

Computers, Materials & Continua DOI: 10.32604/cmc.2022.024618 Article

Artificial Intelligence-Based Fusion Model for Paddy Leaf Disease Detection and Classification

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Received: 25 October 2021; Accepted: 05 January 2022

Abstract: In agriculture, rice plant disease diagnosis has become a challenging issue, and early identification of this disease can avoid huge loss incurred from less crop productivity. Some of the recently-developed computer vision and Deep Learning (DL) approaches can be commonly employed in designing effective models for rice plant disease detection and classification processes. With this motivation, the current research work devises an Efficient Deep Learning based Fusion Model for Rice Plant Disease (EDLFM-RPD) detection and classification. The aim of the proposed EDLFM-RPD technique is to detect and classify different kinds of rice plant diseases in a proficient manner. In addition, EDLFM-RPD technique involves median filtering-based preprocessing and K-means segmentation to determine the infected portions. The study also used a fusion of handcrafted Gray Level Co-occurrence Matrix (GLCM) and Inception-based deep features to derive the features. Finally, Salp Swarm Optimization with Fuzzy Support Vector Machine (FSVM) model is utilized for classification. In order to validate the enhanced outcomes of EDLFM-RPD technique, a series of simulations was conducted. The results were assessed under different measures. The obtained values infer the improved performance of EDLFM-RPD technique over recent approaches and achieved a maximum accuracy of 96.170%.

Keywords: Rice plant disease; classification model; artificial intelligence; deep learning; fusion model; parameter optimization



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1 Introduction

Early detection of pests and plant diseases is one of the challenging problems in the domain of agriculture. In India, rice is accountable for 70% of overall crop production and 93% overall production in terms of grain yield. For this reason, early detection of diseases in paddy plants is of national importance to ensure the productivity of staple in India [1]. Identification and classification of pests and plant diseases remain the main challenge faced in the field of agriculture. Insect damages the crops in almost every stage which in turn affects the yield of crops. Classification of insects that damages the crops is a tedious process, due to high number of connections and complex formation in the emergence of distinct species. However, it is crucial to recognize, categorize and reduce the infection of insects that cause damage to crops, mainly by selecting effective pesticide and biological systems to avoid less yield and heavy damage to the crops [2,3].

In general, few factors that include weather conditions i.e., temperature/moisture, abnormal fertilization unfavourable conditions such as stress, and imbalanced soil nutrient [4] affect the crop yield. In addition to these, if the crops are prematurely harvested and placed in heap, gets damaged by natural phenomena such as cyclone/strong wind, because of collision of panicle and attack of pests at the time of grain formation, these conditions might result in the formation of discoloured grains. Deep Learning (DL) techniques have triggered large expectations in image classification. These methods function on the basis of feature learning from labelled training datasets. Further, these methods have been utilized earlier in recognizing the diseases that affect apple, tea, grapes, tomato, pears, and peaches. Generally, the methods utilize leaves' images to recognize the diseases. In this background, images are utilized from a homogeneous background [5] for this study. Moreover, the data is accessible from different sources on internet.

Every disease has their own special feature. A disease can be identified based on its impact upon the host i.e., plant, in terms of colour, shape, and size of disease symptoms. Some plant diseases induce the plant to exhibit similar colours, but distinct shapes; while some have distinct colours, but similar shapes. Occasionally, farmers get confused owing to these differences and fail to take appropriate action for pesticide selection [6]. One of the ways to avoid crop loss that occurs due to disease infection is by capturing the images of infected leaves, identifying the data about the disease and taking precautionary measures or corrections measures to overcome the disease. As an automated solution to these problems, cameras can be installed across the fields at specific distance for periodic capturing of the images. This image can be transmitted to the main system for disease analysis; the system can identify the disease, provide data about the selection of pesticides and measures to control the disease. These systems, based on their training knowledge, automatically recognize the diseases that occurred earlier. In this background, the aim of this study is to demonstrate how Machine Learning (ML) methods can be made use of in other fields like agriculture [7]. One of the crucial segment in deploying ML fields is the choice of better features. Only, disease-based features can be extracted after the segmentation of non-disease and disease parts. So, it is crucial that the main component of this study is to extract leaf parts from disease portion (most plants have green colour leaves), which is a critical phase in determining the quality of feature.

In order to attain Artificial Intelligence (AI)-enabled accurate and rapid disease recognition, Wang et al. [8] presented the Attention-based Depthwise Separable Neural Network with Bayesian Optimization (ADSNN-BO) algorithm-based augmented attention mechanism and MobileNet structure. Furthermore, Bayesian optimization algorithm was also employed for fine tuning the hyperparameter of the system. Cross-validated classification experiment was carried out, according to public rice disease datasets with four classes in total. The stimulation result demonstrated that the proposed architecture attained 94.65% test accuracy. In Chen et al. [9], MobileNet-V2, pretrained on ImageNet, was selected as the backbone network to improve the learning capacity for minute lesion feature. Further, the study also added attention mechanism to learn the significance of spatial points and interchannel relationships for input feature. Meanwhile, loss function got improved and Transfer Learning (TL) method was implemented twice for training the model.

