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CLASSIC APRIORI: IMPLEMENTATION OF MODIFIED APRIORI ALGORITHM IN MATLAB FOR PADDY CROPS DISEASE PREDICTION USING DATA MINING TECHNIQUES

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ABSTRACT

Automation of essential processes in agriculture is becoming widespread, especially when fast action is required. However, some processes that could greatly benefit from some degree of automation have such difficult characteristics, that even small improvements pose a great challenge. This is the case of paddy disease diagnosis, a problem of great economic, social and ecological interest. Difficult problems like this often require a interdisciplinary approach to be tackled properly, as multifaceted issues can greatly benefit from the inclusion of different perspectives. In this context, this paper presents the most recent advances in research subjects such as data mining algorithm applied to paddy disease diagnosis, and give the solution, and discusses how those advances may be combined to support future developments towards more effective diagnosis tools. The paper finishes suggesting a possible solution to increase the degree of automation of paddy disease diagnosis tools.

Keywords: Classification, Association Set Rules, K-Nearest Neighbor, Apriori Algorithm Artificial Neural

1. INTRODUCTION

The use of automation in agriculture is not new, and as the cost of technology plummets, its use becomes increasingly widespread. However, the technical challenges vary greatly depending on the problem to be tackled. While some simpler automation systems, like timed irrigation, have been solved long ago, others require a much higher level of sophistication in order to become feasible. This is the case of automated disease diagnosis, and particularly so for paddy diseases.

Fast and reliable diagnosis of diseases is essential if the problem is to be handled properly, no matter if we are dealing with a person, an animal, or a plant. The best way to achieve that is by having a specialist readily available to assess the situation and to suggest some line of action. However, this is not always possible, especially in remote places. This is particularly true in the case of agriculture, as those are often located in places far from urban centers.

There are basically three approaches that can be used to overcome this problem. All three employ technological tools, but they possess different degrees of sophistication and automation, as shown below:

- Remote diagnosis system: in this type of approach, the person interested in obtaining a diagnosis sends information through some communication channel (usually internet or telephone), and such an information is analyzed by a human specialist, who sends a diagnosis (or a request for more information) back to that person. Because it has a low level of automation, this approach tends to be time demanding. On the other hand, the involvement of a human specialist improves the chances of a correct diagnosis. This option should be preferred when there are no reliable methods with a higher degree of automation, which still is the case for the vast majority of cases.

- Expert systems: this kind of system tries to emulate the decision-making ability of a human expert (JACKSON, 1998). They employ a series of rules in order to reach a diagnosis. Human experts are essential during the development of the tool, but they usually do not have any involvement once the tool is put into use – in general, the user answers a number of questions about the symptoms of the

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disease, until the system has enough information to provide a diagnosis.

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This may be considered a semi-automatic diagnosis method, as the computer program depends on a number of human inputs in order to work properly. The higher degree of automation with respect to the previous approach means that it will be faster. Another advantage of this kind of system over the remote one is that it is usually cheaper to maintain. On the other hand, the quality of the diagnosis will be only as good as the inference engine and the knowledge base employed in the system, and design flaws may lead to error. Since the system depends on human input to reach a diagnosis, and such an input is usually provided by a non-expert, this may also give rise to incorrect outputs. A discussion about this kind of tool, together with a brief description of some of the most relevant systems proposed so far.

- Image-based automatic systems: this kind of system explores color and morphological characteristics of an image of the symptoms to infer the disease. Since there is no human interaction, this kind of approach tends to be very fast. Moreover, such a high degree of autonomy means that a system like this may be used to monitor vast areas without the need for humans to go to the field. This implies that a really reliable image-based system may be a much more advantageous option in comparison with the other two approaches.

The problem is that achieving such a reliability is very challenging, being the main bottleneck for the dissemination of this kind of system. Despite the challenges involved, automatic diagnosis systems are becoming common in several areas of agriculture and medicine, but the difficulties involved in agriculture have prevented any method with focus on paddy to be proposed so far. To the author's knowledge, the only work that has some relation with this kind of approach was proposed by (PARK; OH; HAN, 2007), which has the objective of providing a diagnosis by identifying pathogens in microscopic images. A deeper discussion about image-based automatic diagnosis systems.

