RDA- CNN: Enhanced Super Resolution Method for Rice Plant Disease Classification

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Abstract: In the field of agriculture, the development of an early warning diagnostic system is essential for timely detection and accurate diagnosis of diseases in rice plants. This research focuses on identifying the plant diseases and detecting them promptly through the advancements in the field of computer vision. The images obtained from in-field farms are typically with less visual information. However, there is a significant impact on the classification accuracy in the disease diagnosis due to the lack of high-resolution crop images. We propose a novel Reconstructed Disease Aware–Convolutional Neural Network (RDA-CNN), inspired by recent CNN architectures, that integrates image super resolution and classification into a single model for rice plant disease classification. This network takes low-resolution images of rice crops as input and employs the super resolution layers to transform low-resolution images to super-resolution images to recover appearance such as spots, rot, and lesion on different parts of the rice plants. Extensive experimental results indicated that the proposed RDA-CNN method performs well under diverse aspects generating visually pleasing images and outperforms better than other conventional Super Resolution (SR) methods. Furthermore, these super-resolution images are subsequently passed through deep classification layers for disease classification. The results demonstrate that the RDA- CNN significantly boosts the classification performance by nearly 4–6% compared with the baseline architectures.

Keywords: Super-resolution; deep learning; interpolation; convolutional neural network; agriculture; rice plant disease; classification

1 Introduction

Rice (Oryza sativa L.) being the primary source of nutrition and the major consumed food crop, demands constant increase with respect to the equivalent growth in population. Depending upon the environmental factors and variety, rice plants take around 3–6 months duration to obtain their yield. The growth stages of rice plant such as vegetative, reproductive, ripening is monitored and production may be affected by diseases which result in yield loss. These diseases are caused by various pathogens like bacteria, fungus or viruses and can affect different parts of crops at any of the stages [1]. Here the study considered most commonly affected different diseases of rice in Asian countries listed in Tab. 1.

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Identifying and diagnosing rice plant diseases and pests is one of the major challenges in the agriculture sector. Timely detection is generally carried out through visual inspection by experts. Thus, it is often time-consuming, detection is not accurate and may lead to misdiagnosis, overuse of pesticides etc. Hence, there is a need for an automatic early disease diagnostics system for classification. This system helps the researcher to monitor the plant growth and also detect the disease symptoms at earlier stages through deep learning and computer vision algorithms.

In recent years, many researchers showed optimistic results in improving the accuracy of plant disease classification using deep learning techniques [2,3]. Similarly, the images acquired in real-time are typically unclear and thus reduces the accuracy of the image classification algorithms. In order to enhance the low-image quality (blur, noise, poor camera focus), SR algorithm needs to be incorporated to reconstruct the quality of input images. Although super-resolution is concerned, the early research on the contribution of deep learning is Super-Resolution Convolutional Neural Network (SRCNN). SRCNN, which aims to restore missing information from low-resolution images and acts as a mapping function between input low resolution ($I_{LR}$) and reconstructed high-resolution images ($I_{SR}$) [4]. Deep learning-based image SR techniques are in increased demand in the field of agricultural imaging which enhances the overall performance of disease recognition and classification through computer vision techniques.

The scarcity of images with high resolution becomes challenging, recovering texture details from input images, poor diagnostic performance are few limitations identified in the related works. In order to overcome this, SR methods generate higher image resolution achieves better disease classification. The proposed RDA-CNN method uses two main architectures to recover the high resolution of input images by integration of CNN-based super resolution layers with the deep classification layers.

During this exploration, the key contributions of this paper can be summarized as follows;

1. A Novel Reconstructed Disease Aware–Convolutional Neural Network (RDA-CNN) is the combination of super resolution layers and classification layers.
2. RDA-CNN uses convolutional layers to improve fine-grained low-resolution images with different scaling factors for image classification.
3. In addition, experimental results of verified RDA-CNN outperform state-of-the-art deep learning CNN architectures are used in terms of visual image quality and classification accuracy on a publicly available rice disease image dataset.