In Zhang et al. [10], a group of automated diagnosis and detection approaches, integrated with spectral and image analysis, was presented to detect different diseases that are broadly deleterious. The method also checked the quality of rice and yields. This research presented a stepwise methodology to detect rice sheath blight. Temniranrat et al. [11] proposed an object refinement process and detection model training to enhance the performance of prior studies on rice leaf disease recognition. This method depends upon examining the model prediction outcomes and can be used recurrently to improve the quality of dataset in training the following model. Deployment framework was chosen from the elected performances in prior studies i.e., You Only Look Once (YOLO) v3 and is trained by the refined trained dataset.

Jiang et al. [12] focused on three types of rice leaves' disease and two types of wheat leaf disease. The researchers gathered a total of 40 images of all the leaf diseases and improved them. This study focused on improving the Visual Geometry Group (VGG-16) method depending on the concept of multitask learning. It utilized pretraining method on ImageNET for alternate learning and TL method. In Krishnamoorthy et al. [13], InceptionResNetV2, a kind of Convolutional Neural Network (CNN) was used by TL method to recognize the diseases in rice leaf images. The parameter of the presented method enhanced the classification task and attained a better performance.

Bhoi et al. [14], proposed an Internet of Things (IoT)-enabled Unmanned Aerial Vehicles (UAVs)enabled rice pest detection method with Imagga cloud to recognize the pest in rice, at the time of crop production. This IoT-enabled UAV model focused on Python programming and AI model to send the rice pest images to cloud and provide the pesticide data. It decides the pesticide after determining the confidence value with tags. Tiwari et al. [15] presented a DL-based method for plant diseases' classification and detection from leaf images, captured in different resolutions. A dense CNN framework undergone training on huge plant leaf image datasets from many nations and was used in this study.

1.1 Objective of the Paper

The objective of the proposed technique is to detect and classify different kinds of rice plant diseases proficiently, thereby avoiding the loss of crop productivity.

1.2 Contribution of the Paper

The current research paper devises an Efficient Deep Learning-based Fusion Approach for Rice Plant Disease (EDLFM-RPD) diagnosis. In addition, EDLFM-RPD technique involves Median Filtering (MF)-based preprocessing and K-means segmentation to determine the infected portions. Further, a fusion of handcrafted Graylevel Co-occurrence Matrix (GLCM) and Inception-based deep feature was conducted to derive the features. Finally, Fuzzy Support Vector Machine (FSVM) model was utilized for classification. In order to validate the enhanced outcomes of the proposed EDLFM-RPD technique, a series of simulations was conducted and the results were reviewed under different aspects.

1.3 Organization of the Paper

The upcoming portions of the study are planned as follows. Section 2 elaborates the proposed EDLFM-RPD technique and a detailed discussion of results is given in Section 3. At last, the concluding remarks are drawn in Section 4.

2 The Proposed Model

In this study, a new EDLFM-RPD technique is presented to proficiently detect and classify different kinds of rice plant diseases, thereby avoiding the loss in crop productivity. The proposed EDLFM-RPD technique encompasses preprocessing, K-means segmentation, fusion-based feature extraction, and FSVM-based classification. A comprehensive work of every module is discussed in succeeding sections.

2.1 Image Preprocessing

MF is a non-linear procedure that is helpful in decreasing impulsive or salt-and-pepper noise. It is also helpful in maintaining the edge from images, but decreases the arbitrary noise. Impulsive noise occurs due to arbitrary bit error under transmission channel [16]. MF smoothens the images and so it is helpful in decreasing the noise. Unlike low pass filter, MF preserves the discontinuity from a step purpose and smoothens some pixels whose value differs considerably, from its surrounding, without affecting another pixel.

2.2 K-Means Segmentation

Clustering is a technique to divide a group of data under particular sets of groups. One of the most famous clustering techniques is k-means cluster. K-means cluster splits a group of data into k number of data collected for the study.

It categorizes the provided group of data under k amount of disjoint clusters. There are two distinct stages present in K-means technique. The primary stage computes k centroid whereas secondary stage gets all the points to cluster i.e., adjacent centroid in respective data points. Several techniques are used to define the distance of nearby centroids whereas the most utilized technique is Euclidean distance. When the grouping is completed, it recomputes a novel centroid of all the clusters. Based on this centroids, a novel Euclidean distance is computed amongst all the centers and all data points. Then, the point is allocated from the cluster with minimal Euclidean distance. All the separated clusters are determined as their member objects and within their centroids. The centroid to all the cluster points to the sum of distance in every object from that cluster gets minimized [17]. Therefore, K-means is an iterative technique that minimizes the sum of distances in all objects to their cluster centroid on every cluster.

