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Segmentation Method for Automatic Leaf Disease Detection

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ABSTRACT: Present paper is an attempt to automatically detect leaf diseases. Automatic Detection of Leaf Disease is an essential research topic in agricultural research as it may prove benefits in monitoring fields, and early detection of leaf diseases by the symptoms that appear on the leaves. This paper provides various methods for leaf disease detection using Image Processing. The developed system consists of Image Acquisition, Image Pre-Processing, Image Segmentation and Feature Extraction. Image Pre-Processing involves HSI transformation, RGB splitting. Image Segmentation involves selecting best segmentation method among Region based segmentation and Mean-Shift Segmentation based on Quality Metric Parameters, Histogram analysis and Computational Time analysis. Feature extraction involves extracting morphological and color features for each disease spot.

KEYWORDS: Region Based Segmentation Methods; Mean-Shift Segmentation Method; Quality Metric Parameters; Hue Value Feature Extraction;

I. INTRODUCTION

India is an agriculture country. Major portion of India's population depend on agriculture as food source and also as an economical income. Agriculture crops suffer losses due to diseases that affect plant leaves, crops, stem and so on. [1] The conventional way of leaf disease identification and detection is naked eye observation by experts to detect the leaf disease by the symptoms that appear on the leaf. The naked eye observation requires continuous monitoring and is not feasible for monitoring large farms. Hence, automatic detection of leaf disease is required. Automatic detection of leaf disease is an essential research topic in agricultural research as it may prove benefits in monitoring fields and early detection of leaf diseases.

Image segmentation refers to the process of partitioning a digital image into multiple regions (sets of pixels). The result of image segmentation is a set of regions that collectively cover the entire image, or a set of contours extracted from the image.Each pixel within a region is uniquely similar with respect to some characteristic or computedproperty, such as colour, intensity, or texture, depth, or motion. Pixels from adjacent regions are significantly different with respect to the same characteristic(s). This makes the object differentiation, isolation and detection task easier. Image segmentation is one of the primary steps in image analysis for object detection, recognition and identification.

Image Segmentation algorithms are based on two principles: discontinuity and similarity. The principle of discontinuity is to extract regions that differ in properties such as intensity, color, texture etc. In the principle of similarity pixels are grouped together to form regions based on some pre-defined criteria. Thresholding based segmentation, clustering based segmentation, Region based segmentation are some of the Similarity based segmentation methods.

The Segmentation algorithms can be classified based on pixel similarity relationship with neighbouring pixels as: region-based algorithms and pixel-based algorithms.

In plant leaf disease detection the following steps are used:

i. The input image (RGB image) is converted to HSI color model.

ii. The diseased regions are extracted using the best segmentation method.



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iii. From the extracted disease regions, the features (morphological, color) are extracted. iv. Using the extracted features the disease is identified using classification methods.

II. VARIOUS SEGMENTATION METHODS

The fig 1 shows the types of segmentation methods used:



Fig 1: Image Segmentation Methods

A. A.Region Based Segmentation Methods

A region denoted by 'r' of an image is defined as a connected homogenoussetofthe image with respect to some criterion such as grey level or texture. Regions in an image are a group of connected pixels with similar properties. In the region approach, each pixels assigned to a particular object orregion. In this segmentation, pixels corresponding to an object are groupedtogether andmarked.

The Main Methods of Region Segmentation [3] are Region Growing, Region Splittingand RegionMerging.

1) Region Growing Segmentation

Regiongrowingisatechniqueforextractingaregionoftheimagethatisconnected

basedonintensityinformation.InRegiongrowingneighbouringpixelsareexaminedand added to a region class of number of edges aredetected.Intensity image of input image is used for region growing segmentation.

Algorithm – Region Growing

Input: Color Image

Output: Greyscale Segmented image

STEPS:

1. Consider the input RGB image. Convert the RGB image into HSI image and consider I buffer.

2. Assign a seed pixel for the input image. Histogram peak value is chosen as the pixel.

3. Fix a homogeneity criterion as the similarity variation tolerance (threshold) which is computed using standard deviation formula.

4. Consider the seed pixel and compare its 8-neighbors with the threshold value. Suppose, the value is less than the threshold, the pixels are added to the existing regions. Otherwise, it forms another region.

Repeat the procedure until every pixel in the frame belongs to any one of the region.

