

RESEARCH ARTICLE

Exploring the risk factors and clustering patterns of periodontitis in patients with different subtypes of diabetes through machine learning and cluster analysis

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ABSTRACT

Aim: To analyse the risk factors contributing to the prevalence of periodontitis among clusters of patients with diabetes and to examine the clustering patterns of clinical blood biochemical indicators.

Materials and methods: Data regarding clinical blood biochemical indicators and periodontitis prevalence among 1804 patients with diabetes were sourced from the National Health and Nutrition Examination Survey (NHANES) database spanning 2009 to 2014. A clinical prediction model for periodontitis risk in patients with diabetes was constructed via the XGBoost machine learning method. Furthermore, the relationships between diabetes patient clusters and periodontitis prevalence were investigated through consistent consensus clustering analysis.

Results: Seventeen clinical blood biochemical indicators emerged as superior predictors of periodontitis in patients with diabetes. Patients with diabetes were subsequently categorized into two subtypes: Cluster A presented a slightly lower periodontitis prevalence (74.80%), whereas Cluster B presented a higher prevalence risk (83.68%). Differences between the two groups were considered statistically significant at a *p* value of ≤ 0.05 . There was marked variability in the associations of different cluster characteristics with periodontitis prevalence.

Conclusions: Machine learning combined with consensus clustering analysis revealed a greater prevalence of periodontitis among patients with diabetes mellitus in Cluster B. This cluster was characterized by a smoking habit, a lower education level, a higher income-to-poverty ratio, and higher levels of albumin (ALB g/L) and alanine aminotransferase (ALT U/L).

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Introduction

Periodontitis, a chronic inflammatory disease caused by plaque microorganisms, is now the leading cause of tooth loss in adults and not only poses a serious socioeconomic burden but also interacts with a wide range of systemic diseases [1]. The relationship between periodontitis and diabetes has been widely reported [2, 3], with chronic periodontitis considered the sixth most common complication of diabetes [4–6]. A study by Dhir et al. revealed that the overall prevalence of periodontitis in patients with and without diabetes was 72% and 27%, respectively [7]. Diabetes increases people's risk of periodontitis by 34%, and the periodontal status of diabetic individuals is worse [8–10]. Additionally, diabetes can promote the development of periodontitis [11]. The current diagnosis of periodontitis usually relies on the detection of clinical parameters (alveolar bone loss and clinical attachment levels) [12,14]. However, before these parameters can be used to detect the disease, severe damage to the periodontal tissue may have already occurred [15]. Therefore, finding a simple, rapid, and reliable way to diagnose and monitor the development of periodontitis in patients with diabetes at an early stage and to conduct early

preventive and personalized interventions is highly clinically important.

Blood biochemical tests are not only routine medical check-ups but also provide a more intuitive and precise reference for diagnosing specific diseases. The analysis of some unique biomarkers present in saliva and blood can further help estimate the speed of disease occurrence and development and facilitate timely monitoring of the patient's condition by healthcare professionals [16–18]. There is limited research on the relationship between blood biochemical markers and periodontitis in people with diabetes, and new evidence suggests that the prevention and management of oral health problems in diabetic individuals can minimize the harmful effects of hyperglycemia, with significant benefits for adults with diabetes [19–21]. Therefore, a total of 50 clinical and blood biochemical factors that may be associated with the prevalence of periodontitis in patients with diabetes were collected in this study; these factors included demographic characteristics, diseases, and lifestyles that may be associated with diabetes mellitus and periodontitis, as well as standard blood biochemical test indices [13, 22]. These factors are shown in Schedule 1. The

machine learning algorithm was used to develop predictive models and perform consensus clustering analysis, which in turn were employed to identify populations of patients with diabetes at high risk for the occurrence of periodontitis and to make clinical decisions for patient management. This was done to provide targeted oral health education and develop a comprehensive periodontal treatment program. The aim of this study was to provide reference information for periodontal disease prevention in patients with diabetes, as well as for early clinical diagnosis and treatment.

Information and methods

Data downloading and processing

The National Health and Nutrition Examination Survey (NHANES) is a multistage, nationally representative survey of the US noninstitutionalized population that has been conducted on a 2-year cycle since 1999, with data collected through interviews and physical examinations [23]. The Ethics Review Board of the CDC's National Center for Health Statistics approved the protocol for collecting oral health data, and all survey participants provided written informed consent [24]. Since the NHANES collected periodontal examination data reflecting periodontal health (periodontal pockets, periodontal atrophy, and loss of attachment) from participants aged 30 years and older only from 2009 to 2014, data from three cycles, that is, 2009–2010, 2011–2012, and 2013–2014, with a total of 28,779 participants, were used in this study (Figure 2). This study followed the Strengthening the Reporting of Observational Studies in Epidemiology (STROBE) guidelines [25]. The data analyzed were publicly available and therefore did not require any Institutional Review Board approval.