Assume an image with resolution $x \times y$ and image with a cluster as to k amount of cluster. Assume p(x, y) be the input pixels with cluster and c_k be the cluster center. K-means cluster technique is given herewith.

- 1. Initializing the amount of cluster k and centre.
- 2. To all the pixels of an image, compute the Euclidean distance *d*, among the center and all pixels of image based on the relation given herewith.

$$d = ||p(x, y) - c_k||$$
(1)

3. Allocate every pixel to adjacent centre, depending upon distance d.

4. Afterward, each pixel is allocated and then the novel place of centre is recomputed based on the relation given herewith.

$$c_k = \frac{1}{k} \sum_{y \in c_k} \sum_{x \in c_k} p(x, y)$$
⁽²⁾

- 5. Repeat the procedure till it fulfills the tolerance or error value.
- 6. Reshape the cluster pixel as to image.

When primary centroid is arbitrarily selected, it attains distinct outcomes to different primary centers. Thus, the primary center is carefully selected so that it can obtain the desired segmentation. Some other terms are also required in addition to computational difficulty while the development of K-means clustering is prominent here. It depends upon the count of data elements, count of clusters, and count of iterations.

2.3 Fusion Based Feature Extraction

In this phase, GLCM and Inception features are fused together to derive a set of feature vectors. In general, GLCM is meant to be a statistical method with spatial connection among the pixels. Here, five varied texture features are explained in *co*-occurrence matrix function as given herewith.

$$Energy = \sum_{a} \sum_{b} C^{2}_{\theta,d}(a, b)$$
(3)

$$Variance = \frac{1}{N \times M} \sum_{b} \sum_{a} (C_{\theta, a}(a, b) - m)^2$$
(4)

$$Entropy = -\sum_{a} \sum_{b} C_{\theta,d}(a, b) \log_2(C_{\theta,d}(a, b))$$
(5)

$$Contrast = \sum_{a} \sum_{b} (a-b)^2 C_{\theta,d}(a, b)$$
(6)

$$Homogeneity = \sum_{a} \sum_{b} \frac{C_{\theta,d}(a,b)}{1+|a-b|}$$
(7)

where *m* indicates the mean value of *co*-occurrence matrix, $C_{\theta,d}$.

CNN has a total of five layers such as input, convolution, pooled, Fully Connected (FC), and output. GoogLeNet network is intended to make CNN, an utilizable element by Google. It implements the Inception network approach, which restricts the amount of network attributes and improves the depth of systems. So, it can be widely utilized in image classification processes. Fig. 1 shows the sample of usual CNN.

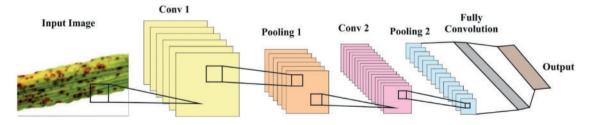


Figure 1: Structure of CNN

Convolutional layer obtains various layers in Neural Network (NN) while not every pixel is connected to future layer with weight and bias. But the entire image gets separated to a small area, while weight and bias are utilized. Here, Pooling layer is executed to decrease the spatial dimensions of image and parameter counts and minimize the processes. It carries out a set function to input due to lack of some parameters. Distinct kinds of pooling layers are available such as average, stochastic, and max pooling. At this point, the flattened outcome of final pooling layer is fed as input to FC layer. It performs the role of CNN in every neuron of the present layer which is connected to present layer. Therefore, the number of parameters should be the maximum than convolutional layer. This FC layer is linked with resultant layer i.e., classifier.

Activation function

The diverse activation function is implemented on various frameworks of CNN. Non-linear activation function is demonstrated with better results than previous sigmoid or tangent functions. Non-linear functions are executed to enhance the training speed. So, distinct activation functions are implemented and Rectified Linear Unit (ReLU) is a remarkable technique compared to other such techniques.

CNN learning technique depends on vector calculus and chain rule. Let z is scalar (for instance, $z \in R$) and $y \in R^{H}$ is a vector [18]. Therefore, if z has a function of y, then the partial derivative of z, with respect to y, is the vector which is defined as follows

$$\begin{pmatrix} \partial z \\ \partial y \end{pmatrix}_i = \frac{\partial z}{\partial y_i}$$
 (8)

Particularly, $\begin{pmatrix} \partial z \\ \partial y \end{pmatrix}$ refers to the vector that compromise of identical size as y, and their *i*-th element

is $\begin{pmatrix} \partial z \\ \partial y \end{pmatrix}_i$. Besides, it can be stated that $\begin{pmatrix} \partial z \\ \partial y^T \end{pmatrix} = \begin{pmatrix} \partial z \\ \partial y \end{pmatrix}^T$. Also, let $x \in R^W$ is another vector, and y represents the function of x. Then, a partial derivative of y with respect to x is defined as follows

$$\begin{pmatrix} \partial z \\ \partial y^T \end{pmatrix}_{ij} = \frac{\partial y_i}{\partial x_i}.$$
(9)