2) Region Merging Segmentation

Merging must start from a uniform seed region. Initially, a suitable seeded region must be chosen. One method is to divide the image into 2x2 or 4x4 blocks and check each one. Another is to divide the image into strips, and then



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subdivide the strips further. In the worst case the seed will be a single pixel. Once a seed has been found, its neighbours are merged until no more neighbouring regions conform to the uniformitycriterion. At this point the region is extracted from the image, and a further seed is used to merge another region.

Algorithm – Region Merging

Input: Color Image

Output: Greyscale Segmented image

STEPS:

1. Consider an input RGB image and convert the RGB image into HSI image and consider I buffer and find its image matrix

2. Begins the segmentation process with regions of 2x2, 4x4 or 8x8 blocks of pixels.

3. Compare the statistical grey level properties of each block with the adjacent block; if they match, they are merged into a larger region. Otherwise regions are marked as non-matching.

4. Merging of adjacent regions continues between all neighbors, including newly formed ones. If a region cannot be merged with any of its neighbors, it is marked as 'final' and the merging process stops when all image regions are so marked.

3) Region Splitting Segmentation

These algorithms begin from the whole image, and divide it up until each sub region is uniform. The usual criterion for stopping the splitting process is when the properties of a newly split pair do not differ from those of the original region by more than a threshold.

Algorithm – Region Splitting

Input: Color Image

Output: Greyscale Segmented image

STEPS:

- 1. Consider an input RGB image and convert the RGB image into HSI image and consider I buffer. Find the image matrix of the 'I' buffer.
- 2. Assume the whole image as a single region.
- Calculate the maximum and minimum intensity values.
 If the difference between minimum and maximum value is greater than the predetermined threshold then split the existing region into new regions.
- 5. Check if all the pixels contained in each of the regions satisfy the similarity constraint. If TRUE, then the area of interest corresponds to existing region. If FALSE split the area of interest (usually into four equal subareas) and consider each of the sub-areas as the area of interest in turn.
- 6. This process continues until no further splitting occurs. In the worst case this happens when the areas are just one pixel in size.

B. B. Mean-Shift Image Segmentation

Mean Shift method is a data clustering method. For each data point, mean-shift defines a window around it and computes the mean of data point, then it shifts the center of the window to the mean and repeats the algorithm till it converges. The Mean-Shift algorithm works based on the histogram of the image.

Algorithm – Mean-Shift Image Segmentation

Input: Color Image

Output: Greyscale Segmented image

STEPS:

The idea of Mean-Shift is to start with windows centered on a random bin and shift the center of the bin according to the data within it until the window converges at the mode of the histogram.

The shifting of the window is as follows:

1. Initialize with a random seed that selects a bin and select the window W centered on that bin.

2. Calculate the center of gravity or weighted mean wm of the histogram values in window W

wm = $\sum b_i f(b_i)$ b_i€ W



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Where f (bi) is the count in bin bi

3. Translate the search window W to the weighted mean wm

4. Repeat step 2 until convergence.

III. QUALITY METRICS FOR SEGMENTATION

The quality metrics are used as the measures, used to define the consistency of the algorithms. Using these measures, the better algorithm for region based segmentation is analysed and the formulae for the different measures are as follows [13]:

A. Grey Level Energy

The grey level energy indicates how grey levels are distributed [16]. It is formulated as:

$$E(x) = \sum_{i=1}^{k} p(i)^{2}$$

WhereE(x)represents the grey level energy with 256 bins and p(i)refers to the probability distribution functions, which contains the histogram counts. The larger energy value corresponds to the lower number of grey levels, which means simple. The smaller energy corresponds to the higher number of grey levels, which means complex.

B. Discrete Entropy

The discrete entropy is the measure of image information content, which is interpreted as the average uncertainty of information source. Discrete entropy is formulated as:

$$H(x) = \sum_{i=1}^{k} p(i) \log_2 \frac{1}{p(i)} = -\sum_{i=1}^{k} p(i) \log_2 p(i)$$

Where p (i) is the probability distribution, considering all histogramcounts.

Resultshaveshownthattheentropyoftheproperlysegmentedimageismuchsmaller than that of the sourceimage.

