



# DATA MINING APPROACH TO PREDICT DENGUE FEVER

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**Abstract:** Data mining approaches are used everywhere in the industry. It permits users to analyze data from many different perspectives. It is also used to classify and build relationships among data. So it is widely used in today's world in every field. It has given very fruitful results in healthcare field also. Many diseases such as High BP, sugar, dengue, COVID, heart attacks etc are taking the lives of community. In this paper dengue data has been analyzed using data mining techniques. Association and classification techniques have been applied to predict dengue fever. The results generated by these techniques will predict the dengue fever in individuals and they can consult doctor and take necessary medicine.

**Keywords:** Data Mining; Classification; Association; Dengue; Prediction; Medical Science.

## 1. INTRODUCTION

The process of data mining analyzes data from different dimensions and summarizing it into useful information. Basically data mining is used to analyze data. It permits users to analyze data from many different perspectives. Scientifically, data mining is a process for finding out an interconnection or patterns in huge amount of relational databases. It is a process of arranging data to find out patterns and identifying relationships. Data mining techniques can be executed quickly on real software and hardware platforms to improve the value of current information resources. These patterns cannot be determined by using traditional methods because of the bulkiness of data.

Data mining is one the important and motivating range of research with the equitable of discovery intelligible knowledge from large scale data sets [8]. Healthcare is most popular field application in data mining to identify hidden and admired knowledge in health data. Data mining is used in health industry to discover fraud in health insurance, treatment at lower cost, cure the disease [9].

### 1.1 Association

Association rule is one of important aspect of data mining. Association rule mining is used to predict correlations, patterns, associations or causal structures from data sets found in several types of databases such as relational databases, transactional databases, and other forms of data repositories. Given set of transactions, the goal of association rule mining is to find the rules which permit us to predict the occurrence of specific item depend on occurrences of the another items in the transaction. In many application areas it is required to discover the frequent patterns, frequent patterns are those data items that frequently occur in the data set. Classical problem of association data mining is super market analysis. Super market contains huge amount of data of their customers and it is required to maintain the data in the well-mannered form.

The association rule mining on "Market Basket Data" is Boolean association rule mining in which only Boolean values are considered. The aim of the association rule mining is to extract interesting patterns in the large datasets of items. Association rules are often used to analyze sales transactions. For example, it might be noted that customers who buy cereal at the grocery store often buy milk at the same time. In fact, association analysis might find that 85% of the checkout sessions that include cereal also include milk. This relationship could be formulated as the following rule: Cereal implies milk with 85% confidence



1.2 Classification

Classification consists of predicting a certain outcome based on a given input. In order to predict the outcome, the algorithm processes a training set containing a set of attributes and the respective outcome, usually called goal or prediction attribute. The algorithm tries to discover relationships between the attributes that would make it possible to predict the outcome. Next the algorithm is given a data set not seen before, called prediction set, which contains the same set of attributes, except for the prediction attribute – not yet known. The algorithm analyses the input and produces a prediction. The prediction accuracy defines how “good” the algorithm is. For example, in a medical database the training set would have relevant patient information recorded previously, where the prediction attribute is whether or not the patient had a heart problem.

Training set

Age	Heart rate	Blood pressure	Heart problem
65	78	150/70	Yes
37	83	112/76	No
71	67	108/65	No

Prediction set

Age	Heart rate	Blood pressure	Heart problem
43	98	147/89	?
65	58	106/63	?
84	77	150/65	?

Among several types of knowledge representation present in the literature, classification normally uses prediction rules to express knowledge. Prediction rules are expressed in the form of IF-THEN rules, where the antecedent (IF part) consists of a conjunction of conditions and the rule consequent (THEN part) predicts a certain predictions attribute value for an item that satisfies the antecedent. Using the example above, a rule predicting the first row in the training set may be represented as following:

**IF (Age=65 AND Heart rate>70) OR (Age>60 AND Blood pressure>140/70) THEN Heart problem=yes**

Knowledge Discovering in Databases is a process of discover knowledge in data and recap the “high-level” application of specific data mining technique. It is used for relevance interest to researcher’s awareness, databases, specific, artificial intelligence and data visualization. The unified aim of the KDD process is to abstract data from data warehouse in the framework of huge databases.

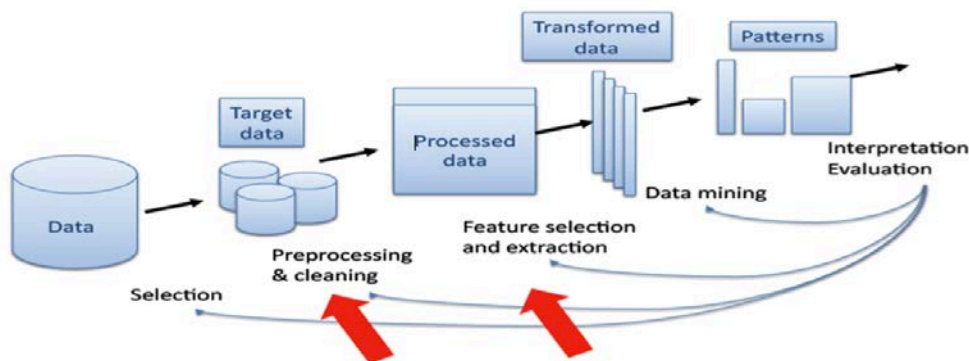


Figure-1: Knowledge Discovering in Databases (KDD)



### 1.3 Dengue Fever

Dengue is a dangerous disease. Dengue fever is one of the important reasons for the increasing deaths globally. It is a mosquito borne life threatening disease caused by breeding of Aedes mosquito. In India numbers of deaths are caused by dengue disease. Dengue is a dangerous disease with concern that is widespread in tropical and sub-tropical countries. It is a disease caused by breeding of Aedes Mosquito and also named as break bone fever. Dengue disease infected by Dengue Hemorrhagic Fever (DHF) whose symptoms including breeding, low level of blood platelets, low blood pressure, metallic taste in mouth, headache, muscle and joint pain and rashes etc. Dengue fever is painful and life threatening mosquito borne disease caused by female mosquitoes. Dengue virus is spread by Aedes mosquito. The mosquito becomes affected when it bites a person with dengue virus in their blood and it can be breed to other normal person. There is no particular medicine or antibiotic convenient to treat it. Dengue fever occurs in form of cycles and this cycle is present inside the body of an affected person for two weeks or less than two weeks. The various symptoms that are associated with dengue disease are:

- Fever Temperature
- Period of Fever
- Muscle and Joint Pain
- Rashes
- Low Heart Rate
- Blood Pressure
- Pain behind Eyes
- Headache

These various symptoms indicate serious illness and are sign that we must consult a doctor.

## 2. LITERATURE REVIEW

This section reviews the current research papers and articles to present the existing knowledge on this particular topic.

