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Machine Learning Approaches For Predicting Patient Severity Levels In T2dm Complications Neuropathy

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1. INTRODUCTION

The most common complication of both type 1 and 2 diabetes is diabetic neuropathy (DN), a type of nerve damage caused by long-term elevated glucose levels, which occurs in more than half of the individuals affected. Over the span of several decades, the disorder normally progresses progressively and occasionally. Many studies have found that the quality of life of those living with DPN has had a substantial negative effect. In comparison, a significantly improved Mortality for people afflicted with peripheral neuropathy (DPN) with diabetes who have undergone a major amputation. This needs decisive action to tackle this growing public health problem [1]. The research also uses the algorithms RF, SVM, KNN and NB to search for accuracy and indicates that in classifying the complexity of the T2DM, SVM has the high est accuracy [2]. Diabetes is a significant metabolic disease that can harmfully annoy patients. It may lead to multiple complications such as heart stroke, diabetic nephropathy and other conditions, if not diagnosed. In keeping a safe life, early diagnosis will help. As cases of diabetes are growing increasingly, these diseases are a cause for global concern. This research also uses the PIMA database and monitors the trend to identify patterns using R-Supervised MLAs-Linear Kernel SVM, RBF-Kernel SVM, KNN, ANN and Reduction of Multifactor Dimensionality (MDR) [3] [4]. The aim of this analysis was to compare the outputs of SVM, Naïve Net, Decision Stump, and Proposed Ensemble process models of predicting Diabetes using common risk factors. The findings show that the Suggested Ensemble Approach worked well in terms of accuracy of classification [5]. The number of diabetic patients (characterised by hyperglycemia) in the world is predicted to exceed 642 million in 2024 and requires a great deal of care. So, it is important to use the effective methods to anticipate and recommend to medical practitioners the proper approach.

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2. RELATED WORKS

This study finding the correlation among those factors using heatmap visualization techniques in Machine Learning. Many researchers have conducting analysis on the performance of various algorithms and similar approaches to find potential gaps in this area of analysis. This chapter summarises several important works. First of all to make the data correct, relevant and accessible, it is most important to specify the parameters that will assist the study in determining the source of the DPN. Of the 20 features recorded for each patient, 13 were found to impact the progress of the DM patient towards DPN. Age, type of diabetes, degree of education, BMI, blood pressure history, systolic blood pressure, foot ulcer history, drug, weight, history of laser photocoagulation, length, average blood glucose and height were the 13 characteristics identified. **[6, 7]**.

Significant Risk Factors associated with DPN	References
Age	[8-11]
BMI (obesity)	[8][13][19]
Blood Glucose	[8][18][19]
HbA1c	[14][18]
Triglycerides	[19-21]
Blood Pressure	[22-24]
Urea Serum Creatinine	[25, 26]
GFR][27,28]
Hyperglycemia	[29-32]
Microalbuminuria	[33-37]
DiabetesMellitus and its duration of years	[10][38-40]

The following Table 1 depicts the various significant factors associated with DPN.

Table 1: Significant Risk Factors associated with DPN along with references

• Age and risk of DPN

The impact of age on diabetic peripheral neuropathy has been described in [8-11] all of the article's reports and they have shown revealed substantial age gaps through the univariate analysis and the multivariate analysis.

• BMI (Obesity) and risk of DPN

The effects of BMI on DPN has been analysed in all the studies **[8][13][19].** Obesity has been described as a risk factor in DPN in the Southern German population. Obesity and the existence of at least 2 cardiovascular risk factors (triglycerides or plasma glucose, decreased HDL, increased waist circumference, hypertension) in the general US population aged 40 years or older raise the possibility of peripheral neuropathy (OR: 2.20, 95 percent CI: 1.43-3.39).