The study interviews included questions regarding demographic data (sex, age, race, marital status, pregnancy status, income-to-poverty ratio, frequency of alcohol consumption, sleep habits, job stress, dietary practices, etc.), education level, and medical history (diabetes, hypertension, hyperlipidemia, osteoporosis, etc.). The clinical examination included medical and dental evaluations and laboratory tests. The participants considered to have diabetes were individuals who self-reported having diabetes, had a glycated hemoglobin level of $\geq 6.5\%$, or were taking oral hypoglycemic medication or insulin [26]; there were a total of 2,648 participants, with 1,931 adults aged 30 years or older who underwent a periodontal health check-up (periodontal pockets, periodontal atrophy, and loss of adhesion). To ensure data quality, 1,804 patients with diabetes aged 30 years and older with complete periodontal examination data were ultimately enrolled in this study after the exclusion of participants with ≥ 0.2 missing interview content data and clinical examination data from the overall survey. The diagnosis of periodontitis was based on the classification of periodontitis developed for the American Academy of Periodontology in collaboration with the Centers for Disease Control and Prevention (CDC), which is applicable to NHANES data according to Eke et al. [27]. Mild periodontitis was defined

as follows: 2 adjacent attachment losses (ALs) ≥ 3 mm, ≥ 2 adjacent probing depths (PDs) ≥ 4 mm (not on the same tooth), or 1 PD ≥ 5 mm [27]. Based on this assessment, 1,401 of the 1,804 participants included had periodontitis, and 403 did not have periodontitis. Missing values in the raw data were filled in via multiple interpolation with the R language 'mice 3.12.0' software package, and the processed data were background corrected and normalized by using MinMaxScaler (the samples were scaled to [0, 1]).

Statistical analysis

Statistical analyses were performed via SPSS 25.0 software. Normality tests were conducted via the Kolmogorov-Smirnov test. Continuous variables are herein presented as medians (interquartile ranges [IQRs]) and were compared via the Mann-Whitney U test. Categorical variables are expressed as n (%) and were compared via the chi-square test or Fisher's exact test. Group comparisons were made through the independent samples t test or one-way ANOVA. Count data are presented as $[n$ (%)], and measurements are expressed as $(x \pm s)$. The relationships between clinical and blood biochemical factors affecting patients with diabetes and periodontitis were analyzed via Spearman's correlation analysis. A significance level of $p < 0.05$ was used.

Feature screening

Some studies have shown that when the model incorporates too many influencing factors, problems such as overfitting, degradation of the generalization ability, and an increase in covariance among independent variables may occur, making the model coefficients difficult to interpret [28]. In recent years, to avoid these problems and formulate simple and easy-to-explain clinical models, researchers have advocated the use of scientific statistical methods to screen variables before modeling when the model contains more variables [29, 30]. This is done to simplify the model as much as possible and improve its statistical efficiency and interpretability [31]. The traditional single-factor analysis screening results reflect only whether there are statistical discrepancies in the relationships between single influential factors and the results, ignoring the covariance among multiple influential factors and the effects of complex interactions on the results [32–34]. The least absolute shrinkage and selection operator (LASSO) compresses some of the regression coefficients by constructing a penalty function, forcing the sum of the absolute values of the coefficients to be smaller than a certain fixed value [35]. It also sets some of the regression coefficients to zero to obtain a more refined model. This technique can improve the prediction accuracy and explanatory power of the model. LASSO is currently one of the most widely used feature selection techniques [36, 37]. The basic principle is to introduce the L1 regularization term on the basis of the ordinary least squares method to achieve feature selection and coefficient sparseness in the model by minimizing the objective function.

The LASSO regression objective function is as follows: minimize $\|Y - X\beta\|^2 + \lambda\|\beta\|_1$, where Y is the observation value vector, X is the feature matrix, β is the regression coefficient vector to be estimated, and λ is a hyperparameter that controls the strength of regularization. The L1 regularization term $\lambda\|\beta\|_1$ plays a key role in the objective function. This term introduces sparsity and the coefficients of some features are reduced to zero, thereby achieving automatic feature selection. Therefore, LASSO regression can not only make predictions but also identify features that have an important impact on the target variable. The data were balanced via the SMOTE (Synthetic Minority Over Sampling Technique) method before feature selection so that the ratio of minority to majority categories was 1:1. The final matching result was periodontitis (1.0) = 1,401 patients with diabetes with periodontitis and periodontitis (0.0) = 1,401 patients with diabetes without periodontitis. LASSO regression was then used for feature selection to build this LASSO regression model on a binomial basis (R4.2.3 version: glmnet4.1.8 software package).

ML modeling and development

Machine learning methods can be categorized into several types, including supervised learning, unsupervised learning, semi-supervised learning, and reinforcement learning. Supervised learning utilizes labeled training data to train models that are capable of making predictions on new data. In contrast, unsupervised learning automatically identifies patterns and regularities from unlabeled data. Semi-supervised learning combines supervised and unsupervised learning techniques, while reinforcement learning develops optimal behavioral strategies through trial and error. In this study, predictive models were developed and compared using nine commonly employed representative algorithms from the domain of supervised learning. The methods used to construct the predictive models include XGBoost, Logistic Regression, LightGBM, Random Forest, AdaBoost, Multi-Layer Perceptron, Support Vector Machines, k-Nearest Neighbours, and Naive Bayesian classification algorithm. All the prediction models were implemented via Python 3.7, with the packages 'xgboost 1.2.1' for XGBoost, 'lightgbm 3.2.1' for LightGBM, and 'sklearn' for the remaining models. The package for XGBoost is 'lightgbm 3.2.1', and the package for the remaining models is 'sklearn 0.22.1'. The bootstrap resampling technique was used to train and classify the ML models. The potential predictor variables selected in the feature selection process of LASSO regression were used in the predictive models. Patients were randomly divided into training and testing groups at an 8:2 ratio. The validation dataset was used to evaluate and compare the performance of each model. The ability of the model to predict periodontitis development in patients with diabetes was evaluated in terms of the area under the working characteristic curve (AUROC) of the subjects, accuracy, sensitivity, specificity, positive predictive value, negative predictive value, and F1 score. Model calibration was assessed via a 'prediction versus observation' plot for periodontitis incidence.