C. Normalized Mutual Information

The mutual information [13] can be applied as another objective metric. The mutual information acts as a symmetric function, which is formulated as:

$$I(X;Y) = \sum_{X|Y} p_{XY}(X, Y) \log_2 \frac{p_{XY}(X, Y)}{p_X(X)p_T(Y)}$$

= $-\sum_X p_X(X) \log_2 p_X(X) + \sum_{X,Y} p_{XY}(X, Y) \log_2 \frac{p_{XY}(X, Y)}{p_T(Y)}$
= $H(X) - H(X|Y)$

Where I(X;Y) represents the mutual information; H(X) and H(X|Y) are entropy and conditional entropy values. It is interpreted as the information that Y can tell about X is equal to the reduction in uncertainty of X due to the existence of Y. In Images egmentation, the less ervalue of the mutual information indicates the better match between the source and processed images.

Normalized Mutual Information [17] is a well-defined measure covering contents from both discrete entropies and mutual information. It is formulated as:

$$N_{MI} = \frac{I(X; Y)}{\sqrt{H(X)H(Y)}}$$

Where I(X, Y) is the mutual information; H(X) and H(Y) are the discreteentropies. Onceagain, similar to that of the mutual information,

thebettermatchbetweenthesourceandprocessedimages, the smaller the normalized mutual information.

D. Information Redundancy

Another symmetric information measure can be used to indicate redundancy [18]in image segmentation. It reaches the minima of zero when all variables are independent, bigger information redundancy, the greater dependence between two images.

$$R = \frac{I(X; Y)}{H(X) + H(Y)}$$



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Where H(X) and H(Y) are entropies of two images and I(X; Y) is the Mutual Information.

IV. PROPOSED METHOD

Based on the analysis done in section V. Region Growing Segmentation method is chosen for separating healthy and diseased regions of the leaf.Fig 2 shows the proposed method using region growing segmentation.



Fig 2: Flow chart of the proposed method

A. Steps Followed

1. Input image is split to R, G and B components. The Green buffer is considered forfurther processing.

2. Again the input image is converted to HSI color model [6], Intensity componentis considered forsegmentation.

3. Region growing segmentation is applied on Intensity component of HSI colormodel.

4. Now, Histogramisplotted [5] on region growing segmented image and the high estpeak is calculated [neglecting the background peak].

5. The peak gives us the intensity of green pixels. This peak is used to segment the diseased spots.

6. The greenbufferfromstep1isconsidered. The peak obtained from step4is used and pixels are compared with the peak value. The pixels that doesn't match the range of peak value are considered to be diseased pixels. In other words masking and removing green pixels.

7. The diseased pixels obtained from step 6 are displayed. This forms the diseased regions.

8. Now, connected component labelling is used to label the diseased spots and diseased spots are pushed onto thestack.



Disease: Tar Spots Disease: Bacterial Leaf Spots

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9. Each diseased spot is popped from stack in order to extract Shape features[area,perimeter, centroid, roundness and compactness] and color features [Mean,Standard deviation, Entropy, Skewness and SpatialFrequency].10. All these features are written into a ".txt" file for further analysis.

V. EXPERIMENTAL RESULTS AND ANALYSIS

A. Sample Database



Leaf 1: Soybean Leaf Leaf 2: Ivy LeafLeaf 3: Maple Leaf Disease: Brown Spots Disease: Leaf Spots Disease: Tar Spots

B. Analysis to Find the Best Region Based Segmentation

1) Quality Metric Analysis

TABLE I. shows the various quality metric parameters used to find the best region based segmentation method. The Quality Metric values in TABLE 1 are for Leaf 4 that is Maple leaf with tar spots disease. RG in TABLE I. stands for Region Growing Segmentation, RM stands form Region Merging Segmentation and RS stands for Region Splitting Segmentation.

Quality Metric Parameters	RG	RM	RS
Discrete entropy	1.136	4.128	3.659
Grey level energy	0.536	0.252	0.354
Mutual Information	0.282	3.752	3.659
Normalized Mutual Information	0.127	0.886	0.719
Image Redundancy	0.051	0.443	0.358

TABLE I. Analysis to Find the Best Region Based Segmentation Based On Quality Metric Parameters

2) Computation Time Analysis

The Computation Time values in TABLE II. are for Leaf 4 that is Maple leaf with tar spots disease. RG in TABLE II. stands for Region Growing Segmentation, RM stands form Region Merging Segmentation and RS stands for Region Splitting Segmentation.