**Mani Shankar, Mayank Pahadia, Divyang Srivastava, Ashwin T S, G. Ram Mohan Reddy (2015) A Novel Method for Disease Recognition and Cure Time Prediction Based on Symptoms:** The study of this paper describes a novel model for identification of disease and prediction of their cure time by considering only the symptoms, they allow different coefficients for each symptoms of a disease and filter out a dataset with rating assigned to each symptom by the user. The disease get identify based on calculate numerical value using reinforcement learning. Algorithm used into account the similarly between the symptoms of one user and other user how have suffered from the same disease. For implementing this approach recommender system was used to read an input to find out pattern, which is based on the dataset given to train the system. This paper provides the solution that takes the symptoms with rating given by the patient to predict the possible disease and possible cure time of the disease.

**Kashish Ara Shakil, Shadma Anis and Mansaf Alam (2017) Dengue Disease Prediction Using Weka Data Mining Tool:** Study of this paper describes dengue is Dangerous disease. It is caused by Aedes mosquito. In this paper, they use available dataset for dengue including the symptoms like fever temperature, WBC, platelets, severe headache, vomiting metallic taste, joint pain, appetite, diarrhea, hematocrit, haemoglobin. This paper discusses various algorithm approaches available for data mining that have been used for the predication of dengue disease. Data mining is a well-known technique used for classification of diseases such as dengue, diabetics and cancer by many health organizations. The proposed approach used WEKA for evaluation. It was concluded that naïve bayes and J48 are the best performance algorithms for classification because they achieve maximum accuracy (100%) with 99 correctly classified instances and least mean absolute error and it takes least time foe building this model through explorer and knowledge flow results.

**P.Manivannan, Dr. P. Isakki Devi (2017) Dengue Fever Prediction using K-Medoid Clustering Algorithm:** The paper describes about dengue is a life threatening disease infection by female mosquitoes. It is searched in hot regions. This paper include the symptoms of dengue like severs joints pains, headache, rashes, thrombocytopenia, leucopenia and muscle ache. It is also called break bone fever because it is due to muscle ache and joint pains. This paper, they include four levels namely preprocessing, attribute selection, k-medoid clustering and prediction of dengue



fever by using of dengue fever by using R 3.3.2 tool for preprocessing the household of dengue dataset. The aim of this paper, they predict those people they affected by dengue categorization of age by using k-medoid clustering algorithm, which has been implemented. The k-medoid algorithm is just like k-means clustering algorithm. The proposed algorithm is used for systematic method of selecting the initial medoids. The performance of algorithm of algorithm is varying on initial medoids. K-medoid clustering is improving the efficiency of output.

**V.Krishnaiah, Dr.G.narsimha, Dr.N. Subhash Chandra (2018) Diagnosis of Lung Cancer Prediction System Using Data Mining Classification Techniques:** The paper describes the cancer is most considerable cause of death for both man and women. Initial detection of cancer can be beneficial in curing the disease completely. Preliminary of techniques to discover the appearance of cancer module in initial stage is expanding. Commonly lung cancer disease is misdiagnosed. Previous diagnosis of lung cancer saves immense lives, failing may lead to other severe headache causing abruptly fatal end. Early detection depends on the cure rate and diagnosis of the disease. In this paper they include following symptoms dyspnea, hemoptysis, chronic coughing or change in regular coughing pattern, wheezing, chest pain or pain in abdomen, cachexia, dysphonia, clubbing of the finger nails, dysphasia, pain in shoulder, chest, arm, bronchitis or pneumonia, decline in health and unexplained weight loss. Data mining and knowledge discovery of database is diverse applications in business and scientific domain. The main purpose of this paper to proposed a model for early detection and correct diagnosis of a disease and they will consult doctor for saving life of the patient.

**Nandini. V and Sriranjitha. R and Yazhini. T. P (2019) Dengue Detection and Prediction System Using Data Mining with Frequency Analysis:** In this paper author(s) defines clinical documents and conditions of patients. Dengue is widespread disease in today's world. This main focus of this paper is performing named entity recognition to extract disorder mentions, time expressions and other related characteristic from clinical data. They proposed a model, which predict about presence or absence of dengue disease and analysis of relationship of occurrence with consider symptoms. The proposed model produces accuracy and estimate tool for medical experts. The proposed system is effective for medical repositories. POS tagging is used for tag the discharge detail of the patient.

**Shameem Fathima, Nisar Hundewale(2019) Comparison of Classification Techniques-SVM and Naives Bayes to predict the Arboviral Disease-Dengue:** In this paper, author(s) define the analysis of several data mining techniques to predict the arboviral disease dengue. It analyzes the real dataset from super specialty hospitals diagnostic laboratories where blood samples were collected. The data set contain 5000 records with 29 parameters. This paper includes data mining techniques SVM and Naïve bayes classifier and evaluates the comparison results between the method by using dengue disease diagnosis. They collect the data record forms in Chennai and tirunelveli from India.

**N.Subitha, Dr.A.Padmapriya (2020) Diagnosis for Dengue Fever Using Spatial Data Mining:** In this paper they describe spatial data and used by accurate method for rule mining. Basically association rule mining is used for finding interestingness patterns from spatial database by using K-means algorithm in which patterns are not exactly stored. Spatial association rule mining is used to calculate several spatial correlations among a huge number of spatial objects. Finding interesting optimization method is called progressive refinement. It adopted by spatial association analysis. In this paper, we have implementation the automatic dengue test kit system to reorganization the dengue fever by using microscope blood image report. Microscope image report as input and signals are filtered and the feature characteristics are extracted by neural networks. Classification and back propagation network gives 98% correct results in short time.

**Nor Shafikah Roslan, Zulkiflee Abd Latif, and Nazri Che Dom (2020) Dengue Cases Distribution Based in Land Surface Temperature and Elevation:** It describes Malaysia is one of the countries that dengue fever crisis. The dengue virus in Malaysia remains high. Cure of dengue is unavailable. In this paper, they use association between land surface temperature, elevation and study of dengue fever. Spatial application is geographical information system and remote sensing image. These applications are used under control of dengue outbreaks as analysis of potentially high risk areas in urban areas. Dengue dataset can disturbed in patterns and factors are visualized. In this paper, it includes the range of temperature from 14-18 degree C at lower level and 35-40 degree C at the upper level. Dengue fever is spread between 10 degree C and not more than 30 degree temperature. It includes four classes of temperature 12 degree C to above 30 degree C. dengue incidence was distributed results at low elevation. It emphasized lower elevation less than 500 meter. It was found dengue is occurring in low land area.

### 3. RESULTS AND ANALYSIS

Various responses from 100 Patients have been collected according the questionnaire framed. The data has been collected from patients suffering and not suffering from dengue disease. The collected response was analyzed on



various parameters as presented below. In order to carry out an implementation R Studio is used as data mining tool. The main aim of the research is to predict the disease based on symptom.

The following are the screenshots to illustrate the results produced from association rule mining in R studio.