Table 1: Types of diseases that affect rice crop

| Disease name | Disease stages | Cause category | Infected area         |
|--------------|----------------|----------------|-----------------------|
| Rice blast   | Nursery        | Fungus         | Leaf, collar, neck, node |
| Bacterial leaf blight | Nursery        | Bacteria       | Leaves                |
| Rice tungro disease | Nursery     | Virus          | Leaves                |
| Bacterial leaf streak | Main field  | Bacteria       | Leaves                |
| Brown spot   | Main field     | Fungus         | Grains and leaves     |
| False smut   | Main field     | Fungus         | Grains                |
| Sheath rot   | Main field     | Fungus         | Stem                  |
| Rice grassy stunt virus | Main field | Virus          | Leaves                |
2 Related Works

Recently, many researchers have extensively explored deep learning techniques for super resolution imaging for computer vision tasks that could become very popular and have also achieved significant performance improvements compared to traditional super resolution methods [4,5]. In this section, a short overview of deep neural networks in the context of super-resolution tasks and image recognition is discussed. In many application domains, real-time input images were provided with a lack of resolution used for experimental study. The use of these images seems to be the primary reason for a relatively low diagnostic performance and therefore, low image quality has been highlighted as an essential aspect that affects classification accuracy [6,7]. The objective is to improve low resolution image quality and a crucial step in the emerging research domain such as security surveillance, agriculture, medical diagnostics, remote sensing, ultrasound imaging, astronomical observation and biometric recognition and so on [8]. Image SR techniques aim to enhance the resolution of images from either single or multiple images. In recent times, SR based on a single image as Single Image Super Resolution (SISR) has shown significant performance using learning and reconstructed methods [9,10]. Dong et.al uses pre- upsampling SR techniques and proposed Convolutional Neural Network based Super Resolution techniques referred to as SRCNN model, which further shows significant prediction performance on mapping of low resolution and reconstructed high-resolution images. Furthermore, use of larger filter size in CNN layers are better and the performance of SRCNN is compared with conventional methods.

SRCNN was employed first in the agricultural sector for automated disease diagnosis of tomato leaf images [10]. Here, SRCNN is designed to enhance the spatial resolution of an image where, interpolated low resolution images are taken as input. it showed promising results when compared to conventional interpolated methods such as bilinear, nearest neighborhood, bicubic, cubic, and lanczos4. In addition, the accuracy of disease classification using AlexNet architecture increased by 20% between low and super resolution images whereas the accuracy by 90%. Few other research papers focused on classical work to explore further details on SRCNN and have shown improved performance in significant research fields such as plant disease diagnosis [11–13], cytogenetics [14], blood cell count [15], CT scan [16], and vehicle detection [17], agricultural pest detection [18,19], etc.

SRGAN is the Generative Adversarial Network (GAN) based super resolution method which combines conventional SRCNN with deep learning-based image classification proposed to generate super resolved images and perceptual loss function [20]. Although GAN showed better ability in super resolution tasks for disease classification in plants. Recently, Enhanced Super-resolution GAN (ESRGAN) was used to resolve low resolution crop image and generates its corresponding SR images based on GAN method. Due to inadequate data, transfer learning is applied to fine-tune the ESRGAN model to avoid underfitting problems. The experiments using this model recovered more realistic image crop information and improved the classification accuracy of tomato plants. Images obtained from real environment are typically not clear and of low quality which results in significant impact on disease identification and classification accuracy. To overcome this issue, Dual Attention and Topology Fusion GAN (DATFGAN) was proposed to convert into high resolution images which improve classification accuracy nearly by 2–3% using different transfer learning models. Similarly, SR method for diagnosing cucumber leaf disease named Leaf Artifact-Suppression Super Resolution (LASSR) was proposed to generate images with high quality which achieved 2% better classification performance compared with ESRGAN [21].

In recent years, plant disease recognition and crop stress using deep learning models are the major challenges for the agricultural industry. A lot of intensive studies on factors affecting crop production and identification of specific plant diseases were developed using the current advancement in the field of deep learning and computer vision. More recently, 60% of the researchers uses conventional deep convolutional neural networks for plant disease classification model that works efficiently on different crops such as wheat, paddy, banana, mango, tomato, maize, soybean sugarcane and cassava plant leaves [22–24].
Different kinds of CNN-based models are being used in the field of disease recognition in plants. Earlier, few researchers explored the CNN model as a feature extractor in which features are extracted in the last convolution layer. Initially, CNN with SoftMax and CNN with Support Vector Machines (SVM) results were compared in terms of efficiency to investigate the CNN deep features [25]. Popularly, CNN-based transfer learning models such as VGG16, ResNet50, InceptionV3, InceptionResNetV2, Xception have been used to recognize crop diseases and pests, and the performance of these models was compared in terms of fine tuning, transfer learning, and training from scratch of rice crop diseases [26]. In addition, VGG16 performed high accuracy for the classification of various diseases in different datasets [27,28]. The author proposed novel a DENS-INCE which is the combination of pre-trained DenseNet and Inception network for disease identification in rice plants and achieved better performance when compared to other pre-trained models [29].