Some disease diagnosis problems have already achieved a high degree of automation. In the case of paddy diseases, however, the successes have been limited to a handful of expert systems, most of them designed to deal with a single species and a limited set of diseases. This is due to the very difficult characteristics surrounding the problem, such as the large variety of species, the symptoms heterogeneity, the difficulties involved in gathering data in the field. The use of digital imaging is particularly challenging: according to (ZION, 2012), "the sensing technology has to overcome limited visibility, temporal and spatial variations in lighting, varying distances and relative orientations between cameras and objects, motion and density of the monitored targets, and even lack of physical stability." Those challenges suggest that an interdisciplinary approach may be needed in order to increase the degree of automation in diagnosis. It is important to emphasize that a fully automated system for disease diagnosis in paddy would probably require very sophisticated artificial intelligence, capable of autonomously performing analysis based on images, biopsy samples, water samples, etc. Such a technology is far beyond what is available today.

In this context, the objective of this manuscript is to discuss some possible ways to gradually increase the automation of the process, by analyzing how the most current advances in areas such as expert systems for disease diagnosis in aquaculture, imagebased disease diagnosis in agriculture (especially paddy), could be combined into a more effective diagnosis tool.

2. DATA MINING FUNCTIONALITIES

Functionalities of data mining are as follows-

- Characterization

Data characterization is a summarization of the general characteristics or features of objects in a target class of data. Relevant data to a user-specified class are collected by a database query and run through a summarization module to extract the essence of the data at different level of abstractions.

- Discrimination

Data discrimination is a comparison of the general features of target class data objects with the general features of objects from one or a set of contrasting class. Discrimination descriptions expressed in rule form are referred to as discriminant rules.

2.1 Association Analysis

- Association analysis is the discovery of association rules. It studies the frequency of items take place together in transactional databases based

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on threshold called support and confidence. Association analysis is commonly used for market basket analysis. The discovery of association rule can help retailers to develop marketing strategies by gaining into which items are frequently purchase together.

For example- age $(X, "20-29")^{F}$ income (X, "10-20k") => dropout (X, "Yes") [Support= 2%, confidence=60%]

Rule implies that 2%(support) are 20-29 year of age with an family income of 10-20k generally drop their study and 60% probability(confidence) that a student in this age and income group will withdraw from their study.

2.2 Classification and Prediction

Classification analyzes a set of training data and builds a model for each class based on the features in the data. This model is used to classify new objects and also known as supervised learning. Derived model may be represented in various forms, such as classification (IF- THEN) rules, decision trees, and neural network etc. A decision tree is a flow-chart like tree structure where each node denotes a test on an attribute value and each branch represents an outcome of the test. A neural network is a collection of linear threshold units that can be trained to distinguish objects of different classes. Classification can be used to predicting the class label of data objects. Classification predicts categorical labels (or discrete values), prediction models continuous- valued function.

2.3 Clustering

Clustering is an unsupervised learning, in which the class labels of the training samples are not known. Cluster is a collection of data objects that are similar to on one another. Similarity can be expressed by distance functions, specified by use experts. A good clustering method produces high quality clusters to ensure that inter- cluster (object of different class) similarity is low and the intracluster (object in a same class) similarity is high.

2.4 Evolution and Deviation Analysis

Evolution and deviation analysis pertain to the study, trend of object whose behavior changes over time. Evolution analysis models evolutionary trends in data, whereas deviation analysis, considers differences between measured values and expected values.

3. OBJECTIVE OF THE STUDY

Data mining is the process of extracting patterns from the data and it is an automated extraction of hidden predictive information from large databases. The main aim of the proposed Classic Apriori Algorithm(MAA) is to maintain a good health in between the paddy are based on the performance provided by those crops by using association rule mining (ARM). Its objectives are to increase paddy plant health. Often, information related to paddy name, disease,Symptoms are analyzed by means of simulations, predictions and classification of customers.

- There are two main characteristics of plant disease detection using machine-learning methods that must be achieved, they are: speed and accuracy. Hence there is a scope for working on development of innovative, efficient & fast interpreting algorithms which will help plant scientist in detecting disease.
- Work can be done for automatically estimating the severity of the detected disease.
- The main outcome of this research is an android based software tool to find and cure paddy diseases for farmers
- This work proposes two databases which contain paddy disease syndromes images and treatment experiences which can further extendable.
- To identify the concepts and techniques using ARM to improve the efficiency and effectiveness in a data mining environment
- To design and develop a software prototype to prove the above concepts
- To test a prototype

3. MOTIVATION OF THE RESEARCH

As mentioned earlier, data mining has been widely used to solve various kinds of discovering a set of association rules problems in numerous applications. However, association rules have turned out to be one of the most persistent problems that encompasses many diverse applications in the data mining field. These problems have attracted

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more active research in order to find efficient approaches to address them, and the outcome of the research is still unsatisfactory.