During fractional derivative of $H \times W$ matrix, the $\frac{\partial y_i}{\partial x_i}$ can be retrieved at juncture of i^{th} row and j^{th} column. It is easy and obvious that z has a function of x from the chain-like arguments. In addition, the function maps x to y, and other function maps y to z. The chain rule is employed here for computing as given herewith.

$$\begin{pmatrix} \partial z \\ \partial x^T \end{pmatrix}, as \begin{pmatrix} \partial z \\ \partial x^T \end{pmatrix} = \begin{pmatrix} \partial z \\ \partial y^T \end{pmatrix} \begin{pmatrix} \partial z \\ \partial x^T \end{pmatrix}$$
(10)

The cost or loss function is employed to measure the alteration amongst the forecast of CNN x^L and the objective $t, x^1 \rightarrow w^1, x^2 \rightarrow \ldots, x^L \rightarrow w^L = z$, employing a simplistic loss function $z = ||t - x^L||^2$. The predictive result is realized as $argmax_i x_i^L$. In convolutional technique, it is demonstrated as follows

$$y_{i^{l+1}j^{l+1},d} = \sum_{i=0}^{H} \sum_{j=0}^{W} \sum_{d=0}^{D} f_{ij,d} \times x_{i^{l+1}+,i,j^{l+1}+j,d}^{L}$$
(11)

Filter f is sized at $(H \times W \times D^{l})$, thus the convolution has a spatial size of $(H^{l} - H + 1) \times (W^{l} - W + 1)$ with D slices and represents that $y(x^{l+1})$ in $R^{H^{l+1} \times W^{l+1} \times D^{l+1}}$, $H^{l+1} = H^{l} - H + 1$, $W^{l+1} = W^{l} - W + 1$, $D^{l+1} = D$.

The probability of every label $k \in \{1, ..., K\}$, implemented for training the sample, is computed as $P(k|x) = \frac{\exp(z_k)}{\sum_{i}^{K} \exp(z_i)}$, where z signifies the non-normalization log possibilities. The ground truth, shared on labels q(k|x), remains the normalization, because $\sum_{k} q(k|x) = 1$. In this technique, the loss is incurred by Cross Entropy (CE) as determined herewith.

$$l = \sum_{k=1}^{K} \log(p(k))q(k)$$
(12)

Cross Entropy (CE) loss can be differentiated with respect to logit z_k . It can be employed for gradient training of deep techniques since the gradients are easier procedures $\frac{\partial l}{\partial z_k} = p(k) - q(k)$ and is bound between -1 and 1. Usually, when CE obtains minimization, it signifies that the log possibility of accurate label gets maximized. Inception V3 is considered as shared on labels, independent of trained samples u(k), with smooth parameter ϵ , for training sample. The label shared $q(k|x) = \delta_{k,y}$ is simply returned as follows.

$$q^{'(k|x)} = (1 - \epsilon)\delta_{k,x} + \frac{\epsilon}{K}.$$
(13)

Then, it can be understood that CE is calculated as given herewith.

$$H(q',p) = -\sum_{k=1}^{K} \log(p(k))q'^{(k)} = (1-\epsilon)H(q',p) + \epsilon H(u,p).$$
(14)

Therefore, label-smoothing regularization is similar to be applied as a single CE loss H(q, p) with a couple of losses H(q, p) and H(u, p), while second loss penalizes the difference of the predicted label, shared p in previous u with comparative weight $\frac{\epsilon}{(1-\epsilon)}$.

The main objective of GoogLeNet network is Inception network framework while GoogLeNet technique is called Inception networks [19]. It encompasses maximal GoogLeNet version and is classified as distinct versions such as Inception v1, Inception v2, Inception v3, Inception v4, and Inception-ResNet. Therefore, Inception usually has three sizes of convolutional and maximal pooling layers. The outcomes of the network from preceding layer are determined as channel has gathered the information, when implementation of convolutional task and non-linear fusion is performed. Also, the express purpose of this network is to have appropriate and improved distinct scales and remove the over-fitting issues. Inception v3 signifies the network framework utilized by Keras. For sample, a 3 * 3 convolutional is separated as 3 * 1 and 1 * 3 convolutional layers. By utilizing this partition technique, the amount of attributes are restricted; so, training speed of the network gets triggered, while removing the spatial feature from efficient model. Concurrently, Inception v3 optimizes the Inception model with the help of three distinct-sized grids namely, 35 * 35, 17 * 17, and 8 * 8.