According to these research studies, deep learning-based classification methods are useful for analyzing plant disease and successful results have been obtained. The challenges based on the limitation of existing methods for disease classification are as follows: i) lack of availability of high-resolution image data, ii) recovery of texture details from low-quality images, iii) images collected from a controlled environment, iv) data scarcity, and v) complex background in real conditions. However, the images retrieved from the farms (both healthy and diseased images) are usually unclear which causes a significant impact on plant disease image classification. Therefore, it is crucial to improve the disease classification accuracy using SR methods. The proposed RDA-CNN extremely useful in real-time applications especially in the agricultural sector help the researchers to make a plant disease diagnosis and classification at an early stage.

3 Proposed Methodology

The proposed network named RDA-CNN integrates the super resolution layers and classification layers are the major contribution addressed in this paper as shown in the Fig. 1. It includes three distinctive phases of techniques: i) initial feature extraction, ii) super resolution layers and iii) classification layers for disease classification. Image quality distortion uses a conventional image scaling algorithm for resizing the input image dataset. Finally, low resolution image can be reconstructed to super resolution image and performance of the different image classifier models is improved. The overview of the proposed methodology is discussed in this section.

Figure 1: Architecture of reconstructed disease aware–convolutional neural network (RDA-CNN)

3.1 Initial Feature Extraction

Consider the input dataset with high resolution images \( I_{HR} = \{I_1, I_2, I_3, \ldots, I_N\} \) of each class in a rice disease samples. Generally, the low-resolution image \( I_{LR} \) is obtained by \( I_{HR} \) using Eq. (1).
\[ I_{LR} = F(I_{HR}; \sigma) \]  

where \( F \) denotes a degradation function and \( \sigma \) degradation parameter as scaling factor (\( r \)) or a noise. In general, the process of image degradation is unknown and only \( I_{LR} \) images are provided. Several works implement downsampling operation with scaling factor (\( r \)) as a degradation process, where \( r \) is always greater than 1 and as discussed in Eqs. (2) and (3).

\[ F(I_{HR}; \sigma) = (I_{HR}) \downarrow r, \ \{r\} \subset \sigma \]  

The \( I_{LR} \) is the low-resolution representation of \( I_{HR} \) image, which would be retrieved by resizing the downsampled images using interpolation methods.

Algorithm 1: DSRCNN

1. **Input:** High resolution images \( \{I_{HR}\}_i, i:1\rightarrow N \), scaling factor \( r \) re \{2, 3, 4\}
2. **Initialize:** \( Z_0 \leftarrow \{I_{HR}\}_i \)
3. \( I_{LR}^{\text{NN}} \leftarrow \text{nearestneighbor}(Z_0) \)
4. **for** \( i \leftarrow 1 \) to 2
5. \( X_i = W_i \ast Z_{i-1} \) // \( \ast \) - Convolution operation
6. \( Z_i = \max (X_i, 0) \)
7. **if** \( i=2 \)
8. \( I_{LR}^{\text{bil}} = \text{bicubic}(Z_2) \)
9. \( I_{LR} = [I_{LR}^{\text{NN}}, I_{LR}^{\text{bil}}] \) // [] - concatenation operation
10. **End**
11. **Output:** \( I_{LR} \)

**Training Phase**

12. **Input:** \( \{\{I_{LR}\}_i\}_i \), i:1\rightarrow N, scaling factor \( r \) re \{2, 3, 4\}
13. **for** \( i \leftarrow 1 \) to \( N \) // \( N \)- no of training image data
14. **for** \( j \leftarrow 1 \) to 3 // three – layered CNN
15. Compute \( L_i \) (\( I_{LR} \)) based on Eqs. (4)-(6)
16. **End**
17. Calculate \( \text{MSE} \leftarrow \sum \text{func} \{\|I_{LR}\|_i - \|I_{SR}\|_i\} \)
18. Calculate \( \text{PSNR} \leftarrow 10 \log_{10} (\text{MAX}^2/\text{MSE}) \)
19. Calculate \( \text{SSIM} \leftarrow \text{stat_func} (\mu, \sigma) \{\|I_{LR}\|_i - \|I_{SR}\|_i\} \)
20. **end**
21. **Output:** Model parameter \( \Theta \leftarrow \{W_1, W_2, W_3, B_1, B_2, B_3\} \)