• The ultimate goal of association rule is to discriminate new rule data into the most likely of the specific predictive models, based on the induction model generated by the classifier. However, the association rule problem has become very difficult and computationally infeasible when the number of possible different combinations of variables are so high.

4. RELATED WORK

Z.Xue et al (2003) proposed Bayesian shape model (BSM) to find contour points in the face. A full-face model consisting of the contour points is designed to describe the face patch, using which the normalization of the extracted face patch can be performed efficiently. In BSM, the prototype of the face contour can be adjusted adaptively according to its prior distribution. Moreover, an affine invariant internal energy term is introduced to describe the local shape deformations between the prototype contour in the shape domain and the deformable contour in the image domain. The face patch is extracted and normalized using the piecewise affine triangle warping algorithm[5].

A.Camarago et al (2009) proposed color transformation method. This method converting the RGB image of the disease infected plant or leaf, into the H, I3a and I3b color transformations. The I3a and I3b transformations are developed from a modification of the original image in to color transformation to meet the requirements of the plant disease data set. The transformed image is then segmented by analyzing the distribution of intensities in a histogram. The threshold cut-off value is determined according to their position in the histogram. This technique is particularly useful when the target in the image data set is one with a large distribution of intensities [2].

S.L.S. Abdullah et al (2012) proposed improved thresholding based technique for image segmentation. Different illuminations may produce different color intensity of the object surface and thus lead to inaccurate segmented images and traditional methods were unable to produce good quality segmented. Therefore, an improved thresholding based segmentation integrated with an inverse technique (TsTN) that was able to partition natural images correctly. The analysis results showed that TsTN has the ability to produce good quality segmented images for dark images. Furthermore, this segmentation technique was proven to be more accurate than the traditional thresholding and clustering techniques [1].

A.K Das et al (2012) proposed SVM and Bayes' classifier to classify the diseases of rice plant leaf. An automated system has been developed to classify the leaf brown spot and the leaf blast diseases of rice plant based on the morphological changes of the plants caused by the diseases. Otsu method is used to isolate the infected region from the background. Radial distribution of the hue from the center to the boundary of the spot images has been used as features to classify the diseases by Bayes' and SVM Classifier [3].

R.Lu et al (2013) proposed Support Vector Machine for classification and Otsu method for segmentation. This method automatically adjusts the classification hyper plane calculated by using linear SVM and requires minimum training and time. It also avoids the problems caused by variations in the lighting condition and color of the fruit. Disadvantage of this method is segmentation is not accurate [6]

Classification Rules

f appearance = spot and color = olive Then disease = blight	
f appearance = spot and color = brown and a 55	ige <=
Then disease =brown-spot	

5.. RESULT AND DISCUSSION

One hundred and fifty samples of each data class are considered to test the system. For the first level classification (i.e. uninfected leaf or a diseased paddy leaf), it has been found that success rate is 92%, 96.4% and 84% respectively for uninfected paddy leaf, leaf with brown spot and leaf with blast, summarized in Table 1.

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Table 1. Result Of Classification Of Uninfected And

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	Diseased Le	raves
Image Type		Success Rate (%)
N	CT.	02

Normal Paddy leaf Image	92
Brown Spot Image	97.5
Blast Paddy Image	86

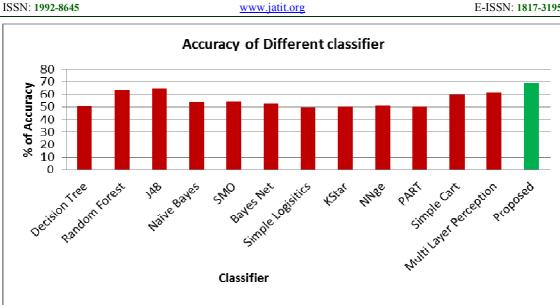
Table 2: Comparison Of Classification Error Of Algorithms