Data fusion is applied in different computer vision and ML fields. Feature fusion is an important process, which incorporates the maximum number of feature vectors. The proposed technique is based on fusion of features using entropy. Furthermore, the features are attained and integrated with individual vectors. It is expressed as follows.

$$f_{Inception_v3\times m} = \{Inception_v3_{1\times 1}, Inception_v3_{1\times 2}, Inception_v3_{1\times 3}, \dots, Inception_v3_{1\times n}\}$$
(15)

$$f_{GLCM\times p} = \{GLCM_{1\times 1}, GLCM_{1\times 2}, GLCM_{1\times 3}, \dots, GLCM_{1\times n}\}$$
(16)

Next, feature extraction is integrated as an individual vector.

Fused (feature vector)_{1×q} =
$$\sum_{i=1}^{2} \{ fInceptionV3_{1×m}, fGLCM_{1×p} \}$$
 (17)

whereas f indicates a fused vector. Entropy is performed on features vector to elect the features as follows.

$$B_{He} = -NHe_b \sum_{i=1}^{n} p(f_i) \tag{18}$$

$$F_{select} = B_{He}(\max(f_i, 1186)) \tag{19}$$

In Eqs. (18) and (19), p represents the feature likelihood and He determines entropy. At last, the elected feature is provided to classification model to detect and classify the plant diseases.

2.4 Image Classification

Finally, FSVM model is applied to determine the classes that exist in rice plant image. In traditional Support Vector Machine (SVM), each data point is considered to have equally importance and is allocated a similar penal variable in its objective function. To address this issue, FSVM system is proposed. Fuzzy memberships are proposed to each sample point; therefore, distinct instance points can create different contributions to the creation of decision surfaces. Fig. 2 depicts the hyperplane of SVM model.

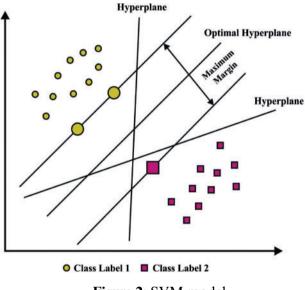


Figure 2: SVM model

Consider training instance as follows

$$S = \{(x_1, y_i, s_i), i = 1, \dots, N\},$$
(20)

Whereas $x_1 \in \mathbb{R}^n$ signifies *n*-dimensional instance point, $y_i \in \{-1, +1\}$ represents class label, and $s_i (i = 1, ..., N)$ implies fuzzy membership that fulfills $\sigma \le s_i \le 1$ with adequately smaller constant

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 $\sigma > 0$. Re denotes the quadratic optimization problem for classification and is expressed as follows.

$$\frac{\min_{w,s,\xi} \frac{1}{2} w^{T} w + C \sum_{i=1}^{l} s_{i} \xi_{i}}{s.t. y_{i}(w^{T} x_{l} + b) \ge 1 - \xi_{i}, \xi_{i} \ge 0, i = 1, \dots, l,}$$
(21)

While *w* signifies the standard vector for splitting hyperplane, *b* means bias term, and *C* implies the variable that should be described before controlling the trade-offs between cost of misclassification error and classification margin [20]. Subsequently, s_i represents the attitude of equivalent point x_1 to single class; slack parameter ξ_i signifies the measure of error, then $s_i\xi_i$ denotes the deliberated measures of error with distinct weights. It is stated that larger the s_i is, more prominent, the equivalent point is processed; lesser the s_i is, lesser prominent the equivalent point is processed; hence, dissimilar input points might produce several contributions to learn the decision surface. Later, FSVM detects stronger hyperplanes by growing the margin and allowing few misclassifications of lesser significant points. To address Finite State Machine (FSM) optimal problem, (21) is transformed to two problems given herewith by offering Lagrangian multiplier, α_i :

$$\max \sum_{i=1}^{N} \alpha_{i} - \frac{1}{2} \sum_{i=1}^{N} \sum_{j=1}^{N} \alpha_{i} \alpha_{j} y_{i} y_{j} x_{i} x_{j}$$

$$s.t. \sum_{i=1}^{N} y_{i} \alpha_{i} = 0, 0 \le \alpha_{i} \le s_{i} C, i = 1, \dots, N.$$
(22)

Like regular SVM, the aforementioned representation has a small variation, i.e., upper bound of α_i . By resolving these two problems for optimal α_i , w, and b are recovered in the same way as in standard SVM.

3 Experimental Validation

This section investigates the performance of the proposed EDLFM-RPD technique in terms of rice plant disease detection and classification. The model was validated against a benchmark dataset [21] that comprises of 40 images under Bacterial Leaf Blight (BLB), Brown Spot (BS), and Leaf Smut (LS).

Fig. 3 demonstrates the confusion matrices generated by the proposed EDLFM-RPD technique on classification of rice plant disease images under five runs. With run-1, EDLFM-RPD technique classified 35 images under class 0, 36 images under class 1, and 36 images under class 2. Similarly, with run-3, EDLFM-RPD technique classified 40 images under class 0, 34 images under class 1, and 37 images under class 2. Lastly, with run-5, EDLFM-RPD technique classified 39 images under class 0, 36 images under class 2.