**Testing Phase**

22. **Input:** Low resolution images(\( \{I_{LR}\}_i \), i:1\rightarrow N, scaling factor \( r \) re \{2, 3, 4\},
Model parameter \( \Theta \leftarrow \{W_1, W_2, W_3, B_1, B_2, B_3\}\)
23. **for** \( j \leftarrow 1 \) to 3
24. Compute \( L_i \) (\( I_{LR} \)) based on Eqs. (4)-(6)
3.2 Super-Resolution Layers

The ultimate goal of this work is to reconstruct a super-resolution image \( I_{SR} \) from the low-resolution image \( I_{LR} \) and learn a function mapping between \( I_{LR} \) and \( I_{HR} \). To recover quality of the image, Dong et.al considered a Convolutional Neural Network (CNN) and proposed a model named SRCNN for SISR. The proposed Deformed SRCNN (DSRCNN) uses a three-layer convolution network, and the upsampling layer is applied before the convolutional layers to upscale the \( I_{LR} \) image to the \( I_{SR} \) size. The architecture of the DSRCNN is shown in Fig. 1.

3.3 Upsampling Layer

Given an input color image \( I_{LR} \) is the low-resolution version of \( I_{HR} \) which includes tensor size as width \((w)\), height \((h)\) and RGB channel \((c)\). In the upsampling layer, image size is scaled to \( rh \times rw \times c \), where \( r \) is the scaling factor, \( r \in \{2, 3, 4\} \) and \( c \) is the color channel. This can be expressed as follows in Eq. (3).

\[
r = \frac{w^{HR}}{w^{LR}} = \frac{h^{HR}}{h^{LR}}
\]

Generally, upsampling methods such as interpolation (bilinear, bicubic, nearest neighborhood, cubic, and lanczos4), deconvolution, and unpooling are used in computer vision is employed with CNN. The original SRCNN uses bicubic interpolation for upscaling the image to its desired size. In the proposed work, upsampling layers uses bilinear and nearest neighbor interpolation methods to achieve better results for the rice disease dataset. The output of bilinear interpolated images is concatenated with the nearest neighbor interpolated input before being fed to the convolutional block.

3.4 Convolutional Block

This section explores the concepts of convolutional neural networks that are applied to super resolution problem. Consider the input image in which the RGB color space is converted to \( YCbCr \) and the DSRCNN network trains the luminescence \((Y)\) channel exclusively. With an input low-resolution image, final super resolved image is the output resulting from three layered convolutional network includes the phases of operation such as feature extraction, non-linear mapping and reconstruction as follows.

3.4.1 Feature Extraction

The upscale low-resolution image as \( I_{LR} \) will be used as an input for first layer Convolution Neural Network. Consider the input image \( I_{LR} \) has size \( w, h \) represents the width and height respectively. \( I_{LR} \) is convoluted with kernels of the size \( f_1 \times f_1 \) and the depth of this layer is \( n_1 \), i.e., the number of filters in this layer represents \( n_1 \). Formally, the first layer is given in Eq. (4) as follows.

\[
L_1(I_{LR}) = \max(0, W_1 * I_{LR} + B_1)
\]

where \( W_1 \) denotes kernels in filters, \( B_1 \) is the bias vector with \( n_1 \) dimensions and ‘\(*\)’ represents the convolution operation. The \( W_1 \) is applied to each filter and size of the \( W_1 \) is \( c \times f_1 \times f_1 \times n_1 \), where \( c \) is the
channel \((c = 1)\), \(f_1 \times f_1\) is the spatial size of filter \((f_1 = 9)\), \(n_1\) is the number of filter \((n_1 = 128)\) and \(B_1\) is \(n_1\). A Rectified Linear Unit (ReLU, \(\max (0, x)\)) used as an activation function on each filter response.

### 3.4.2 Non-Linear Mapping

The first layer in the network is initially scaled down and then scaled up to the input RGB image using bilinear interpolation technique. This layer extracts \(n_1\) dimensional features maps \((n_1 = 128)\) using the \(9 \times 9\) kernel function. Similarly, second layer maps \(n_1\) to \(n_2\) dimensional features maps with \(n_2\) filters and kernel sizes such as \(f_2 \times f_2\). The second layer operation is specified in Eq. (5) as follows

\[
L_2(I_{LR}) = \max(0 W_2 * L_1(I_{LR}) + B_2)
\]  

(5)

\(W_2\) and \(B_2\) are the kernels size and bias vector respectively in the second layer, \(L_1(I_{LR})\) is the feature maps from the first layer. The size of the \(W_2\) is \(n_1 \times f_2 \times f_2\) as \(f_2 = 3\) and bias vector \(n_2\) where \(n_2 = 64\). The output of second layer is considered as \(n_2\)-dimensional vectors and thus represents the high-resolution patch that to be used for image reconstruction.