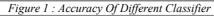
Classifier	Accuracy(%	True) Positive Rate	False Positive Rate	Precision(%)	Recall(%)	Classification Error (%)	n Kappa Statistics	RMS Error
Decision Tree	50.68	0.507	0.23	0.478	0.507	49.32	0.211	0.404
Random Forest	63.34	0.633	0.254	0.57	0.633	36.66	0.354	0.313
J48	64.45	0.5	0.544	0.521	0.5	35.55	0.344	0.31
Naïve Bayes	53.75	0.571	0.594	0.528	0.571	46.25	0.484	0.539
SMO	54	0.643	0.465	0.629	0.643	46	0.589	0.632
Bayes Net	52.5	0.681	0.502	0.625	0.681	47.5	0.585	0.536
Simple Logisitics	49.8	0.547	0.45	0.52	0.547	50.2	0.58	0.598
KStar	50.25	0.564	0.459	0.561	0.564	49.75	0.48	0.654
NNge	51.2	0.655	0.5	0.54	0.655	48.8	0.49	0.655
PART	49.99	0.652	0.55	0.65	0.652	50.01	0.5	0.654
Simple Cart	60.16	0.6	0.49	0.682	0.6	39.84	0.47	0.654
Multi Layer Perception	61.58	0.546	0.5	0.855	0.546	38.42	0.495	0.356
Proposed	68.8	0.65	0.545	0.42	0.65	31.2	0.301	0.212

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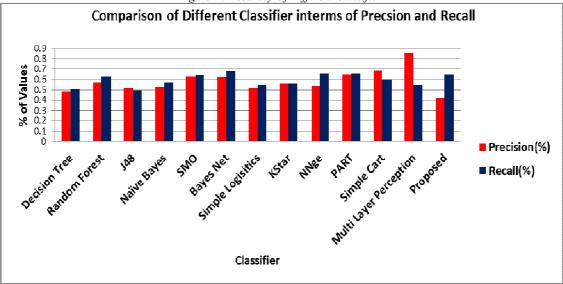


Figure 2 : Comparison Of Different Classifier In Terms Of Precision And Recall Values

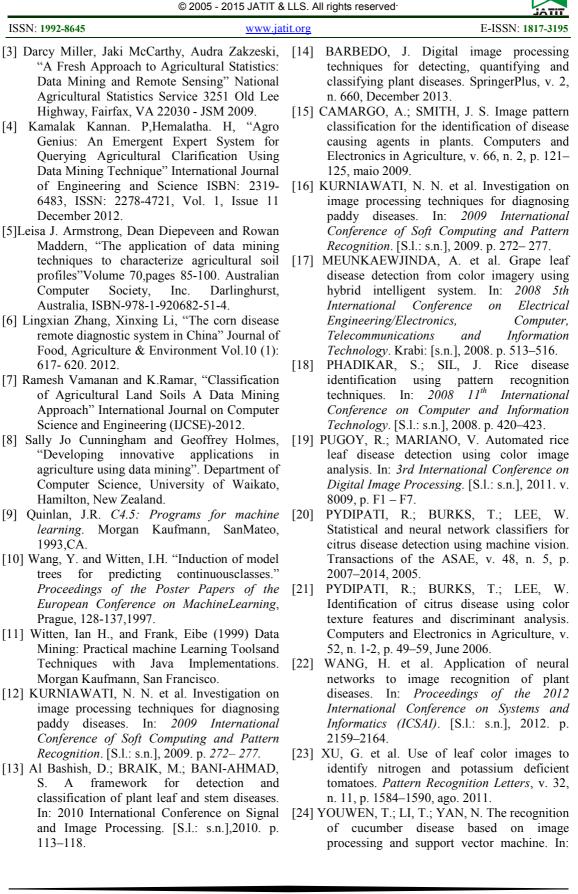
6. CONCLUSION

In the work, an automated system has been [1] Ashok Kumar. D,Kannathasan. N, "A Survey on developed for detecting different types of paddy crop diseases. In the first stage, uninfected and the diseased leaves are classified based on the number of peaks in the histogram. Miss classification may occur due to shadow effect and color distortion of aging leaves. To compare the other, our classic [2] Chandrakanth. Biradar, Chatura S Nigudgi, "An apriori algorithm is better performance.

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