Model optimization and evaluation

To ensure the stability of the model, tenfold cross-validation was used to evaluate its predictive ability. The training set was randomly divided into 10 groups, with eight groups used for training and two groups used for validation in each iteration. During training, 20% of the datasets from the training set were randomly selected to test the model's performance. The models were then quantified individually via receiver operating characteristic (ROC) curve analysis, and their prediction accuracy was evaluated via area under the curve (AUC) and calibration plots. Clinical usefulness and net benefit were assessed via decision curve analysis (DCA). Shapley additive explanation (SHAP), a feature-importance metric based on game theory, was used to calculate the contribution of each feature to the model output. The SHAP was used to assess feature importance, with higher absolute SHAP values indicating features with the greatest impact on the model's predictive scores. The distribution of feature values and their correlation with model predictions were also assessed, and the most reliable model was selected to construct a clinical prediction model for predicting the occurrence of periodontitis in patients with diabetes via blood test indices. An online web calculator was developed for applying the model and predicting the likelihood of periodontitis in specific individual patients with diabetes.

Consensus clustering analysis

The K-means clustering algorithm is an unsupervised learning algorithm that is commonly used to divide the data set into K clusters. The K-means clustering algorithm can classify patients with similar disease risks into the same group, identify the characteristics of different disease risk patterns in patient groups, and help conduct risk assessments and customize personalized treatment plans. This study was based on the feature variables screened out by baseline analysis and LASSO regression, using the K-means clustering algorithm to calculate the distance between features and automatically aggregate and classify patients with diabetes to identify those with clinical and blood biochemical parameters at a high risk of periodontitis. The optimal number of groups (K) was determined by comparison with the flexed elbow method. The above statistical analyses were completed using the R package (version 4.2.3) and Python (version 3.11.4). The performance of clustering models is commonly evaluated by the silhouette coefficient and the Calinski-Harabasz index. The value of the silhouette coefficient ranges from -1 to 1. The closer the value is to 1, the better the model performance. The Calinski-Harabasz index ranges from 0 to $+\infty$. The larger the value is, the better the model performance.

Continuous variables are expressed herein as the means $\pm$ standard deviations or medians $\pm$ IQRs. Differences between diabetes clusters were compared via the unpaired t test or Mann-Whitney U test. Categorical variables are expressed as absolute numbers (n) and relative frequencies (%) and were analyzed via the chi-square test or Fisher's exact test. All the

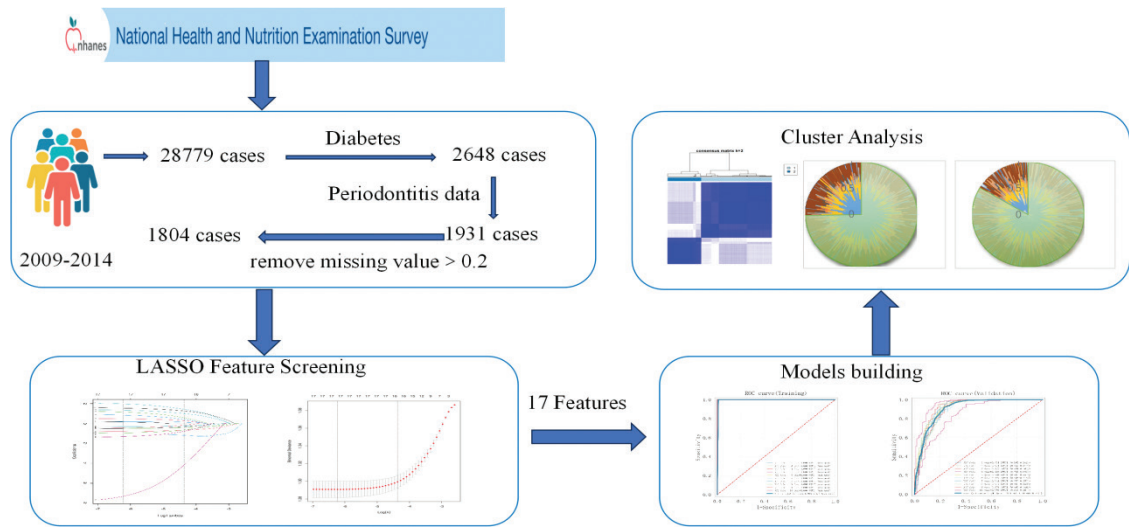


Figure 1. Flow chart of the study.

statistical analyses were performed via SPSS Statistics for Windows (version 25, IBM Corp., Armonk, NY, USA). Differences with $p \leq 0.05$ were considered statistically significant.