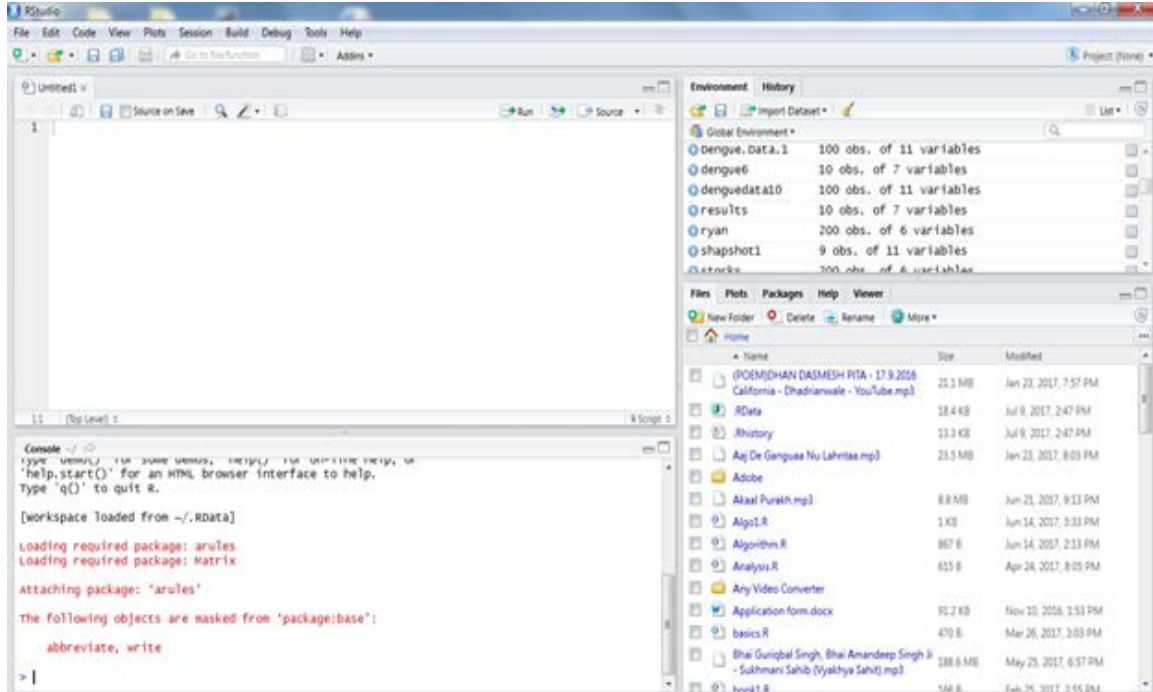


Figure-2: R studio tool window.

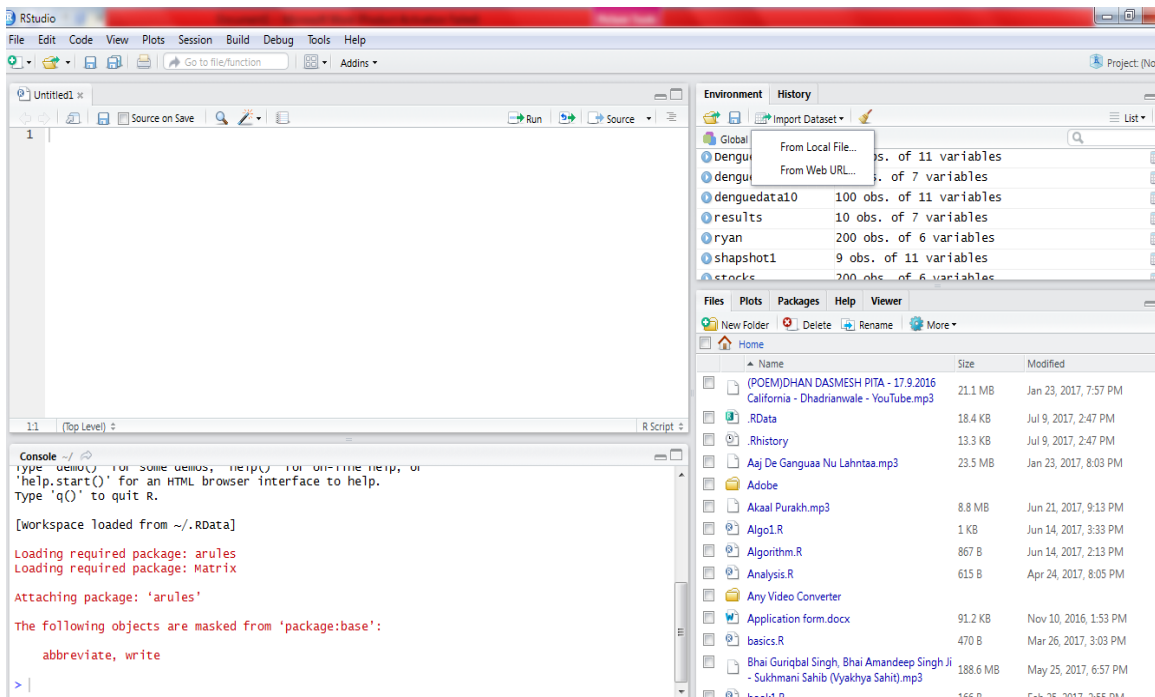


Figure-3: Import database file into R studio.

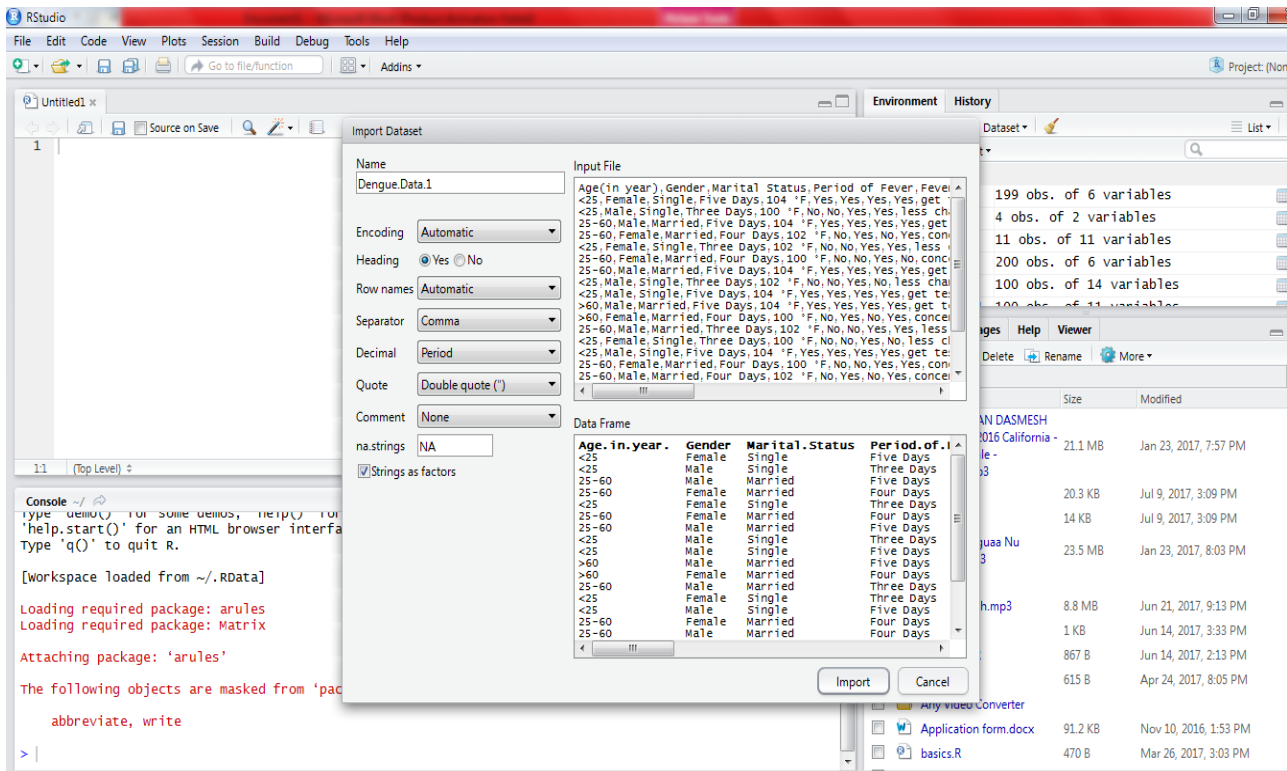


Figure-4: Import window of R studio.