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• Blood Glucose and risk of DPN

Fasting plasma glucose (FPG) **[8][18][19]** High blood sugar (glucose) can injure nerves throughout your body. The results of the studies have been established a syndromic link between neuropathy associated with impaired glucose tolerance and the sensory-predominant neuropathy commonly observed in early diabetes.

• HbA1c and Risk of DPN

The results of the Nisar et al and Fargol Booya et al., [16][17] studies confirm reports regarding the association of neuropathy with glycemic control (HbA1C), age, BMI and duration of disease.

• Triglycerides and Risk of DPN

The studies **[19][21]** have shown that the presence of triglycerides in diabetics' normal blood functions raises the risk of neurological damage.

• Blood Pressure and Risk of DPN

Federica Di Gennaro et al., and Vincenza Spallone assesses the clinical correlates of Morning Blood Pressure Surge (MBPS) in a diabetic population, with particular attention paid to diabetic complications and diabetic neuropathy. Also found that the framework highlights the relevance of assessing BP variability in people with diabetes. These research reports confirm that vascular disease is associated with increased MBPS diabetes in the population, diastolic blood pressure as well as cardiovascular autoimmune disease (CAN) [22][24]

• Urea Serum Creatinine and Risk of DPN

Diabetic patients with high urea levels, high cholesterol and triglycerides levels should therefore be considered at higher risk for peripheral neuropathy [25]. The researchers found that if the incidence of DPN developed, abnormal lipid profile, elevated urea and reduced RBC levels point to the coexistence of cardiovascular and renal comorbid conditions [26].

• GFR and Risk of DPN

The variety of neurologic conditions affecting the central nervous system and the peripheral nervous system are caused by chronic renal failure. The kidneys do not perform well as they should, frequently due to a mismatch of salts and chemicals. The risks of damages to the peripheral nerve may be raised by this mismatch, leading to neuropathy. The research was exploring the relationship between the Glomerular Filtration Rate (GFR) and microvascular complications in type II diabetes mellitus such as diabetic neuropathy and nephropathy Patients of (DM) [27,28]

• Hyperglycemia and risk of DPN

The central nervous system (CNS) damage related to hyperglycemia. The obvious hyperglycemia present in diabetes can development of abnormalities and it has been discussed the evidence for insulinopenia in type 1 and insulin resistance in type 2 diabetes as causal factors in the development of DPN. Moreover, data suggest that other factors may also

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contribute. The study suggested that these two cases actually represent a single cause of impaired insulin signalling [29-32].

• Microalbuminuria and risk of DPN

Diabetic neuropathy observed in this study have been using distal symmetrical sensory motor neuropathy. This design of this study revealed that microalbuminuria can be an essential cause for microvascular complications in type 2 diabetes. The authors concluded that the incidence of microalbuminuria in patients with type-2 diabetes is considered significant. More it also indicates an early stage of diabetic nephropathy. Therefore, strict control of hyperglycemia and hypertension should deal early. It helps to prevent the other host of diabetes related complications occurring in future. Hence, it is clear that microalbuminuria is significantly related to the neurological presence [33-37].

• Duration of Diabetes and Risk of DPN

The research framework [10] [38-40] assessed Diabetic neuropathy is strongly associated with duration of diabetes and other factors such as HbA1c and GFR.