Tab. 1 shows the results from overall classification analysis accomplished by EDLFM-RPD technique. The table values denote that the proposed EDLFM-RPD technique identified the class labels effectually under distinct test runs. For instance, with run-1, the EDLFM-RPD technique achieved *sens_y*, *spec_y*, *prec_n*, *accu_y*, and *F_{score}* values such as 0.8750, 0.9733, 0.9459, 0.9391, and 0.9091 respectively. In addition, with run-2, EDLFM-RPD technique attained *sens_y*, *spec_y*, *prec_n*, *accu_y*, and *F_{score}* values namely, 0.9209, 0.9609, 0.9212, 0.9478, and 0.9201 respectively. Moreover, with run-3, the proposed EDLFM-RPD technique gained *sens_y*, *spec_y*, *prec_n*, *accu_y*, and *F_{score}* values such as 0.9642, 0.9826, 0.9653, 0.9768, and 0.9644 respectively. Furthermore, with run-4, EDLFM-RPD technique produced *sens_y*, *spec_y*, *prec_n*, *accu_y*, and *F_{score}* values being 0.8750, 0.9733, 0.9459, 0.9391, and 0.9091

respectively. Moreover, with run-4, the EDLFM-RPD technique accomplished $sens_y$, $spec_y$, $prec_n$, $accu_y$, and F_{score} values such as 0.9478, 0.9740, 0.9478, 0.9652, and 0.9477 respectively. At last, with run-5, the EDLFM-RPD technique produced $sens_y$, $spec_y$, $prec_n$, $accu_y$, and F_{score} values namely, 0.9476, 0.9736, 0.9505, 0.9652, and 0.9481 respectively.

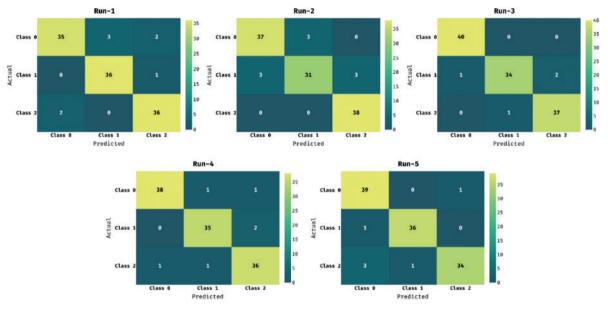


Figure 3: Confusion matrix of EDLFM-RPD model

Classes	$Sens_y$	$Spec_{y}$	$Prec_n$	Accu _y	F_{score}
Run-1					
BLB	0.8750	0.9733	0.9459	0.9391	0.9091
BS	0.9730	0.9615	0.9231	0.9652	0.9474
LS	0.9474	0.9610	0.9231	0.9565	0.9351
Avg.	0.9318	0.9653	0.9307	0.9536	0.9305
Run-2					
BLB	0.9250	0.9600	0.9250	0.9478	0.9250
BS	0.8378	0.9615	0.9118	0.9217	0.8732
LS	1.0000	0.9610	0.9268	0.9739	0.9620
Avg.	0.9209	0.9609	0.9212	0.9478	0.9201
Run-3					
BLB	1.000	0.9867	0.9756	0.9913	0.9877
BS	0.9189	0.9872	0.9714	0.9652	0.9444
LS	0.9737	0.9740	0.9487	0.9739	0.9610
Avg.	0.9642	0.9826	0.9653	0.9768	0.9644

(Continued)

Table 1: Continued						
Classes	<i>Sens</i> _y	Spec _y	$Prec_n$	Accu _y	F_{score}	
Run-4						
BLB	0.9500	0.9867	0.9744	0.9739	0.9620	
BS	0.9459	0.9744	0.9459	0.9652	0.9459	
LS	0.9474	0.9610	0.9231	0.9565	0.9351	
Avg.	0.9478	0.9740	0.9478	0.9652	0.9477	
Run-5						
BLB	0.9750	0.9467	0.9070	0.9565	0.9398	
BS	0.9730	0.9872	0.9730	0.9826	0.9730	
LS	0.8947	0.9870	0.9714	0.9565	0.9315	
Avg.	0.9476	0.9736	0.9505	0.9652	0.9481	

Fig. 4 illustrates the results of accuracy graph analysis attained by EDLFM-RPD technique on test dataset. The figure depicts that the proposed EDLFM-RPD technique improved both training and validation accuracies.