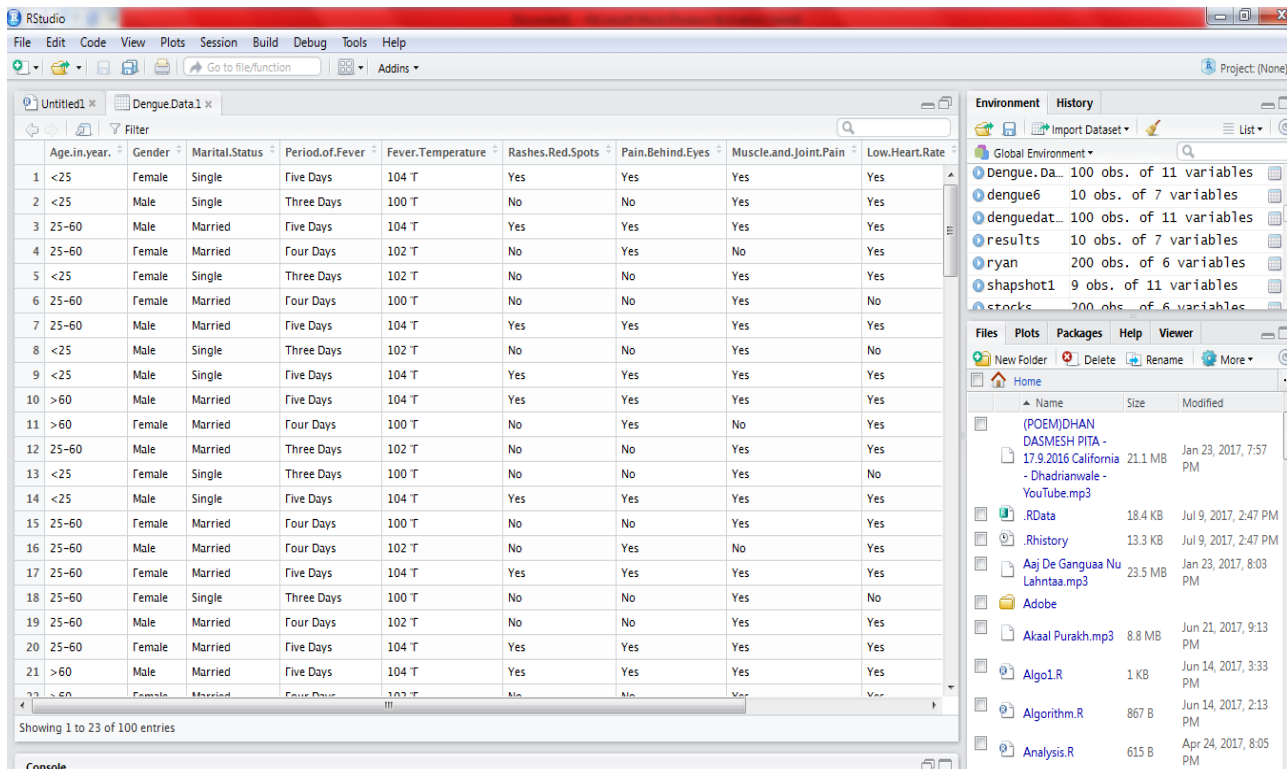


Figure-5: Database with number of rows, columns and attributes.



- ❖ Use apriori algorithm with support = 0.2, confidence = 0.5, minimum length = 4, it shows values of parameters and results in 108 rules in following Figure-5 and Figure-7..

```

Apriori

Parameter specification:
confidence minval smax arem aval originalsupport maxtime support minlen maxlen target ext
0.5 0.1 1 none FALSE TRUE 5 0.2 4 10 rules FALSE

Algorithmic control:
filter tree heap memopt load sort verbose
0.1 TRUE TRUE FALSE TRUE 2 TRUE

Absolute minimum support count: 20

set item appearances ... [2 item(s)] done [0.00s].
set transactions ... [19 item(s), 100 transaction(s)] done [0.00s].
sorting and recoding items ... [18 item(s)] done [0.00s].
creating transaction tree ... done [0.00s].
checking subsets of size 1 2 3 4 5 6 7 8 done [0.00s].
writing ... [108 rule(s)] done [0.00s].
creating s4 object ... done [0.00s].

```

Figure-6

lhs	rhs	support	confidence	lift
[1] {Rashes.Red.Spots=Yes, Pain.Behind.Eyes=Yes, Low.Heart.Rate=Yes}	=> {DengueStatus=positive}	0.52	1.0000000	1.923077
[2] {Rashes.Red.Spots=Yes, Muscle.and.Joint.Pain=Yes, Low.Heart.Rate=Yes}	=> {DengueStatus=positive}	0.52	0.9629630	1.851852
[3] {Rashes.Red.Spots=Yes, Pain.Behind.Eyes=Yes, Muscle.and.Joint.Pain=Yes}	=> {DengueStatus=positive}	0.52	0.9811321	1.886792
[4] {Pain.Behind.Eyes=Yes, Muscle.and.Joint.Pain=Yes, Low.Heart.Rate=Yes}	=> {DengueStatus=positive}	0.52	0.9629630	1.851852
[5] {Rashes.Red.Spots=Yes, Pain.Behind.Eyes=Yes, Muscle.and.Joint.Pain=Yes, Low.Heart.Rate=Yes}	=> {DengueStatus=positive}	0.52	1.0000000	1.923077
[6] {Period.of.Fever=Five.Days, Fever.Temperature=104 °F, FeverStatus=get tested}	=> {DengueStatus=positive}	0.51	1.0000000	1.923077
[7] {Fever.Temperature=104 °F, Rashes.Red.Spots=Yes, FeverStatus=get tested}	=> {DengueStatus=positive}	0.51	1.0000000	1.923077
[8] {Fever.Temperature=104 °F, Low.Heart.Rate=Yes, FeverStatus=get tested}	=> {DengueStatus=positive}	0.51	1.0000000	1.923077
[9] {Fever.Temperature=104 °F, Pain.Behind.Eyes=Yes, FeverStatus=get tested}	=> {DengueStatus=positive}	0.51	1.0000000	1.923077
[10] {Fever.Temperature=104 °F, Muscle.and.Joint.Pain=Yes, FeverStatus=get tested}	=> {DengueStatus=positive}	0.51	1.0000000	1.923077