• Heatmap Visualization Technique

An expansion to the Heat Map has been presented in this work. Algorithm Based Feature Selection. This facilitates the automated collection of threshold parameters that help to enhance the efficiency of generalisation of high-dimensional data such as mass spectrometry. Using several cancer datasets, the authors have conducted a comparative analysis and compared the well-known Recursive Feature Elimination algorithm. The findings showed better efficiency of the grouping, which is very comparable with other related tests. Because of these results, it is possible to create datasets of visual images that are mapped to the original space by generating a heatmap representation of the data, and this will aid in the quest for regions of interest, not just from the point of view of the feature selection, but also for instance selection [41]. The authors illustrated the efficiency of Complex Heatmap to easily expose trends and links between different sources. Four databases of information related to multidimensional genomic evidence in the real world [42]. A new local dimension reduction algorithm suggested by the researchers was used in feature discovery and extraction of features. This article concentrates on extracting the high-dimensional hidden potential structure inMicroarray data for various categories, and explanation and comprehension. The effects of the potential structure information provided with the heatmap graph are also used. The suggested algorithm for the exploration of gene co-expression and coregulation can be efficiently extended to microarray data analysis [43]. A novel research analysis, univariate feature selection, feature importance, and correlation matrix with heat map were applied to find the optimum data subset of erythematosquamous disease in three feature extraction techniques. For calculating model efficiency, four classification techniques are used: Gaussian Naïve Bayesian (NB), Decision Tree (DT), Support Vector Machine (SVM), and Random Forest. The stacking ensemble approach is then introduced to boost the model's predictive efficiency [4]. The heat map image that reveals 11 distinct proteins for HT carcinoma cells that assist in cell survival/apoptosis was considered in this study. Three kinds European Journal of Molecular & Clinical Medicine

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of feature selection techniques namely filter, wrapper, and embedded were used and obtain seven marker protiens. But only one of these protiens was used for classification using the KNN and SVM classifiers and yield the results from these protiens [44].

From the related works and the American Diabetes Association recommends aiming for the normal and abnormal range of blood sugar level, Hemoglobin levels, GFR etc which are shown in the following Table 2.

Diabetes Mellitus Factors	Ranges				
Diabetes Menitus Factors	Normal	Abnormal			
Blood Glucose	<140 mg/dL	>140 mg/dL			
Fasting Plasma Glucose (Pre- breakfast) 2-hours in an oral glucose tolerance test	70-130 mg/dL	>130 mg/dL			
Post breakfast Glucose	130-180 mg/dL	>180 mg/dL			
HbA1C	5.7 % mg/dL	>6.5 % mg/dL			
Age	30-90 (in years)				
BMI (Body Mass Index)	18-25 kg/m 2	>25 kg/m 2			
Triglycerides	<150 mg/dL	150-199 mg/dL			
Blood Pressure (BP)	130/80 mm hg	below or higher the normal range at risk			
Urea-serum-creatinine	0.6 to 1.4 mg/dL	>1.4 mg/dL			
Glomerular Filteration Rate (GFR)	Male: 97 to 137 mL/min Female: 88 to 128 mL/min	Male: >137 mL/min Female: > 128 mL/min			
Gender (Male/Female)					
hyperglycemia-Symptoms	No(0)	Yes(1)			
hypoglycemic_symptoms	No(0)	-1			
microalbuminuria	30 -299 mg	>=300 mg			
Diabetes Mellitus (Duration of	f >5 years & el				
Years)	<5 years with control	level			

 Table 2: Diabetes Mellitus risk factors ranges according to American Diabetes

 Association and through Review of various journals mentioned in the Related Works.

These research framework findings confirm that if their diabetes continues for a long time, patients will soon develop a neurological disorder and other diabetic complications.

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3. METHODOLOGY

The following diagrammatic representation figure 1 depicts an overview of a general system framework for proposed work