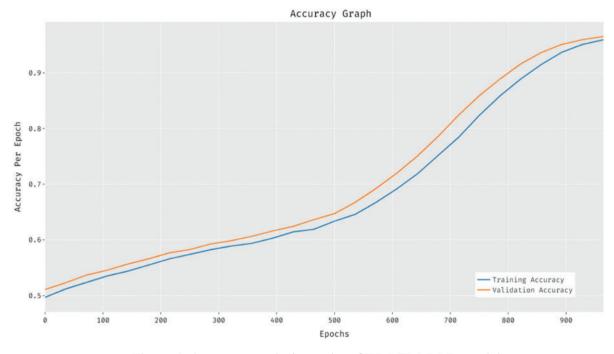


Figure 4: Accuracy analysis results of EDLFM-RPD model

Fig. 5 demonstrates the loss graph analysis results accomplished by EDLFM-RPD technique on test dataset. The figure portrays that the proposed EDLFM-RPD technique reduced both training and validation losses.

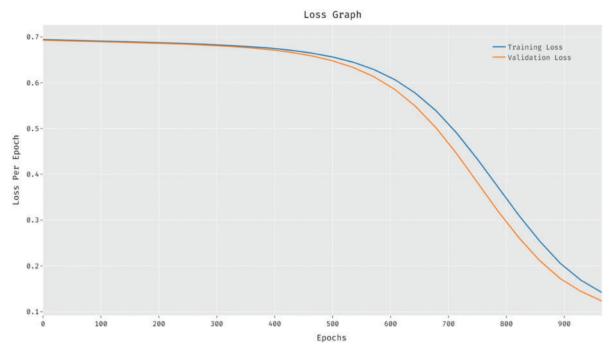


Figure 5: Loss analysis results of EDLFM-RPD model

Fig. 6 reports the Receiver Operating Characteristic (ROC) analysis results achieved by the proposed EDLFM-RPD technique on test dataset. The figure implies that EDLFM-RPD technique gained a higher ROC of 99.2300.

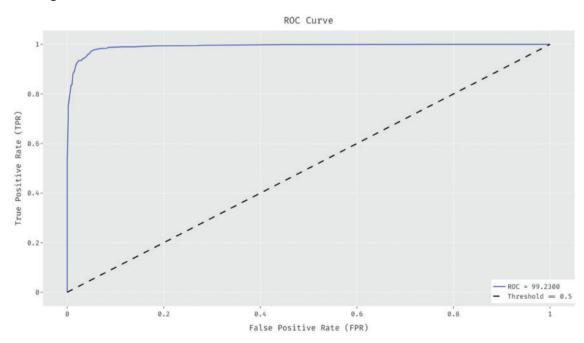


Figure 6: ROC analysis results of EDLFM-RPD model

Tab. 2 offers the results for overall classification analysis accomplished by EDLFM-RPD technique under five different runs. The figure depicts that the proposed EDLFM-RPD technique achieved increased average $sens_y$, $spec_y$, $prec_n$, $accu_y$, and F_{score} values such as 0.9425, 0.9713, 0.9431, 0.9617, and 0.9422 respectively.

No. of runs	Sens _y	$Spec_{y}$	<i>Prec</i> _n	Accu _y	F_{score}
Run-1	0.9318	0.9653	0.9307	0.9536	0.9305
Run-2	0.9209	0.9609	0.9212	0.9478	0.9201
Run-3	0.9642	0.9826	0.9653	0.9768	0.9644
Run-4	0.9478	0.974	0.9478	0.9652	0.9477
Run-5	0.9476	0.9736	0.9505	0.9652	0.9481
Average	0.9425	0.9713	0.9431	0.9617	0.9422

 Table 2: Classification analysis results of EDLFM-RPD model under different runs

Finally, a comprehensive comparative classification analysis was conducted upon EDLFM-RPD technique under different measures and the results are shown in Tab. 3. The proposed model was compared with CNN-based inception with ResNset v2 model and Optimal Weighted Extreme Learning Machine (CNNIR-OWELM), VGG-16, Deep NN, deep autoencoder (AE), Artificial neural networks (ANN), Conv. NN, K-nearest neighbor (KNN), Scale-invariant feature transform with support vector machine (SIFT-SVM), and SIFT-KNN.

Methods	Sens _y	$Spec_{y}$	<i>Prec</i> _n	$Accu_y$	F_{score}
Proposed method	94.250	97.130	94.310	96.170	94.220
CNNIR-OWELM	90.500	96.100	92.100	94.200	91.300
VGG-16	89.800	93.200	90.200	92.900	89.900
Deep NN	73.500	89.400	74.900	90.000	81.500
Deep AE	68.000	87.200	67.600	86.000	77.000
ANN	63.300	81.600	60.900	80.000	68.300
Conv. NN	94.000	94.000	94.000	93.800	94.000
KNN	65.000	78.000	72.000	70.000	65.000
SIFT-SVM	86.700	83.200	86.700	91.100	86.700
SIFT-KNN	90.000	86.400	90.600	93.300	90.100