### 3.4.3 Reconstruction

The third layer in the network aggregates the high-resolution patches from the previous output that results in generating a final super resoluted image. The operation of this layer is expressed in Eq. (6) as

\[
I_{SR} = (W_3 * L_2(I_{LR}) + B_3)
\]  

(6)

Here \(W_3\) represents kernel size in filter \(n_3\) and size is given as \(n_2 \times f_3 \times f_3 \times c\) where \(f_3 = 5\) and \(B_3\) is \(c\)-dimensional vector size and \(c = 1\).

### 3.5 Image Quality Assessment

Mean Square Error, Peak Signal-to-Noise Ratio and Structural Similarity can be used to evaluate the quality of an image. The performance of the super-resolution method is evaluated between high-resolution images \((I_{HR})\) and low-resolution images \((I_{LR})\) and super-resolution images \((I_{SR})\) for quantitative evaluation.

#### 3.5.1 Peak Signal-to-Noise Ratio (PSNR)

Given the high-resolution image \(I_{HR}\) and a reconstructed super resolution image \(I_{SR}\) with \(N\) pixels, the PSNR between \(I_{HR}\) and \(I_{SR}\) is defined as maximum pixel value of gray scale \((\text{MAX}-\text{ranges to 255})\) and Mean Squared Error between images. As shown below in Eqs. (7) and (8), MSE and PSNR are derived between the reference and reconstructed image of \(N\) pixels and PSNR computation is associated with the pixel-level MSE.

\[
MSE = \frac{1}{N^2} \sum_{w=1}^{N} \sum_{h=1}^{N} (I_{SR}(w,h) - I_{HR}(w,h))^2
\]  

(7)

\[
PSNR = 10 \log_{10} \frac{\text{MAX}^2}{MSE}
\]  

(8)

where \(I_{SR}\), \(I_{HR}\) and \(N\) are super resolution image, high resolution image and image size respectively. Here the PSNR is measured in terms of Decibel (dB).

#### 3.5.2 Structural Similarity (SSIM)

For the assessment of structural similarity between images, the structural similarity index (SSIM) is proposed based on independent comparisons of an image with \(N\) pixels in terms of luminance \((\mu)\) as mean, contrast \((\sigma)\) as standard deviation of image intensity and structures are calculated using Eq. (8) as follows.
SSIM = \frac{(2 \mu_{ISR} \mu_{IHR} + C_1)(2 \sigma_{ISR} \sigma_{IHR} + C_2)}{\left(\mu_{ISR}^2 + \mu_{IHR}^2 + C_1\right)\left(\sigma_{ISR}^2 + \sigma_{IHR}^2 + C_2\right)}

where \( \mu_{ISR} \) and \( \mu_{IHR} \) represent the mean, \( \mu_{ISR}^2 \) and \( \mu_{IHR}^2 \) represents variance, \( \sigma_{ISR} \) and \( \sigma_{IHR} \) is the covariance of two images. Assume \( C_1 \) and \( C_2 \) are the constants where \( C_1 = (0.01 \cdot \text{MAX})^2 \) and \( C_2 = (0.03 \cdot \text{MAX})^2 \) are used to determine the SSIM values. Lower MSE values and higher PSNR and SSIM values represent better image quality. If the SSIM value is closer to 1, results in efficient image reconstruction. When these images are identical, the SSIM value is one.

### 3.6 Classification Layers

The third phase of the classification layers is combinational of convolutional and fully connected layers with high quality images from the previous super-resolution layers. For image classification, the number of CNN frameworks have been proposed. Furthermore, the five popular Convolutional Neural Networks are used for the rice crop disease classification. All CNN's architecture with different depths typically consists of Convolutional, ReLU, pooling and fully-connected layers.