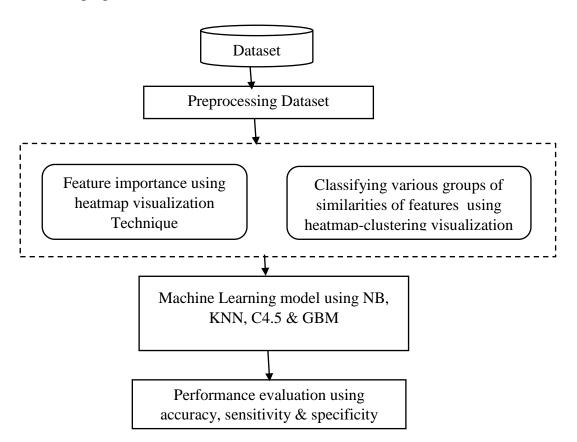


Figure 1. Flowchart of general system framework for proposed work

A. Feature Selection with Heatmap Visualization

Heatmap gives the opportunity to obtain useful information insights from the extensive datasets. It's the perfect method for data exploration strategy. The strong and weak correlations that can distinguish better using this. Simplify the exclusion of obsolete data as well. It may also be possible to do multicollinearity testing. The cell colour in the heatmap relates to the value. It is the best multivariate regression technique. The hue of the gradient allows the values in between to vanish, concentrating on the average [41]. When two variables are correlated, changes in their values have the same cause or an analogous explanation. The growth of one component may be the direct cause of a decrease of another factor in a negative correlation. A weak correlation means that there is a smaller chance of there being an association with the second variable as one variable increases or decreases. Labelling the frequency of the relationship, 0- 0.39 is known to be weak for absolute values of Pearson r, 0.40-0.69 to be mild or moderate, >0.7 to be high and 0.8-1to be a very strong correlation and the background of the findings should be considered. The negative correlated variables were not required for further analysis of the DPN forecast from the heatmap results

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of this analysis. Weak, average and high associations between DPN predictor factors were considered in this analysis. For example, from the figure 1, pre_breakfast_bloodGlucose and blood_Glucose, Microalbuminuria and urea-creatinine, GFR and urea-creatinine associations have been the strong chances the patients affected by DPN. Also, the correlations Triglycerides association with BMI, BMI connection with HbA1c were also moderately likely to increase the DPN.

Likewise, the interaction of BP with any BloodGlucose, Age and Triglycerides was rarely likely to raise the DPN. The table 3 summarizes the correlation features and its value that is associated with DPN provided by the heatmap graph.

-0.16 -0.59 0.09 -0.084 -0.01 -0.26 -0.1 -0.56 akfast_blood_g -0.18 -0.066 -0.16 -0.26 0.044 0.051 0.0077 -0.028 0.097 -0.033 hbalc 0.26 0.26 -0.33 0.0091 0.11 age -0.01 0.049 0.1 -0.32 -0.33 -0.049 -0.08 -0.14 -0.36 0.13 0.0091 -0.063 trialy -0.18 0.11 0.049 0.023 -0.006 -0.11 0.041 -0.33 -0.23 bp 0.044 -0.049 -0.023 urea serum creatinine 0.0 -0.006 -0.13 0.051 -0.086 -0.043 gfr -0.1 -0.043 -0.11 -0.13 -0.059 -0.084 -0.022 0.0077 -0.063 0.0023 0.068 -0.11 -0.03 -0.2 0.1 -0.22 -0.028 0.0023 -0.32 -0.33 0.22 -0.56 -0.36 -0.38 -0.64 -0.59 0.097 -0.4 -0.11 -0.14 0.041 0.11 0.088 -0.23 -0.011 -0.043 -0.38 righbalc ge Ē 8 ender blood gluco: preakfast blood R

The following figure 2 depict the Heatmap Visualization Graphs for DPN factors severity

Figure 2. DPN risk factors visualization using Heatmap

Correlation Features of DPN	Correlation	
	Values	
Pre_breakfast_bloodGlucose & Blood_Glucose	0.91	
Micro_albuminuria & Urea_Serum_Creatinine	0.89	
GFR & Urea_Serum_Creatinine	0.85	
Post_breakfast_bloodGlucose &	0.84	
Pre_breakfast_bloodGlucose		
GFR & Micro_albuminuria	0.81	
Post_breakfast_bloodGlucose & BloodGlucose	0.81	
Micro_albuminuria & Age	0.78	
Urea_Serum_Creatinine & Age	0.71	