Results

Statistical analysis

As shown in **Schedule 1**, 1,804 patients with diabetes were analyzed: 1,401 with periodontitis and 403 without periodontitis. The results indicate that age, race, education level, moderate

leisure, high work intensity, income-to-poverty ratio, albumin (ALB g/L), alanine aminotransferase (ALT U/L), blood urea nitrogen (BUN mmol/L), total cholesterol (TC mmol/L), creatinine (Cr μ mol/L), lactate dehydrogenase (LDH U/L), total protein (TP g/L), potassium (K mmol/L), osmolality (Osm mmol/kg), globulin (GLB g/L), and glycohemoglobin (GHb %) were among the 19 potential characteristics that were significantly different ($p < 0.05$) between the two groups of patients with diabetes with and without periodontitis. These characteristics may have an impact on the prevalence of periodontitis in patients with diabetes.

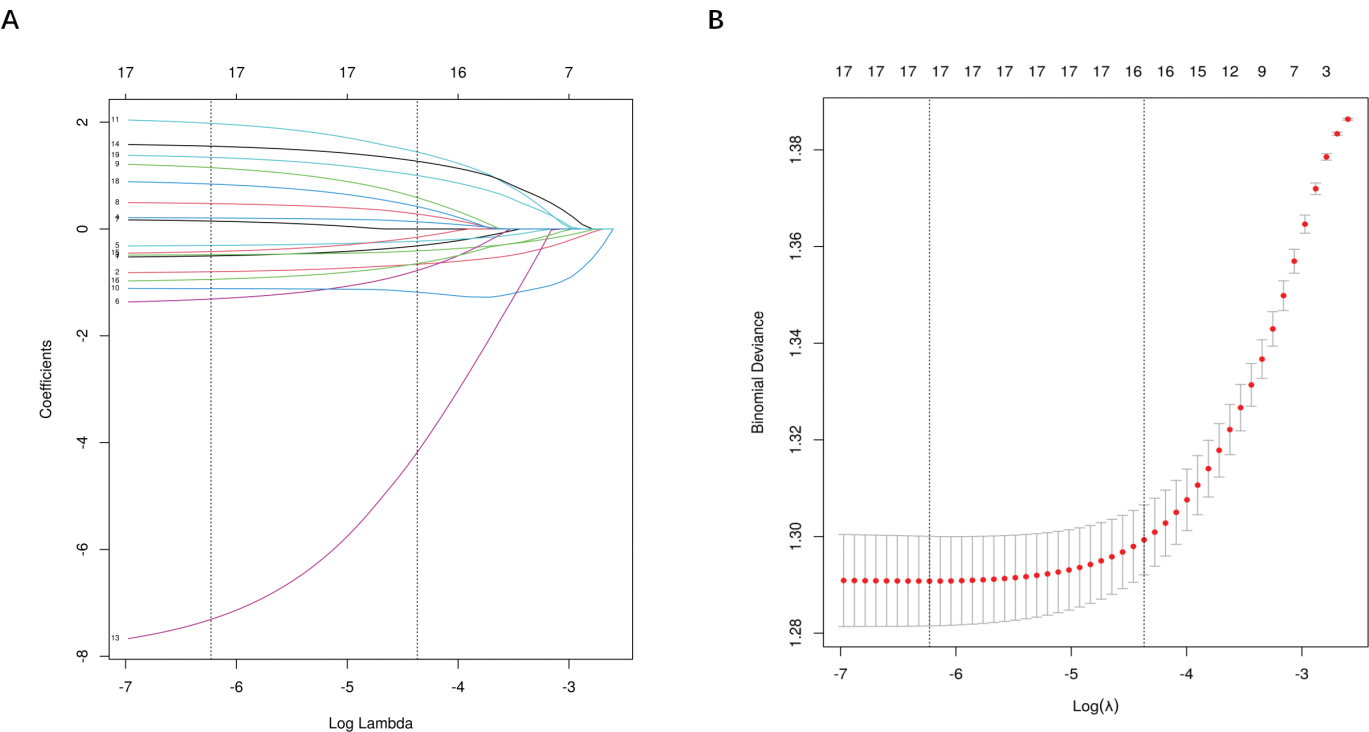


Figure 2. LASSO regression feature selection: coefficient profiles for Lasso regression (A) and plot of the optimal tuning parameter (λ) determined by 10-fold cross-validation of the Lasso regression (B).

Table 1. Predictive performance of nine machine learning algorithms in the development of periodontitis in diabetic patients.