Figure-7

- ❖ In Rule [1], if Rashes.Red.Spots=Yes, Pain.Behind.Eyes=Yes, Low.Heart.Rate=Yes then it predicts DengueStatus=positive with support=0.52 and confidence=1.00.
- ❖ If we increase the value of parameters with support = 0.3, confidence = 0.9, minimum length = 4, it shows results in 99 rules, that means order of rules get decreased. It is depicted in Figure-8.



```

Apriori

Parameter specification:
confidence minval smax arem aval originalSupport maxtime support minlen maxlen target ext
0.9 0.1 1 none FALSE TRUE 5 0.3 4 10 rules FALSE

Algorithmic control:
filter tree heap memopt load sort verbose
0.1 TRUE TRUE FALSE TRUE 2 TRUE

Absolute minimum support count: 30

set item appearances ...[2 item(s)] done [0.00s].
set transactions ...[19 item(s), 100 transaction(s)] done [0.00s].
sorting and recoding items ... [10 item(s)] done [0.00s].
creating transaction tree ... done [0.00s].
checking subsets of size 1 2 3 4 5 6 7 8 done [0.00s].
writing ... [99 rule(s)] done [0.00s].
creating s4 object ... done [0.00s].

```

Figure-8

- ❖ If we increase the value of parameters with support = 0.4, confidence = 1.0, minimum length = 4, it shows 96 rules as in Figure -9, that means order of rules get decreased.

```

Apriori

Parameter specification:
confidence minval smax arem aval originalSupport maxtime support minlen maxlen target ext
1 0.1 1 none FALSE TRUE 5 0.4 4 10 rules FALSE

Algorithmic control:
filter tree heap memopt load sort verbose
0.1 TRUE TRUE FALSE TRUE 2 TRUE

Absolute minimum support count: 40

set item appearances ...[2 item(s)] done [0.00s].
set transactions ...[26 item(s), 100 transaction(s)] done [0.02s].
sorting and recoding items ... [16 item(s)] done [0.00s].
creating transaction tree ... done [0.00s].
checking subsets of size 1 2 3 4 5 6 7 8 done [0.00s].
writing ... [96 rule(s)] done [0.00s].
creating s4 object ... done [0.00s].

```

Figure-9

The following figures predict the classification result in Tanagra tool.

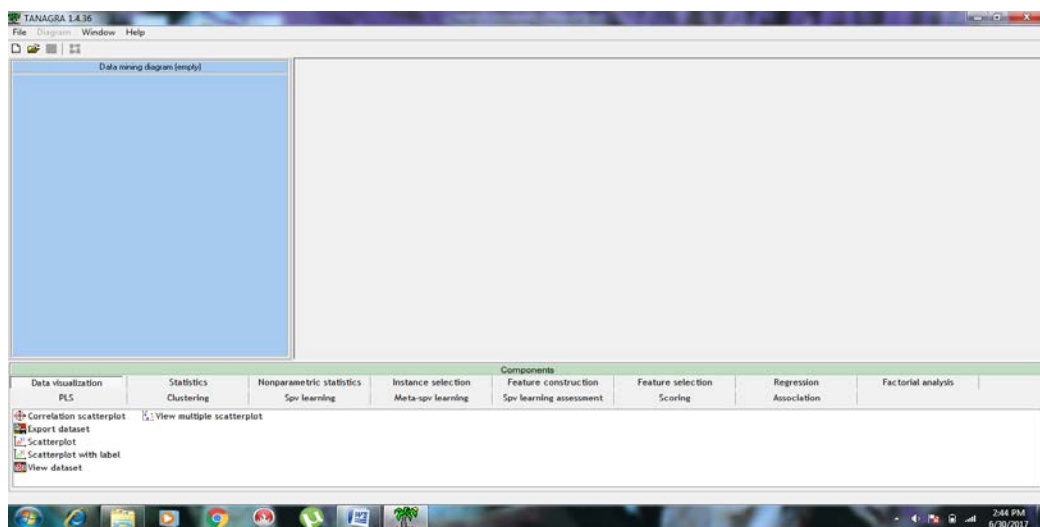


Figure-10 Tanagra tools window.



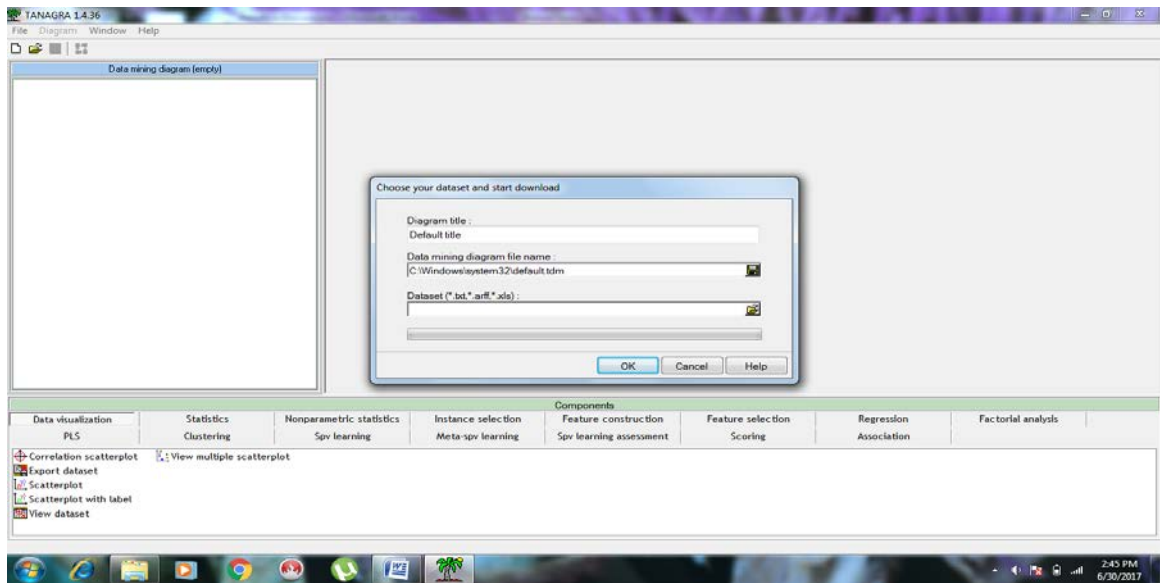


Figure-11 Import database file into tanagra.