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GFR & Age	0.63
Hyperglycemia & BloodGlucose	0.62
Hyperglycemia & Post-breakfast_bloodGlucose	0.61
Micro_albuminuria & Hyperglycemia	0.59
Hyperglycemia & urea_serum_creatinine	0.56
Hyperglycemia & GFR	0.56
Triglycerides & BMI	0.51
BMI & HbA1C	0.50
Age & Post_breakfast_bloodGlucose	0.50
DM_Duration & bloodGlucose	0.50
Hyperglycemia & Age	0.49
DM_Duration & Pre_breakfast_bloodGlucose	0.44
DM_Duration & Post_breakfast_bloodGlucose	0.43
Microalbuminuria & Post_breakfast_bloodGlucose	0.43
Hypoglycemia & Urea_serum_creatinine	0.39
Micrialbuminuria & pre-breakfast-bloodGlucose	0.38
Triglycerides & BloodGlucose	0.37
Age & Prebreakfast_bloodGlucose	0.36
Hypoglycemia & GFR	0.35
Triglycerides & Pre_breakfast_bloodGlucose	0.32
Microalbuminuria & Hypoglycemia	0.32
Microalbuminuria & Blood Glucose	0.32
BloodPressure & Post-breakfast_bloodGlucose	0.31
Urea_Serum_creatinine & Post-breakfast_bloodGlucose	0.31
Urea_Serum_creatinine & Pre-breakfast_bloodGlucose	0.29
GFR & Pre-breakfast_bloodGlucose	0.28
GFR & Triglycerides	0.28
Triglycerides & Post-breakfastGlucose	0.27
Age & BloodGlucose	0.26
DM_Duration & Triglycerides	0.24
BloodPressure& Triglycerides	0.19
BloodPressure & Pre_breakfast_BloodGlucose	0.18
BloodPressure & Blood Glucose	0.16
BloodPressure & Age	0.10
Table 3 Correlation and Heatman feature impor	11

Table 3. Correlation and Heatmap feature important table

B. Classification of DPN severity levels using Heatmap with Hierarchical Clustering

Heatmaps are related with hierarchical clustering as well. The rows or columns are organised based on similarity by hierarchical clustering. This makes comparisons convenient to see in the data. Heat maps also come with dendrograms. It is a diagrammatic representation the hierarchical association of objects [42, 43]. It is generated most commonly as an output from

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hierarchical clustering. This is used to find out the two parameters from the different variations are the most similar and merge them into a cluster.

The figure 3 depicts the pictorial visualization of variety of similarity levels of DPN factors. There are three clusters formed from the given DPN parameters. The clusters with dendrogram and its variety of colour bright-fading with values shown DPN factors that are high severity level group, moderate level group and lower-level affected group as it has been differentiated.

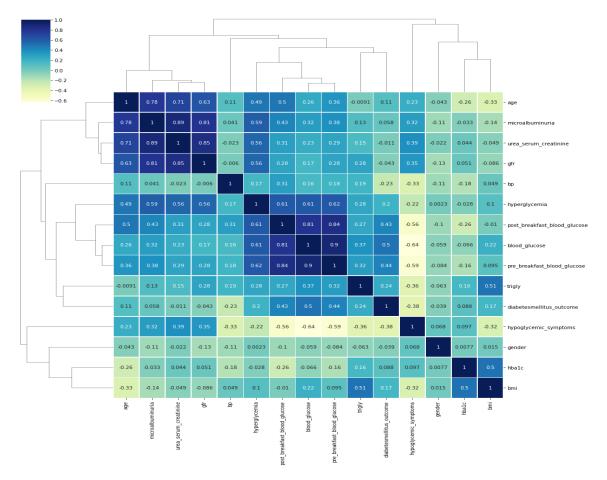


Figure 3. Various group of DPN Criticality level and its risk factors visualization using Cluster-Heatmap

The following table 4 shows a description of the very critical categories of features provided by the cluster of Heatmaps.