Models	AUC	Accuracy	Sensitivity	Specificity	PPV	NPV	F1 score	Kappa
XGBoost	0.917 (0.884–0.951)	0.803 (0.786–0.820)	0.719 (0.687–0.752)	0.887 (0.869–0.904)	0.864 (0.846–0.882)	0.761 (0.741–0.781)	0.784 (0.763–0.806)	0.606 (0.572–0.640)
logistic	0.671 (0.608–0.734)	0.628 (0.612–0.644)	0.592 (0.576–0.607)	0.665 (0.641–0.688)	0.639 (0.621–0.657)	0.619 (0.605–0.633)	0.614 (0.599–0.629)	0.256 (0.225–0.288)
LightGBM	0.917 (0.883–0.950)	0.834 (0.819–0.849)	0.828 (0.813–0.843)	0.84 (0.815–0.865)	0.839 (0.818–0.861)	0.83 (0.817–0.843)	0.833 (0.819–0.847)	0.668 (0.639–0.697)
Random Forest	0.907 (0.872–0.942)	0.827 (0.809–0.845)	0.804 (0.786–0.821)	0.85 (0.822–0.879)	0.844 (0.818–0.871)	0.813 (0.797–0.828)	0.823 (0.806–0.840)	0.654 (0.618–0.690)
AdaBoost	0.888 (0.850–0.927)	0.811 (0.794–0.828)	0.882 (0.855–0.908)	0.74 (0.712–0.769)	0.773 (0.755–0.792)	0.864 (0.839–0.889)	0.823 (0.807–0.840)	0.622 (0.587–0.656)
CNB	0.638 (0.573–0.702)	0.61 (0.597–0.622)	0.482 (0.450–0.514)	0.737 (0.701–0.773)	0.651 (0.624–0.679)	0.588 (0.578–0.598)	0.551 (0.532–0.571)	0.219 (0.195–0.244)
MLP	0.722 (0.663–0.781)	0.664 (0.647–0.682)	0.662 (0.621–0.702)	0.667 (0.625–0.708)	0.667 (0.646–0.689)	0.665 (0.645–0.685)	0.662 (0.640–0.684)	0.328 (0.293–0.364)
SVM	0.820 (0.771–0.868)	0.733 (0.719–0.748)	0.712 (0.673–0.751)	0.755 (0.723–0.787)	0.746 (0.725–0.767)	0.726 (0.703–0.749)	0.726 (0.707–0.746)	0.467 (0.437–0.496)
KNN	0.833 (0.787–0.879)	0.747 (0.740–0.755)	0.632 (0.615–0.650)	0.862 (0.838–0.886)	0.823 (0.802–0.845)	0.701 (0.695–0.708)	0.714 (0.706–0.722)	0.495 (0.480–0.510)

AUC: area under the curve; PPV: positive predictive value; NPV: negative predictive value; XGBoost: extreme gradient boosting; LR: logistic regression; LightGBM: light gradient boosting; RF: random forest; AdaBoost: AdaBoost Classifier; CNB: Complement NB; MLP: multi-layer perceptron; SVM: support vector machine; KNN: k-nearest neighbors.

Feature screening

After the data of 1804 patients (periodontitis (1.0) = 1401 patients in the periodontitis group of patients with diabetes and periodontitis (0.0) = 1401 patients in the nonperiodontitis group of patients with diabetes) were balanced via the SMOTE 1:1 (Synthetic Minority Oversampling Technique) method, the 19 potential characteristics ($p < 0.05$) were further screened via the LASSO regression algorithm. The results were further screened, as shown in Figure 2 (A and B), when the λ of the minimum mean square error in LASSO regression was 0.002. The 17 predictors (age, race, education level, moderate leisure, high work intensity, income-to-poverty ratio, drinking frequency, smoking status, ALB, ALT, TC, Cr, LDH, TP, K, Osm, and GHb levels) were considered the most valuable characteristics for developing a predictive model for periodontitis development in patients with diabetes, which could be used to construct predictive models.

Predictive modeling

The performance of the nine ML classification models for predicting the risk of periodontitis development in patients with diabetes was compared in the training set and validation set. This comparison is shown in Table 1 and Figure 3. Multiple evaluation metrics were employed to assess the performance of the prediction models. The ROC curves of the different ML models that predicted the presence or absence of periodontitis in patients with diabetes in the training and validation sets are shown in Figure 3 A and B. Calibration plots (Figure 3C) were used to determine the accuracy of the models. Furthermore, the ROC curve results for periodontitis prediction for each model are presented in the forest plot in Figure 3D. We observed that the XGBoost model performed better in both the training and validation sets, and it can be argued that XGBoost is the best

model choice for this dataset and performs best in predicting the occurrence of periodontitis in patients with diabetes.

The test set consisted of 560 cases (20.00%), which were randomly selected from a sample dataset of 2,802 cases (1,401 with periodontitis and 1,401 without periodontitis). The remaining samples were used for the training set in a 10-fold cross-validation. The XGBoost algorithm was used to predict the risk of patients with diabetes developing periodontitis. In the final test set, the model had an AUC of 0.9165 and an accuracy of 0.8021, demonstrating the best predictive performance, as shown in Table 2 and Figure 4A–C. The learning curve (Figure 4D) indicated that the model successfully predicted the risk of periodontitis development in patients with diabetes. The difference in error between the training and validation sets converged as the number of training samples increased, as shown in the learning curve. The calibration plot (Figure 4E) demonstrated that the XGBoost model accurately predicted the probability of the observed incidence of periodontitis in patients with diabetes. Additionally, a decision plot (Figure 4F) was constructed via a DCA, which showed the net gain in threshold probability on the basis of the model's output decision. To further explain our model, we analyzed the results of the XGBoost algorithm via SHAP, as shown in Figure 4G. This analysis revealed the relationship between high and low eigenvalues, as well as the SHAP values in the dataset. Each point on the plot represents the eigenvalue and SHAP value for each patient. Figure 4H displays bar charts indicating the importance of the 17 features assessed via the SHAP values. The impact of each feature on the predictive model is represented by the bars with the mean absolute SHAP value. Figure 4I and J provide an individual-level breakdown of how these model features led to changes in predicted periodontitis risk scores for two example patients with diabetes. These figures identify which feature values strongly influenced the final risk prediction score by