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TABLE II. Analysis To Find The Best Region Based Segmentation Based On Computation Time

Computation Time (seconds)	RG	RM	RS
Leaf 4: Maple Leaf Disease: Tar Spots	0.015	0.046	3.234

3) Observation

Region based different segmentation timing analysisis is applied on images and the doneonthesemethodsasshownabove. Basedontiming andQuality Metrics, it is clear that Region growing has the maximum peaks representing distinct regions with least discrete entropy and highest grey level energy when compared withother regionbasedsegmentationmethods.

C. Analysis To Find The Best Segmentation Method Among Region Growing And Mean-Shift Segmentation

1) Quality Metric Analysis

a) Grey Level Energy

The larger energy value corresponds to the lower number of grey levels, which means simple. The smaller energy corresponds to the higher number of grey levels, which means complex.

TABLE III. gives the grey level energy for region growing and mean-shift segmentation for 10 diseased leaves few of which are listed in section V (A).

Sl. No.	Region Growing	Mean-Shift
Leaf 1	0.536	0.494
Leaf 2	0.437	0.405
Leaf 3	0.446	0.420
Leaf 4	0.599	0.590
Leaf 5	0.505	0.439
Leaf 6	0.675	0.681
Leaf 7	0.640	0.628
Leaf 8	0.912	0.915
Leaf 9	0.587	0.591
Leaf 10	0.812	0.829

TABLE III. Quality Metric Analysis Based On Grey Level Energy

b) Discrete Entropy

The entropy of the properly segmented image is much smaller than that of the source image, which are much simpler, when the image pixels are distributed among more grey levels, the entropy value will increase.

TABLE IV. gives the discrete entropy for region growing and mean-shift segmentation for 10 diseased leaves few of which are listed in section V (A).



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Sl. No.	Region Growing	Mean-Shift
Leaf 1	1.136	1.582
Leaf 2	1.377	1.743
Leaf 3	1.359	1.697
Leaf 4	1.025	1.171
Leaf 5	0.993	1.442
Leaf 6	0.971	0.957
Leaf 7	0.876	1.071
Leaf 8	0.327	0.325
Leaf 9	1.178	1.274
Leaf 10	0.658	0.610

TABLE IV. Quality Metric Analysis Based On Discrete Entropy

c) Mutual

In Image segmentation, the lesser value of the mutual information indicates the better match between the source and processed images.

TABLE V. gives the mutual information for region growing and mean-shift segmentation for 10 diseased leaves few of which are listed in section V (A).

Sl. No.	Region Growing	Mean-Shift
Leaf 1	0.282	0.070
Leaf 2	0.357	0.105
Leaf 3	0.394	0.107
Leaf 4	0.322	0.054
Leaf 5	0.429	0.067
Leaf 6	0.259	0.031
Leaf 7	0.296	0.028
Leaf 8	0.079	0.011
Leaf 9	0.303	0.047
Leaf 10	0.148	0.022



d) Normalized Mutual Information

Similar to the mutual information, smaller normalized mutual information results the better match between the source and processed images.

TABLE VI. gives the discrete entropy for region growing and mean-shift segmentation for 10 diseased leaves few of which are listed in section V (A).



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Sl.No.	Region Growing	Mean-Shift
Leaf 1	0.051	0.012
Leaf 2	0.059	0.017
Leaf 3	0.071	0.018
Leaf 4	0.089	0.014
Leaf 5	0.083	0.010
Leaf 6	0.084	0.008
Leaf 7	0.089	0.011
Leaf 8	0.080	0.011
Leaf 9	0.070	0.011
Leaf 10	0.068	0.011

TABLE VI. Quality Metric Analysis Based On Normalized Mutual Information

2) Computation Time Analysis

TABLE VII. Computation Time Analysis

	Execution Time in Seconds			
Sl. No.	Region Growing	Mean-Shift		
Leaf 1	0.344	1.235		
Leaf 2	2.406	0.875		
Leaf 3	1.910	4.547		
Leaf 4	0.344	1.438		
Leaf 5	1.968	3.438		
Leaf 6	1.125	2.172		
Leaf 7	0.750	1.656		
Leaf 8	0.235	0.735		
Leaf 9	1.531	2.750		
Leaf 10	0.250	0.985		

3) Observation

Region growing segmentation and mean-shift segmentation is applied on different images and the timing analysis is done on these methods as shown above. Based on timing analysis and Quality Metrics, it is clear that Region growing has the maximum peaks representing distinct regions with least discrete entropy and highest grey level energy when compared with mean-shift segmentation methods. Hence Region Growing Segmentation is used as the segmentation method for further processing.