Mild -Level (0.1-0.39)		Moderate-Level (0.4-0.69)	Critical-Level (0.7-0.99)			
Hypoglycemia	&	GFR & Age	Pre_breakfast_bloodGlucose			
Urea_serum_creatinine			& Blood_Glucose			
Micrialbuminuria &	pre-	Hyperglycemia &	Micro_albuminuria &			
breakfast-bloodGlucose		BloodGlucose	Urea_Serum_Creatinine			
Triglycerides	&	Hyperglycemia & Post-	GFR &			
BloodGlucose		breakfast_bloodGlucose	Urea_Serum_Creatinine			
Age	&	Micro_albuminuria &	Post_breakfast_bloodGlucose			

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Prebreakfast_bloodGlucose	Hyperglycemia	&			
		Pre_breakfast_bloodGlucose			
Hypoglycemia & GFR	Hyperglycemia &	GFR & Micro_albuminuria			
	urea_serum_creatinine				
Triglycerides &	Hyperglycemia & GFR	Post_breakfast_bloodGlucose			
Pre_breakfast_bloodGlucose		& BloodGlucose			
Microalbuminuria &	Triglycerides & BMI	Micro_albuminuria & Age			
Hypoglycemia					
Microalbuminuria & Blood	BMI & HbA1C	Urea_Serum_Creatinine &			
Glucose		Age			
BloodPressure & Post-	Age &				
breakfast_bloodGlucose	Post_breakfast_bloodGlucose				
Urea_Serum_creatinine &	DM_Duration &				
Post-	bloodGlucose				
breakfast_bloodGlucose					
Urea_Serum_creatinine &	Hyperglycemia & Age				
Pre-breakfast_bloodGlucose					
GFR & Pre-	DM_Duration &				
breakfast_bloodGlucose	Pre_breakfast_bloodGlucose				
GFR & Triglycerides	DM_Duration &				
	Post_breakfast_bloodGlucose				
Triglycerides & Post-	Microalbuminuria &				
breakfastGlucose	Post_breakfast_bloodGlucose				
Age & BloodGlucose					
DM_Duration &					
Triglycerides					
BloodPressure&					
Triglycerides					
BloodPressure &					
Pre_breakfast_BloodGlucose					
BloodPressure & Blood					
Glucose					
BloodPressure & Age					

 Table 4. Summary of greatly important categories of features provided by the heatmap cluster.

C. Predictive Analytics Algorithms for DPN:

After selecting the best feature subset, we used various machine learning algorithms to build a predictive model. In this research, we used different algorithms such as Naïve Bayes, C4.5, K-Nearest Neighbor (KNN) and GBM using the programming language Python.

Naïve Bayes:

Naïve Bayes is a fast and eager learning classifier. By using it, forecasts can be made in real time. It helps to capture the complexity of the model in a principled manner by assessing the probabilities of the findings. By doing this, diagnosis and statistical problems can be overcome. It can measure probabilities by measuring the frequency and variations of values in the results [2] [4].

KNN:

It is categorised under the lazy prediction approach. On the basis of comparisons, the simplified methodology encourages new work to be coordinated. The training data was organised into this algorithm. Define k - the number of neighbours close by. Among Distance Away Training and samples. Inaccessibility calculation of the learning sections were grouped and the nearby neighbour was Focused on the minimum - the distance is fixed in the successive phase [44, 45].

GBM:

Gradient boosting is a form of boosting machine learning. It is based on the intuition that when paired with previous ones, the next best possible model minimises the total prediction error. The error is minimised by projections from the current model that are similar to its targets [46].

C4.5:

C4.5 Algorithm extended features of ID3 algorithm suggested by Ross Quinlan et.al al. [46]. The decision tree uses the same knowledge for training as ID3, which contains the learning function. The method of learning should be used for medical data diagnosis in order to predict the importance of the decision. C4.5 in each of the tree divisions nodes, Choose the most efficient data attribute value that separates the data evaluated into class-enriching subset data. The forest is created by organised awareness. To make the highest value attribute decision, the normalised data gain is chosen from the C4.5 tree of judgement [4].