### 3.7 Network Training

Consider the input dataset with high-resolution, low-resolution, and super-resolution images \( \{(I_{HR})_i, (I_{LR})_i, (I_{SR})_i\} \), where \( i = 1 \) to \( N \) images and class labels \( (C_1 – C_7) \). In these Rice Plant Disease (RPD) classification, input dataset is divided into training and testing data with ratio of 0.8 and 0.2 respectively. The architecture of Reformed CNN (ReCNN) for RPD classification has convolutional, pooling and fully connected layers connected with ReLU activation function as shown in Fig. 2. The performance of the proposed model is compared with traditional deep learning models such as AlexNet, Inception_ModelV3, ResNet-50 and VGGNet. The number of epochs used to train ReCNN is 150, Adam optimizer with learning rate of \( 10^{-3} \) and remaining parameters are set to default values. The loss function of this model uses Cross entropy. The algorithm for the classification layers is given as below.

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**Algorithm 2: ReCNN**

1. **Input**: \( I \subset \{(I_{HR})_i, (I_{LR})_i, (I_{SR})_i\} \), \( i:1\rightarrow N \), Classes \( \leftarrow [\text{‘Tungro’, ‘Bacterialblight’, ‘Blast’, ‘Brownspot’, ‘Leaf smut’, ‘healthy’, ‘Hispa’}] \) // \( N \)- Number of images
2. **Initialize**: width \( \leftarrow \) 75, height \( \leftarrow \) 75, classes \( \leftarrow \) 7, EPOCHS \( \leftarrow \) 150
3. **for** \( j\rightarrow (0, \text{EPOCHS}) \)
4. x\_train\_data, y\_test\_data \( \leftarrow \) train\_test\_split (dataset, test\_size = 0.2)
5. **for** c\_data \( \leftarrow \) x\_train\_data
6. X \( \leftarrow \) c\_data [0], Y \( \leftarrow \) c\_data [1]
7. pred\_results \( \leftarrow \) ReCNN\_MODEL(X)
8. cost \( \leftarrow \) CALCULATE\_CROSS\_ENTROPY\_MEAN (pred\_results, Y)
9. optimizer \( \leftarrow \) ADAM\_OPTIMIZER (lr = \( 10^{-3} \)). MIN (cost) // lr- learning rate
10. **End**
11. **end**
12. **Output**: Performance Analysis report

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4 Experimental Results and Analysis

4.1 Data Acquisition

Rice Crop disease images were acquired from sources extracted from papers authored by [29,30]. This work encompasses seven classes of rice disease images in the dataset which includes i) Blast (*Pyricularia grisea* (*P. oryzae*)), ii) Brown Spot (*Helminthosporium oryzae*), iii) Bacterial Leaf Blight (*Xanthomonas oryzae* *pv. oryzae*), iv) Sheath Blight (*Rhizoctonia Solani*), v) Rice Tungro virus (*RTSV, RTBV*), vi) Hispa, vii) False Smut (*Ustilaginoidea virens*) and viii) Healthy. The number of images of each disease class of different dataset is shown in Tab. 2.

![Proposed ReCNN network architecture](image)

**Figure 2:** Proposed ReCNN network architecture

| Class label | Rice diseases     | Total no of image samples |
|-------------|-------------------|---------------------------|
| C1          | Leaf blast        | 2219                      |
| C2          | Brown Spot        | 2163                      |
| C3          | Tungro            | 1308                      |
| C4          | Bacterial leaf blight | 1624                  |
| C5          | Hispa             | 1488                      |
| C6          | Healthy           | 490                       |
| C7          | Leaf smut         | 565                       |
| Total       |                   | 9857                      |

The images of rice plant affected with above mentioned diseases are shown in Fig. 3.

4.2 Results and Discussions

The performance of proposed RDA-CNN super-resolution over conventional methods and thus results in improving the spatial resolution of both healthy and diseased images of rice plants are discussed in this section. However, the experimental results show better performance of generated super-resolution images and also analyze the importance of SR images for rice disease classification.
4.2.1 Evaluation of DSRCNN

Experimental analysis of the proposed super-resolution method for improving the resolution of low-quality input images is considered and image quality assessment measures like PSNR and SSIM are analyzed. The experiment conducted on different classes of rice plant diseases clearly depicts that DSRCNN successfully reconstructed the low-resolution images. On the other hand, reconstructed SR images helps the researchers to recognize detailed characteristics of a diseased images like lesion, rust and spots on different parts of the plant. The images of rice plant dataset are split into two sets using 80% for training and 20% for testing as a split ratio. During this training phase, low resolution images are generated by downsampling method. Then these interpolated images are up sampled by using nearest neighbor with two skip connections and bilinear interpolation with varying upscaling factor 2–4 respectively. In training phase, proposed DSRCNN model is trained for 200 epochs with batch size of 128. Next, in testing phase DSRCNN model is to produce the super-resolution images of rice plant images. In addition, the performance of super resolution network architecture is examined based on the factors like number of filters, layer depth and filter sizes. Generally, adding more filter would improve the performance of the network. For the experimental analysis, the network model is trained with varying number of filters are shown in Tab. 3 and results indicates that increasing the width would result in better performance. The architecture of DSRCNN is modelled with three layers depth, filter numbers as $n_1 = 128$, $n_2 = 64$ and filter size (9-3-5) to greater results.