D. Results of the Proposed Method

Fig 3 shows the results of proposed method using Region Growing Segmentation. The input image Fig 3(a) isconverted to HSI image and I component is considered for segmentation. The segmented image is as in Fig 3(b).Input ImageRegion Growing SegmentationHistogram of Segmented ImageGreen Buffer

on Intensity Image

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Fig 3: (a) Maple leaf with tar spots as input image, (b) Region Growing Segmentation on Intensity Image, (c) Histogram of segmented image, (d) Green buffer of input image, (e) Segmented Lesions (diseases spots), (f) Healthy region, (g) Total area of healthy and diseased regions and percentage of infection.

E. Hue Value Feature Extraction

The segmented lesions from step 3 is converted to HSI image and the hue image is considered for processing. The hue value of each disease spot is extracted. This is done because the hue value gives the actual color information of the image. Once the hue value is extracted, the area of each diseases spot is calculated and using this area the percentage of area infection is calculated as shown in TABLE VIII.

The hue values are clustered using **K-Means clustering**[18]. K value is chosen to be 3 that is hue values are clustered into 3 clusters. Cluster 1 indicates moderately infected regions, cluster 2 indicates less infected regions and cluster 3 indicates highly infected regions. In order to visually express the clusters, the clusters are assigned false colors. Cluster 1 is assigned red color, cluster 2 is assigned blue color and cluster 3 is assigned yellow color.

1) Experimental Results of Hue Value Feature Extraction

Fig 4 shows the Hue value feature extraction. The input image Fig 4(a) is segmented using Region Growing segmentation. The segmented lesions is mapped back to Fig 4(a) to get the RGB lesions. These lesions are converted to HSI image and the Hue image is considered for processing, this gives Fig 4(c). The segmented lesions are clustered using K-Means clustering to indicate different infected regions and for visually expressing this clustering false coloring is used.



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Blue Color : Less Infected Regions

Red Color : Moderately Infected Regions

> Fig 4: (a) Maple leaf with tar spots as input image, (b) Segmented lesions from inputimage, (c) Hue image of segmented lesions, (d) K-Means clustered lesions with false coloring

I ABLE VIII. Hue value and % of Area Infection					
Spot No.	Spot Area	Hue Value	% of Area Infection		
1	201	172	0.373148		
2	65	144	0.12067		
3	258	189	0.478966		
4	231	190	0.428842		
5	42	142	0.077971		
6	204	171	0.378718		
7	205	161	0.380574		
8	175	176	0.32488		

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IABLE VIII.	Hue	value	and	% O	i Area	Intection

TABLE IX. (A), (B), (C) Shows The K-Means Clustered Lesions Initial Cluster Center are <189.0>, <190.0>, <171.0>

TABLE IX (A)	TABLE IX (B)	TABLE IX (C)
Cluster 1: Red Color	Cluster 2: Blue Color	Cluster 3: Yellow Color
Final Cluster Centre:	<170.0>Final Cluster Centre: <189.0>	Final Cluster Centre: <143.0>

Spot#	Hue Value				
1	172				
6	171	Spot#	Hue Value		
7	161	3	189	Spot#	Hue Value
8	176	4	190	2	144
		-		5	142

VI. CONCLUSION AND FUTURE WORK

The work proposed in this project is a very useful work for agriculture application. A number of different research work will be emerging from the basic work done in this project. Some of the future deductions will be as follows:

1. Segmentation is one of the most important image processing methodology which is used in this project to separate the disease regions of the leaf from healthy area. In this proposed work we have used Region Based segmentation methods to find out the disease regions. Out of the three basic Region based segmentation methods such as Region Growing, Region Merging and Region Splitting we have derived methodologies how to select the best one. The best Region based segmentation is Region Growing, details of which is explained in the work.

A large number of segmentation methods are coming up from researchers. The future deductions of research is to use different kinds of segmentation and different Quality Metric parameters to find which the best is.

2. In the present work the K-Means Clustering is been used. A large number of clustering methods under Data Mining Techniques like Agglomerative, CURE, OPTICS, Chameleon etc. are available in the literature. A future research work would be using various Data Mining Techniques.

3. One of the most useful application would be to link the damage area due to disease with its remedial procedure.

4. The entire system would be used as an Expert System to deliver the farmers some useful information.



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