4. RESULTS AND DISCUSSION

The proposed study on the Diabetes data collection obtained from different hospitals from 2017 to 2019 has been applied using the programming language of Python. The meaning of the role and correlations gained using the visualisation technique of Heatmap and are described in Figure 1 and Table 3. This feature-relevant heatmap methodology provides a clear picture of data analysis and interpretation of data to enhance the efficiency of the forecast. Larger values were represented by dark colour squares and smaller values by light colour squares. The feature importance determined by the intensity and shades of the colours. Shades of dark blue colour used for strong positive effect, whereas light blue shades are used for average effect and shades of yellow are used for negative effect. The clustered heatmap helpful to identify the various critical levels of DPN patients' conditions. However, the authors sometimes though, with the help of the Clinicians and scientific papers were consulted for the chosen attributes have been medical significance.

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After selecting the attributes based on the heatmap visualization technique, applied four ML classifiers namely NB, KNN, C4.5 and GBM. The algorithms are evaluated on the basis of the performance metrics accuracy, sensitivity and specificity and which are presented in Table 3. Table 3 shows that the highest accuracy achieved is 0.82 for NB, 0.72 yields for KNN and almost similar level accuracy achieved for C4.5 and GBM. From this could be concluded, the performance of the KNN, GBM and C4.5 algorithms was almost similar and Naïve Bayes performed better.

The following Table 5 depicts the performance measures accuracy, sensitivity and specificity of DPN prediction using different MLAs.

The DPN prediction performances using various Machine Learning algorithms											
NB KNN C4.5 GBM											
Accuracy	Sensitivity	Specificity	Accuracy	Sensitivity	Specificity	Accuracy	Sensitivity	Specificity	Accuracy	Sensitivity	Specificity
0.82	0.91	0.90	0.72	0.82	0.8 0	0.75	0.81	0.7 9	0.7 7	0.8 0	0.7 9

Table 5. The DPN prediction performances using various MLAs

Accuracy, sensitivity, and specificity are used to assess the performance of classifiers which are represented by the following equations (1-3):

Accuracy = samples correctly classified
total samples classified

$$= TP + TN / (TP + FP + FN + TN)$$
,
Sensitivity = samples correctly classified as positive
total positive samples in the dataset

$$= TP / (TP + FN)$$
eqn.(2)

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Specificity = samples correctly classified as negative total negative samples in the dataset eqn (3) = TN / (TN + FP)

5. CONCLUSION

A graphic guide to interpreting numeric values is given by heatmaps. They are easy to interpret, so it can be helpful for physicians to reasonably grasp the significance of the features. A better predictive model of health care treatment for people at different risk thresholds, such as low, moderate and high levels of rapid development of DPN, was provided by the clustering heatmap. In a heat map, the outcome of a hierarchical clustering calculation is seen as a dendrogram that can be thought of as a tree. Where they intersect, the lines form nodes that belong to a group. In the given dataset, dendrograms represent the distance or similarity between the parameters [42,43]. Diabetic patients with various classes of DPN severity thresholds and distances between latency and amplitude were seen in this study. In addition, it was possible to create a prediction model using machine learning techniques for the DPN patients. Classifiers accuracy proved the Machine Learning techniques Naïve Bayes, C4.5, KNN and GBM can be preferable to use for this study. The system has achieved an accuracy of 0.82, sensitivity of 0.91 and specificity of 0.90 using NB, an accuracy of 0.72, sensitivity of 0.82 and specificity of 0.80 using KNN, an accuracy of 0.75, sensitivity of 0.81 and specificity of 0.79 using C4.5 and an accuracy of 0.77, sensitivity of 0.80 and specificity of 0.79 using GBM. Other classifiers can also be used to enhance the performance of the proposed framework. In addition, this study encourages clinicians to rapidly identify patients with intensive critical conditions and enables doctors in some cases to reduce the different severity of patient categories.

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