The performance of DSRCNN model is evaluated between high-resolution, low resolution and super resolution images of different diseases using PSNR and SSIM. These measures are analyzed and also compared with conventional interpolation and SR methods as shown in the Tab. 4. Although the results of DSRCNN deep learning-based method outperforms for all the classes (C1–C7) with scaling factor (2–4). The generated SR images of the DSRCNN model resulted in higher PSNR and SSIM than low resolution images.

The average results of performance metrics like MSE, PSNR and SSIM for the evaluation of image quality assessment with different scaling factor (2X, 3X and 4X) are shown in the Fig. 4.
In Fig. 4, the graph provides insights average values of MSE, PSNR and SSIM for different SR algorithms over seven different classes of RPD with 2X scaling shows better results compared to other scaling factors. These results indicate that DSRCNN outperformed all other methods in terms of PSNR and SSIM at all magnification scales. In DSRCNN, PSNR were higher by at least 1.5 to 2 dB with conventional methods and 0.5 to 1 dB with baseline and transfer learning based SRCNN. Similarly, SSIM also increased by average of 0.01 to 0.025. Although the proposed super-resolution method performed very well, less than 2–3% of the test images resulted in lower performance measure than the other low-resolution images.

### 4.2.2 Evaluation on Disease Classification

In disease classification experiment, the input image samples (high, low and super resolution image) are chosen randomly with split into train and test sets with a split ratio of 80% and 20%. The proposed model RDA-CNN with ReCNN shows the overall better accuracy of rice disease classification using low, high and super resolution images with different scaling factor is discussed in. Furthermore, classification accuracy on super resolution images is quite higher compared with low resolution images for all scaling factor. These

| Scaling factor | Class label | Bilinear PSNR/ SSIM | Nearest neighbor PSNR/SSIM | SRCNN$_b$ PSNR/SSIM | SRCNN$_f$ PSNR/SSIM | DSRCNN (proposed) PSNR/ SSIM |
|---------------|-------------|----------------------|-----------------------------|----------------------|----------------------|-----------------------------|
| 2X            | C1          | 36.278/0.942         | 36.127/0.932                | 37.425/0.920         | 37.434/0.921         | 38.434/0.959                |
|               | C2          | 38.171/0.938         | 38.182/0.918                | 39.283/0.941         | 39.299/0.948         | 39.303/0.952                |
|               | C3          | 35.666/0.921         | 35.587/0.902                | 37.187/0.937         | 37.246/0.951         | 37.177/0.943                |
|               | C4          | 32.706/0.928         | 32.665/0.911                | 35.501/0.943         | 35.512/0.949         | 35.528/0.956                |
|               | C5          | 39.291/0.926         | 39.183/0.916                | 39.452/0.925         | 39.453/0.925         | 40.064/0.938                |
|               | C6          | 39.584/0.931         | 39.132/0.916                | 39.392/0.927         | 39.405/0.934         | 40.312/0.941                |
|               | C7          | 40.362/0.951         | 40.356/0.952                | 40.799/0.952         | 40.836/0.950         | 40.848/0.956                |
| 3X            | C1          | 34.674/0.852         | 34.681/0.867                | 36.117/0.818         | 36.125/0.824         | 36.145/0.872                |
|               | C2          | 35.363/0.866         | 35.382/0.873                | 37.497/0.873         | 37.549/0.878         | 37.899/0.880                |
|               | C3          | 33.717/0.896         | 34.102/0.902                | 34.128/0.905         | 35.412/0.917         | 35.423/0.928                |
|               | C4          | 29.396/0.841         | 30.016/0.852                | 30.721/0.854         | 31.432/0.861         | 32.432/0.864                |
|               | C5          | 37.909/0.851         | 37.912/0.854                | 38.872/0.859         | 38.953/0.868         | 39.853/0.868                |
|               | C6          | 36.154/0.871         | 36.250/0.883                | 37.700/0.887         | 37.644/0.891         | 38.784/0.906                |
|               | C7          | 37.132/0.907         | 37.232/0.911                | 38.110/0.912         | 38.835/0.921         | 39.721/0.921                |
| 4X            | C1          | 31.778/0.729         | 32.233/0.834                | 32.423/0.841         | 32.431/0.844         | 32.991/0.854                |
|               | C2          | 33.078/0.692         | 33.234/0.673                | 34.276/0.725         | 34.176/0.698         | 35.220/0.715                |
|               | C3          | 31.156/0.753         | 31.252/0.765                | 32.673/0.789         | 32.992/0.789         | 33.562/0.798                |
|               | C4          | 31.014/0.706         | 31.118/0.716                | 32.403/0.727         | 32.410/0.731         | 33.003/0.758                |
|               | C5          | 34.389/0.733         | 34.397/0.741                | 35.749/0.759         | 35.851/0.764         | 36.513/0.794                |
|               | C6          | 33.726/0.702         | 33.741/0.735                | 34.195/0.777         | 35.687/0.783         | 36.478/0.759                |
|               | C7          | 35.099/0.823         | 35.163/0.836                | 36.982/0.893         | 36.684/0.887         | 37.726/0.912                |
results clearly indicates that the low classification accuracy on LR images and reveals these images holds less detailed information for RPD classification.