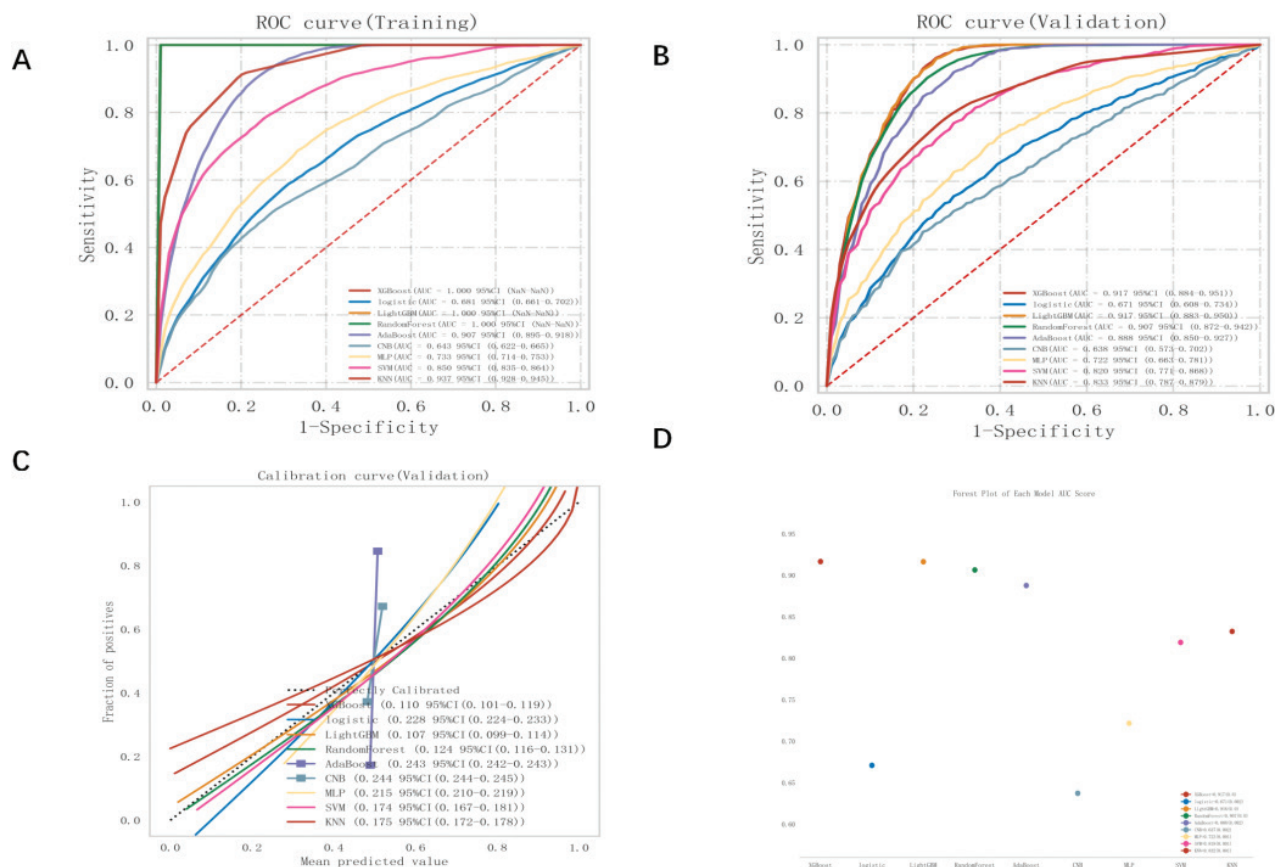


Figure 3. ROC curve analysis was conducted on nine machine learning models to predict the risk of periodontitis in diabetic patients. The analysis compared the performance of these models on the training set (A) and validation set (B). Calibrated typology plots (C) were used to evaluate the predictive models, and a forest plot (D) was created to display the AUC scores of the nine models.

either increasing or decreasing it. On the basis of the feasibility of using the XGBoost algorithm and the 17 predictors mentioned above to predict the risk of periodontitis development in patients with diabetes, an online network calculation prediction model was developed, as shown in Figure 5.

Consensus cluster analysis

To further analyze the prevalence of periodontitis in patients with diabetes, a consensus clustering algorithm was used in this study to identify 1,804 samples of patients with diabetes (with periodontitis: $N = 1,401$, without periodontitis: $N = 403$) with periodontitis. The optimal number of clusters was found when $k = 2$ (Figure 6A and B), and the 1,804 diabetic patient samples were divided into two clusters: Cluster A ($n = 1,222$) and Cluster B ($n = 582$) (Figure 6C). The periodontitis and smoking status, education level, income-to-poverty ratio, ALB level, and ALT level