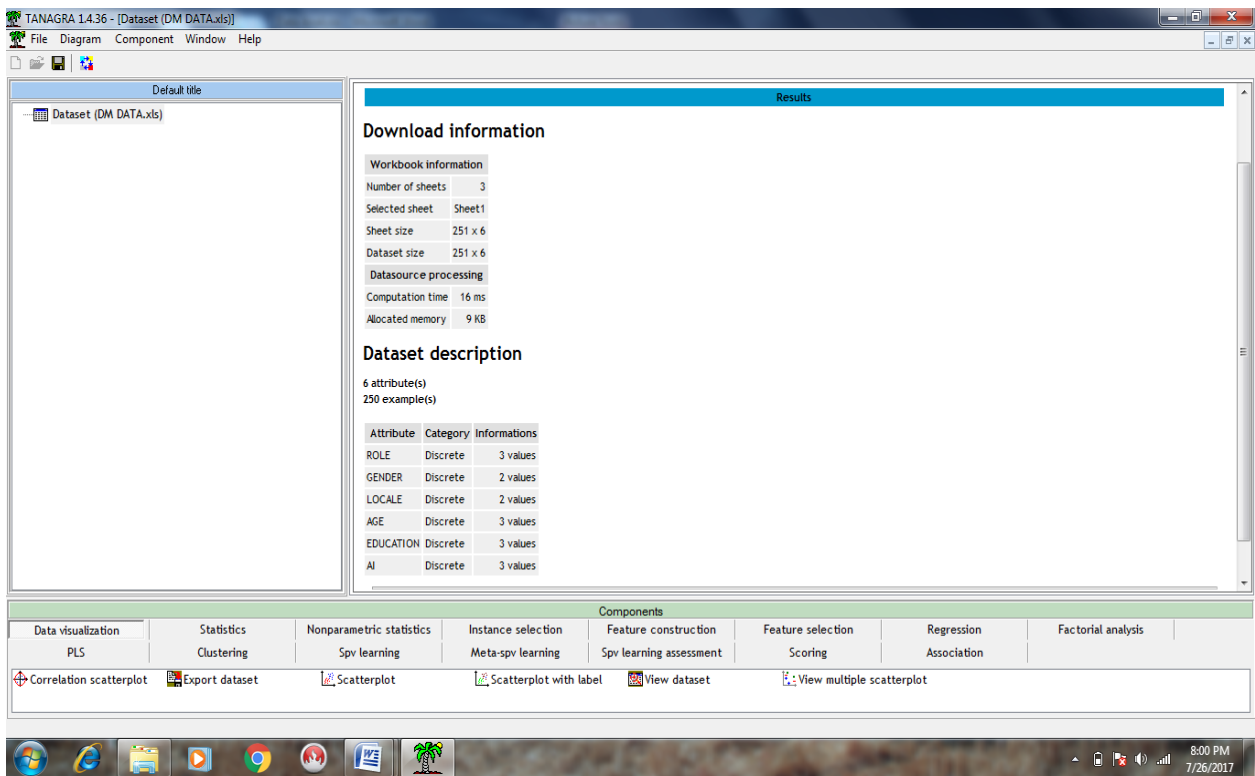


Figure-12 Dataset description

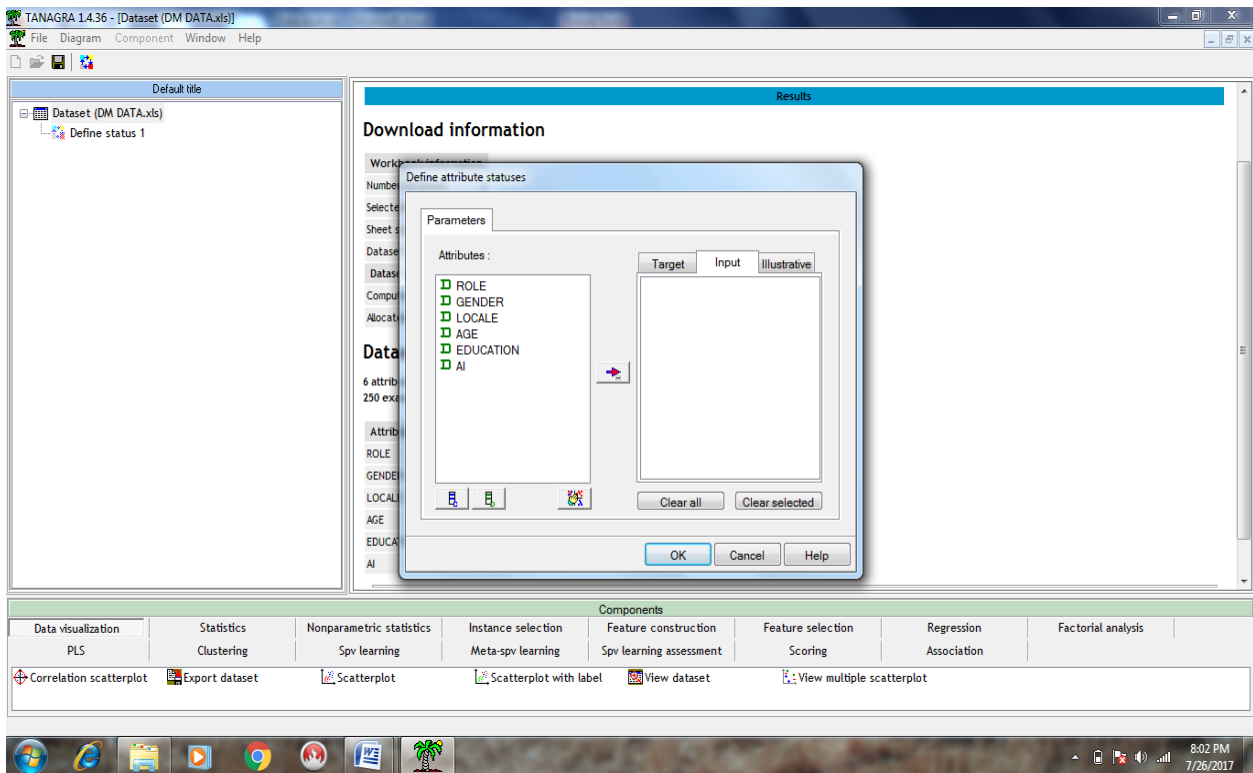


Figure-13 Define status on parameters

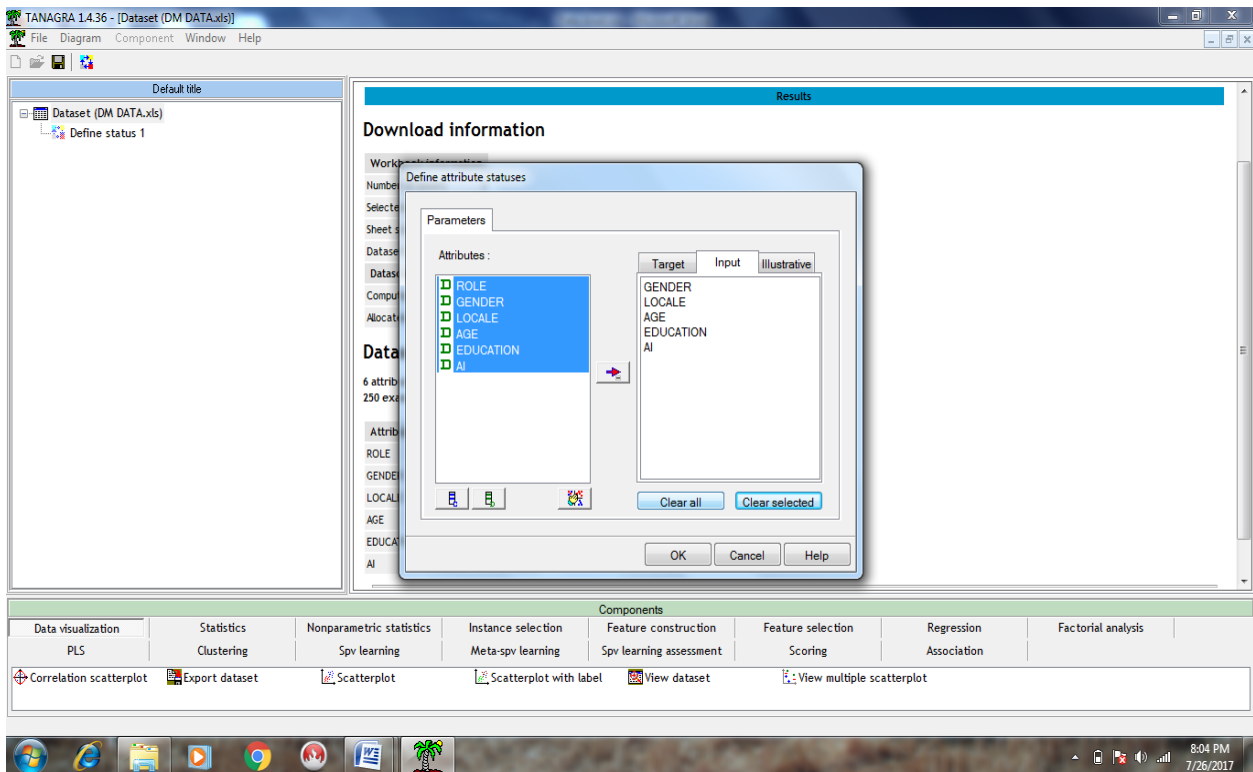


Figure-14 Setting up input values

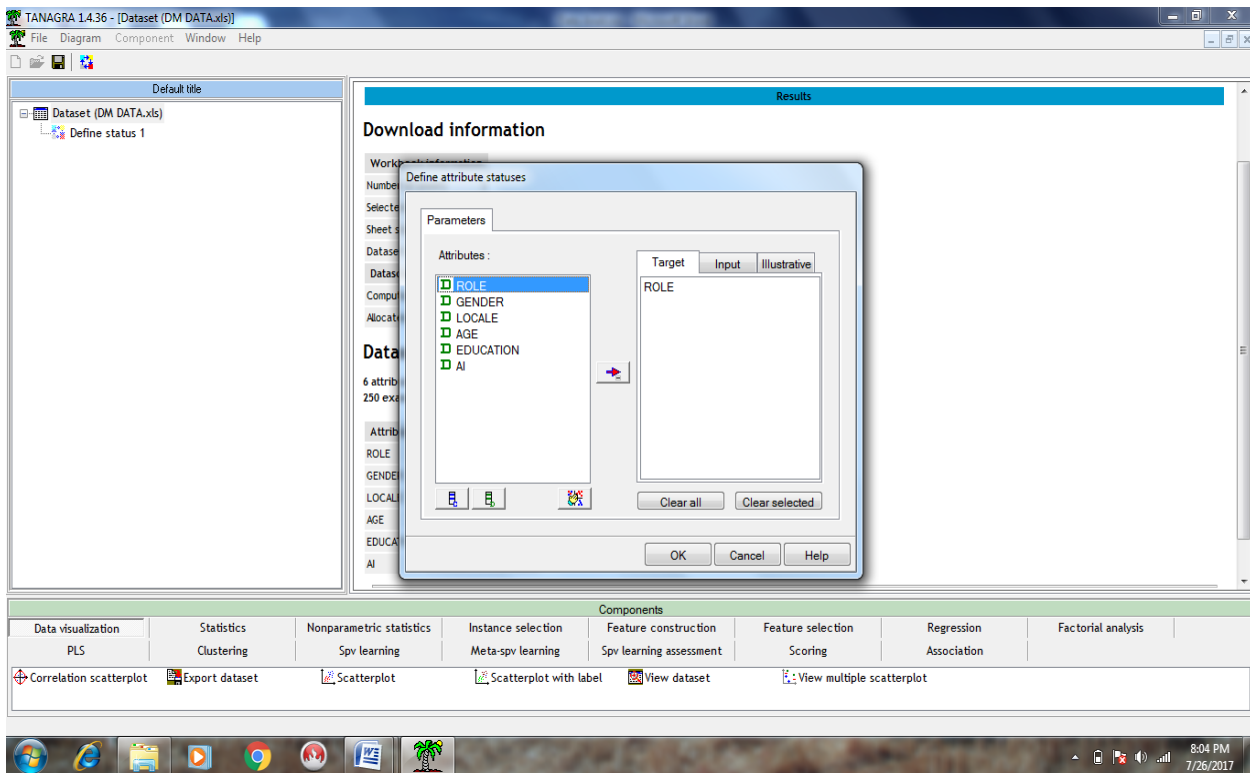