The overall classification accuracy of RPD classification using deep learning-based CNN classification models for the images with 2–4 scaling factor is shown in Fig. 5. This graph indicates the clear perception that classification accuracy for super resolution images with 2X scaling is always higher than all CNN models and super-resolution images outperform for all scaling factor. In terms of deep learning CNN models ReCNN shows greater results for all kind of images along with varying scaling factors.

Figure 4: Comparison of MSE, PSNR, and SSIM for different magnification scales a) MSE b) PSNR c) SSIM

Figure 5: Comparison of average accuracy value over different classification models with varying scaling factors (2–4)
Subsequently, the integration of convolution based super-resolution layers before the traditional convolution layers shows better improvement in terms of classification accuracy results. Therefore, proposed RDA-CNN method motivates to use the super resolution layers prior to any CNN based network architectures (convolution layers) for automatic rice plant disease classification with detailed identification of disease global characteristics (lesions, spots, and rot so on). Finally, the performance of RDA-CNN with VGG19 architecture achieves the highest disease classification accuracy of 93.257% and thereby compares with the different deep learning architectures for automatic RPD classification.

5 Conclusion

This paper proposes and verifies the effective Reconstructed Disease Aware–Convolutional Neural Network (RDA-CNN) to accelerate the image-based plant disease classification in rice plants. First, RDA-CNN highlighted the importance of super-resolution layers to enhance the quality of the image. The results of super-resolution layers indicated that DSRCNN has significantly outperformed for the magnification scales of 2X, 3X and 4X in terms of various image quality assessments like MSE, PSNR and SSIM for all seven classes of rice diseases. Experimental analysis shows that the proposed DSRCNN can improve the PSNR value by 2%, SSIM value by 0.05% and similarly MSE value also reduced compared to conventional SR methods. Next, classification layers perform disease classification using ReCNN and results are compared with different CNN architectures. In addition, RDA-CNN outperforms with the average classification accuracy of 96.595% and thus helps the experts to detect lesions, discoloration, rust and spots and so on in earlier stages of the rice plants. These results indicate that the classification accuracy of generated super-resolution images can be significantly improved for the proposed network architecture. Although RDA-CNN has several advantages compared to baseline SR methods, application of proposed method to detect multiple diseases in a single leaf is still a limitation that needs redressal. In future, super-resolution algorithm can be integrated with object detection, segmentation of blobs etc. with pest identification applications for optimizing yield.

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Table 5: Comparison of average accuracy of RPD classification between traditional deep learning and SR based deep learning methods

| Classification models | Traditional deep learning methods Accuracy (%) | SR based deep learning methods Proposed DSRCNN based classification models Accuracy (%) |
|-----------------------|-----------------------------------------------|-------------------------------------------------------------------------------------|
| AlexNet               | 86.58                                         | AlexNet 93.24                                                                     |
| VGGNet                | 84.52                                         | VGGNet 91.57                                                                       |
| DCNN                  | 95.04                                         | ResNet 94.53                                                                       |
| Simple CNN            | 94.33                                         | Inception 90.05                                                                   |
| Inception             | 72.09                                         | RDA-CNN 96.59                                                                     |
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