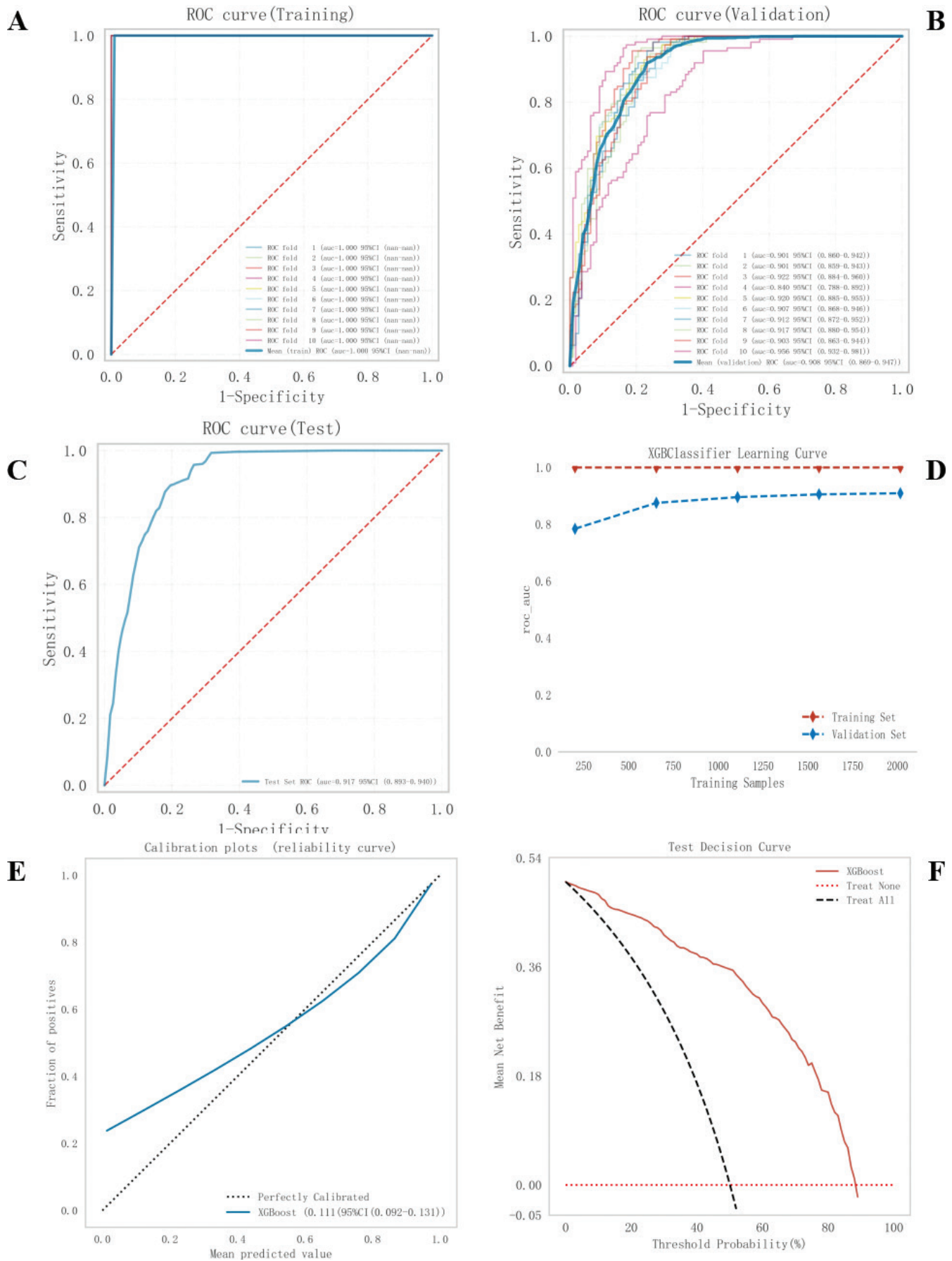
significantly differed across the various diabetes subclusters, as shown in Table 3 and Figure 6D–F.

Discussion

In this study, we found that the prevalence of periodontitis among patients with diabetes was as high as 77.66%. We established a predictive model for periodontitis risk in patients with diabetes and conducted a consistent consensus Cluster A analysis. For the risk of periodontitis in patients with diabetes, age, race, education level, moderate leisure, high work intensity, income-to-poverty ratio, alcohol consumption frequency, smoking status, and the levels of ALB, ALT, TC, Cr LDH, TP, K, Osm, and glycohemoglobin (GHb%) were predicted by a comprehensive assessment of 17 risk predictors. The periodontitis risk was significantly greater in diabetes patients in Cluster B (83.68%) than in those in Cluster A (74.80%). There were also

Table 2. Performance of XGBoost model in predicting the risk of periodontitis in diabetic patients.

Model	AUC	Accuracy	Sensitivity	Specificity	PPV	NPV	F1 score
Training set	1.000 (NaN–NaN)	1.000 (1.000–1.000)	0.999 (0.999–0.999)	1.000 (1.000–1.000)	1.000 (1.000–1.000)	0.999 (0.999–0.999)	1.000 (1.000–1.000)
Validation set	0.908 (0.869–0.947)	0.784 (0.761–0.807)	0.683 (0.650–0.716)	0.884 (0.860–0.908)	0.855 (0.829–0.882)	0.737 (0.714–0.760)	0.908 (0.869–0.947)
Test set	0.917 (0.893–0.940)	0.802	0.705	0.9	0.876	0.752	0.781



(Figure 4 Continued)



Figure 4. Construction and evaluation of the XGBoost model. (A–C) XGBoost ROC curves using 10-fold cross-validation on the training set (A), validation set (B), and test set (C). (D) Machine learning curves and (E) Calibration plots for XGBoost. (F) Decision curve analysis plot. (G–J) Feature importance SHAP summary charts and bar graphs. (G) The dot plots on the left indicate the direction of the contribution of each value of each variable, with red representing the larger value and blue representing the smaller value. (H) The bars on the right represent the importance of the variables and their overall contribution to the model predictions. (I), (J) SHAP score explains the predicted risk of periodontitis prevalence in two diabetic subjects.

significant differences in clinical and blood biochemical indices among patients with diabetes in both clusters. Diabetes Subcluster B, which had a higher risk of periodontitis, was characterized mainly by a smoking habit, a lower education level, a higher income-to-poverty ratio, and higher ALB and ALT levels.