Figure-15 Setting up target values

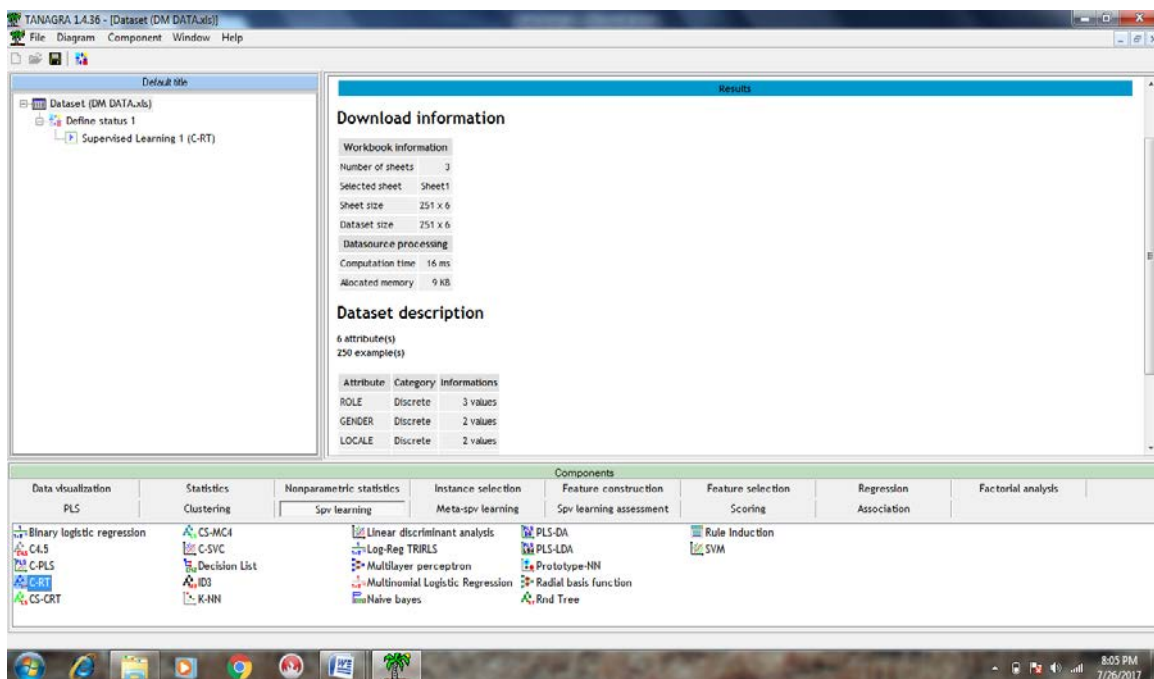


Figure-16 Supervised Learning C-RT

MIN SIZE OF NODE TO SPLIT means the minimum number of instances needed for performing a splitting of a node. For our dataset, we do not perform a split if there are less than 10 instances.

