
- 5. X. Yan, Y. Liu, & M. Jia, "Health condition identification for rolling bearing using a multi-domain indicator-based optimized stacked denoising autoencoder", Structural Health Monitoring, vol. 19, no. 5, p. 1602-1626, 2019. https://doi.org/10.1177/1475921719893594
- 6. G. Yang, Y. He, Y. Yang, & B. Xu, "Fine-grained image classification for crop disease based on attention mechanism", Frontiers in Plant Science, vol. 11, 2020. https://doi.org/10.3389/fpls.2020.600854
- 7. W. Zhang, X. Sun, L. Zhou, X. Xie, W. Zhao, Z. Lianget al., "Dual-branch collaborative learning network for crop disease identification", Frontiers in Plant Science, vol. 14, 2023. https://doi.org/10.3389/fpls.2023.1117478
- 8. J. Lin, X. Chen, R. Pan, T. Cao, J. Cai, Y. Chenet al., "Grapenet: a lightweight convolutional neural network model for identification of grape leaf diseases", Agriculture, vol. 12, no. 6, p. 887, 2022. https://doi.org/10.3390/agriculture12060887



- 9. X. Liu, X. Bai, L. Wang, R. Bingyuan, S. Lu, & L. Li, "Review and trend analysis of knowledge graphs for crop pest and diseases", Ieee Access, vol. 7, p. 62251-62264, 2019. https://doi.org/10.1109/access.2019.2915987
- 10. K. Nagasubramanian, S. Jones, A. Singh, S. Sarkar, & B. Ganapathysubramanian, "Plant disease identification using explainable 3d deep learning on hyperspectral images", Plant Methods, vol. 15, no. 1, 2019. https://doi.org/10.1186/s13007-019-0479-8
- 11. H. Orchi, M. Sadik, & M. Khaldoun, "On using artificial intelligence and the internet of things for crop disease detection: a contemporary survey", Agriculture, vol. 12, no. 1, p. 9, 2021. https://doi.org/10.3390/agriculture12010009
- 12. R. Kamath, M. Balachandra, & S. Prabhu, "Paddy crop and weed discrimination: a multiple classifier system approach", International Journal of Agronomy, vol. 2020, p. 1- 14, 2020. https://doi.org/10.1155/2020/6474536
- 13. N. Loti, M. Noor, & S. Chang, "Integrated analysis of machine learning and deep learning in chili pest and disease identification", Journal of the Science of Food and Agriculture, vol. 101, no. 9, p. 3582-3594, 2020. https://doi.org/10.1002/jsfa.10987
- 14. A. Ramcharan, P. McCloskey, K. Baranowski, N. Mbilinyi, L. Mrisho, M. Ndalahwaet al., "A mobile-based deep learning model for cassava disease diagnosis", Frontiers in Plant Science, vol. 10, 2019. https://doi.org/10.3389/fpls.2019.00272
- 15. B. Liu, C. Tan, S. Li, J. He, & H. Wang, "A data augmentation method based on generative adversarial networks for grape leaf disease identification", IEEE Access, vol. 8, p. 102188-102198, 2020. https://doi.org/10.1109/access.2020.2998839
- 16. Y. Li, H. Wang, L. Dang, A. Sadeghi-Niaraki, & H. Moon, "Crop pest recognition in natural scenes using convolutional neural networks", Computers and Electronics in Agriculture, vol. 169, p. 105174, 2020. https://doi.org/10.1016/j.compag.2019.105174