Table 3: Comparison analysis results of EDLFM-RPD technique against recent approaches

Fig. 7 shows the results of $sens_y$, $spec_y$, and $prec_n$ analysis achieved by EDLFM-RPD technique against recent approaches. The figure portrays that Deep AE, ANN, and KNN models obtained ineffective outcomes with least values in terms of $sens_y$, $spec_y$, and $prec_n$. Along with that, Deep NN, VGG-16, and SIFT-SVM models reached moderate $sens_y$, $spec_y$, and $prec_n$ values. Moreover, CNNIR-OWELM and SIFT-KNN techniques accomplished reasonable $sens_y$, $spec_y$, and $prec_n$ values. However, the proposed EDLFM-RPD technique achieved a maximum performance with $sens_y$, $spec_y$, and $prec_n$ values.

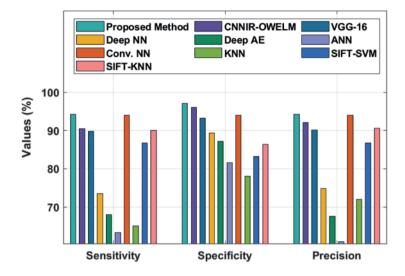


Figure 7: Comparative analysis of EDLFM-RPD model with varying measures

Fig. 8 shows the results of accuracy analysis attained by EDLFM-RPD technique against other techniques. The results demonstrate that Deep AE, ANN, and KNN techniques resulted in low *accu_y* values namely, 86%, 80%, and 70% respectively. In line with these, VGG-16, Deep NN, Conv. NN, SIFT-KNN, and SIFT-SVM techniques too attained moderately closer *accu_y* values such as 92.90%, 90%, 93.80%, 93.30%, and 91.10% respectively. Moreover, CNNIR-OWELM technique accomplished a reasonable *accu_y* of 94.20% while the proposed EDLFM-RPD technique reached the highest *accu_y* of 96.17%.

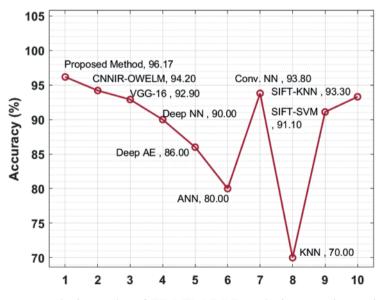


Figure 8: Accuracy analysis results of EDLFM-RPD technique against existing approaches

At last, a brief F_{score} analysis was conducted between EDLFM-RPD technique and other techniques and the results are shown in Fig. 9. The results confirm that Deep AE, ANN, and KNN

techniques gained low F_{score} values such as 77%, 68.30%, and 65% respectively. Besides, VGG-16, Deep NN, Conv. NN, SIFT-KNN, and SIFT-SVM techniques too accomplished somewhat improved F_{score} values namely, 89.90%, 81.50%, 94%, 90.10%, and 86.70% respectively. Though CNNIR-OWELM technique accomplished a reasonable F_{score} of 91.30%, the proposed EDLFM-RPD technique reached the highest F_{score} of 94.22%. Based on the results and discussion, it is apparent that the proposed EDLFM-RPD approach can acts as a vital method for rice plant disease detection and classification.

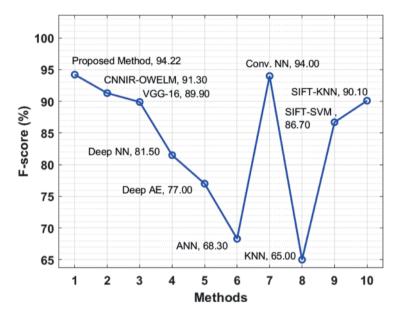


Figure 9: F-score analysis results of EDLFM-RPD technique against existing approaches

4 Conclusion

In this study, a novel EDLFM-RPD technique is derived to detect and classify different kinds of rice plant diseases proficiently, thereby avoiding the loss in crop productivity. The proposed EDLFM-RPD technique encompasses preprocessing, K-means segmentation, fusion-based feature extraction, and FSVM-based classification. In order to validate the improved outcomes of EDLFM-RPD approach, a series of simulations was conducted and the results were inspected under different dimensions. The experimental values establish the improved performance of EDLFM-RPD technique against recent approaches. Therefore, the proposed EDLFM-RPD technique can be employed as a proficient tool for rice plant disease detection and classification. In future, EDLFM-RPD technique can be realized in IoT and cloud environments to assist farmers in real-time setting.

Acknowledgement: The authors would like to acknowledge the support of Prince Sultan University for paying the Article Processing Charges (APC) of this publication.

Funding Statement: The authors extend their appreciation to the Deanship of Scientific Research at King Khalid University for funding this work under Grant Number (RGP.1/127/42). www.kku.edu. sa.

Conflicts of Interest: The authors declare that they have no conflicts of interest to report regarding the present study.

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