PRUNING SET SIZE means the part of the learning set used for the post pruning process. The default value is 33% i.e. 67% of the learning set is used for the growing phase (67% of 300 = 201 instances), 33% for the pruning phase (33% of 300 = 99 instances).



We have the sequences of trees table, with the number of leaves, the error rate calculated on the growing set and pruning set.

The error rate computed on the growing set decreases as the number of leaves increases. We cannot use this information to select the right model. We use the pruning sample to select the "best" model. The tree which minimizes the error rate on the pruning set is highlighted in green.

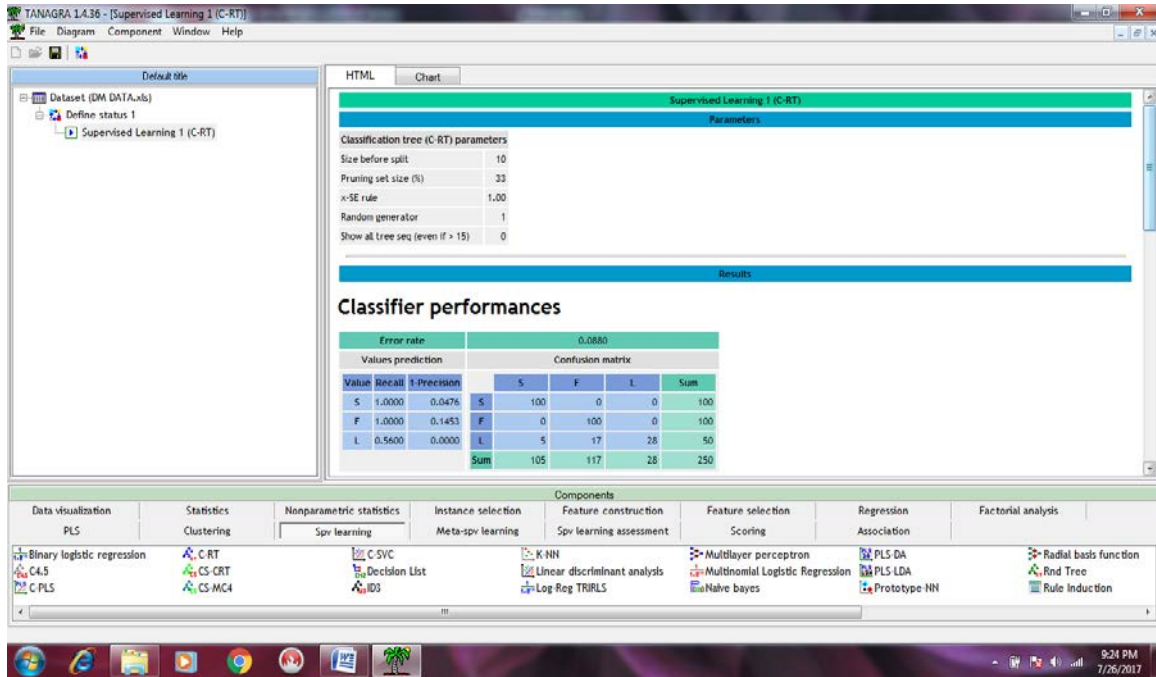


Figure-16 Classifier Performance

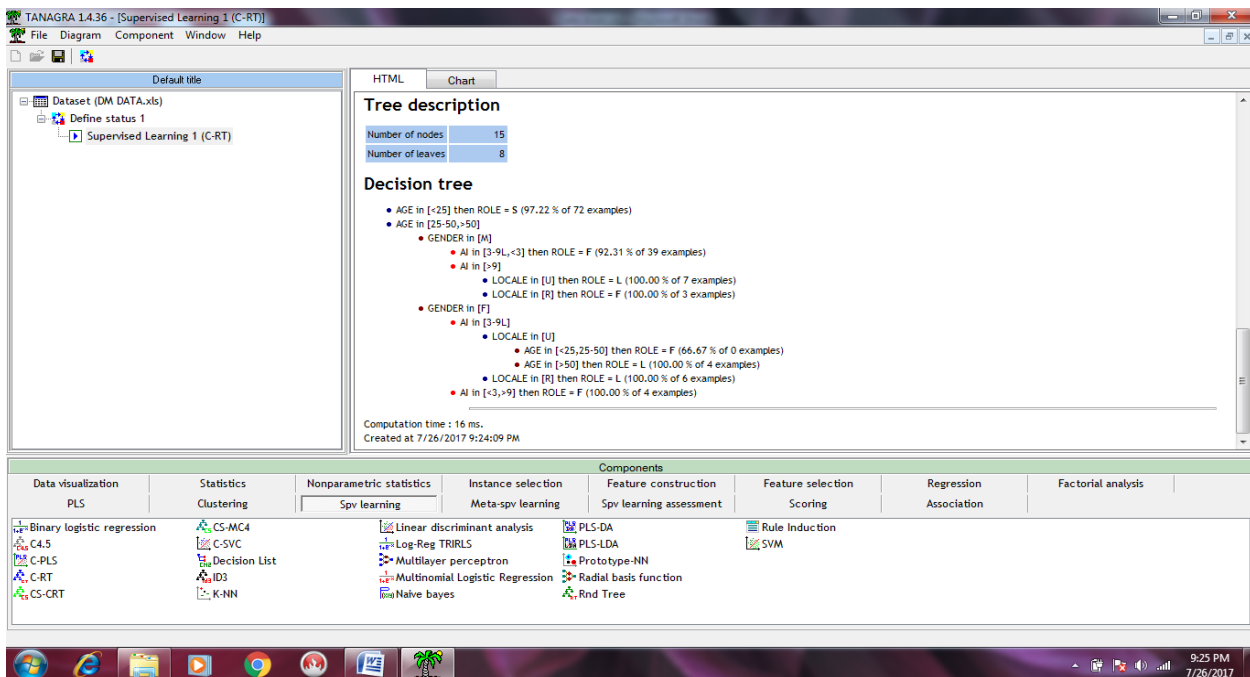


Figure-16 Tree Description



#### 4. CONCLUSION AND FUTURE SCOPE

##### Conclusion

- The prediction of dengue disease is carried out by observing various symptoms related to dengue disease.
- From the above results it is concluded that, the main focus of the research is on the association among symptoms of dengue disease like fever temperature, rashes, muscle and joint pain.
- The importance of knowledge identifies that disease.
- Dengue and from the observed symptoms it is analyzed that dengue fever mostly varies on fever temperature followed by Muscle and Joint pain and Rashes/red spots.
- Private hospital service to treat dengue fever is better than government hospital according to the concluded results.

##### Future Scope

The responses gathered here are limited to 100 patients but we increase the sample size by considering more responses to get more accurate results. More powerful tools such as neural network, fuzzy logic or the other data mining techniques can be applied.

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