Smoking is recognized as an important risk factor for periodontitis. However, in a study by Han et al. [38], no synergistic effect of smoking and diabetes mellitus on periodontitis was

observed. In other words, the combined effect of diabetes mellitus and smoking on periodontitis did not appear to be significant. However, in the present study, there were significantly more patients with diabetes in Subcluster B who were habitual smokers (75.10%) than there were in Subcluster A (65.08%). These findings suggest that smoking in patients with diabetes also leads to an increased risk of periodontitis, which is similar to the results of previous studies [39–41].

Figure 5. XGBoost constructs an online network calculator to predict the risk of periodontitis in diabetic patients.

Education level and the income-to-poverty ratio can reflect an individual's socioeconomic status to some extent. In this study, patients with diabetes with lower education levels were at greater risk of periodontitis, which is consistent with the findings of Walther and Zini et al. [42, 43]. Socioeconomic status inequality is associated with oral health inequality [39, 44, 45]. The study also revealed that a higher income-to-poverty ratio in diabetes patients in Cluster B was associated with a greater risk of periodontitis development, as shown by Buchwald et al. [46]. By examining the relationship between socioeconomic status factors, systemic inflammation, and the progression of periodontitis, previous studies have shown that a low level of education and lower economic income factors can impact the development of periodontitis. However, some scholars have argued that education level and economic factors do not seem to be significantly associated with the risk of periodontitis [47, 48]. Therefore, there is inconsistency in the results of previous studies regarding their influence on the risk of periodontitis. The results of this study may have been influenced by the complex disease course of diabetes, and further investigation is needed to determine the influence of education level and the income-to-poverty ratio on the risk of periodontitis in patients with diabetes.

Cui et al. [49] compared salivary proteins between healthy and periodontitis populations and reported that the concentrations of TP, ALB, and GLB were greater in the saliva of patients with periodontitis than in the saliva of healthy individuals. Banu et al. [50] reported that the concentrations of ALT and aspartate aminotransferase (AST U/L) in the serum and saliva were significantly greater in individuals with periodontitis than in healthy individuals. Widita et al. [51] analyzed the relationships between hepatic enzyme levels and periodontal clinical parameters and concluded that elevated ALT levels may be associated with changes in periodontal clinical parameters

parameter description

Race:1: Mexican American; 2: Other Hispanic; 3: Non-Hispanic White; 4: Non-Hispanic Black; 5: Other Race;
 Educationlevel:1:Less Than 9th Grade;2:9-11th Grade; 3:High School Grad/GED or Equivalent;4:Some College or AA degree;5:College Graduate or above;
 Drinkingfrequency:1:week;2:month;3:year;
 Highworkingintensity:1:one day;2:two days; 3:three days;4:four days;5:five days; 6:sx days; 7:seven days;
 Moderateleisure:1:Yes; 2:No
 Smoking:1: Every day;2: Some days;3:No;
 ALB:Albumin (U/L);
 ALT:Alanine aminotransferase (U/L);
 BUN:Blood Urea Nitrogen (mmol/L) ;
 TC:Total Cholesterol (mmol/L) ;
 Cr:Creatinine (umol/L) ;
 LDH:Lactate dehydrogenase (U/L);
 TP:Total Protein (g/L) ;
 K:Potassium (mmol/L);
 Osm:Osmolality (mmol/Kg);
 GHb:Glycohemoglobin (%);

[52]. The present study also obtained similar results from the analysis of blood standard biochemistry and the periodontitis prevalence in patients with diabetes [16]. Cluster B patients with diabetes with a higher risk of periodontitis were found to have higher levels of ALB and ALT, suggesting that elevated levels of ALB and ALT remain important predictive risk factors for periodontitis development in patients with diabetes.

Machine learning models are becoming increasingly popular in precision medicine [53]. In this study, the use of machine learning to construct a clinical prediction model with k-means clustering helped to identify homogeneous groups and revealed stronger associations between potential risk factors for periodontitis development in patients with diabetes mellitus. These findings suggest that constructing a clinical prediction model and a clustering model using machine learning can aid in identifying high-risk groups of diabetes mellitus patients for periodontitis development and, it is hoped, improve the prevention, identification, treatment, and management of this common disease.

The present study had a cross-sectional study design, and issues such as temporally related causality and some recall bias could not be avoided despite our best efforts. Additionally, because only the 2009–2014 dataset from the NHANES included full-mouth periodontal clinical examination measurements and because that dataset primarily pertained to the US population, with a lack of similarly large samples of complete datasets since then, the generalizability of the present results for predicting the risk of periodontitis and the clustering of the current diabetic population is limited.

Conclusion

The periodontitis risk prediction model for patients with diabetes, which was constructed on the basis of machine learning,

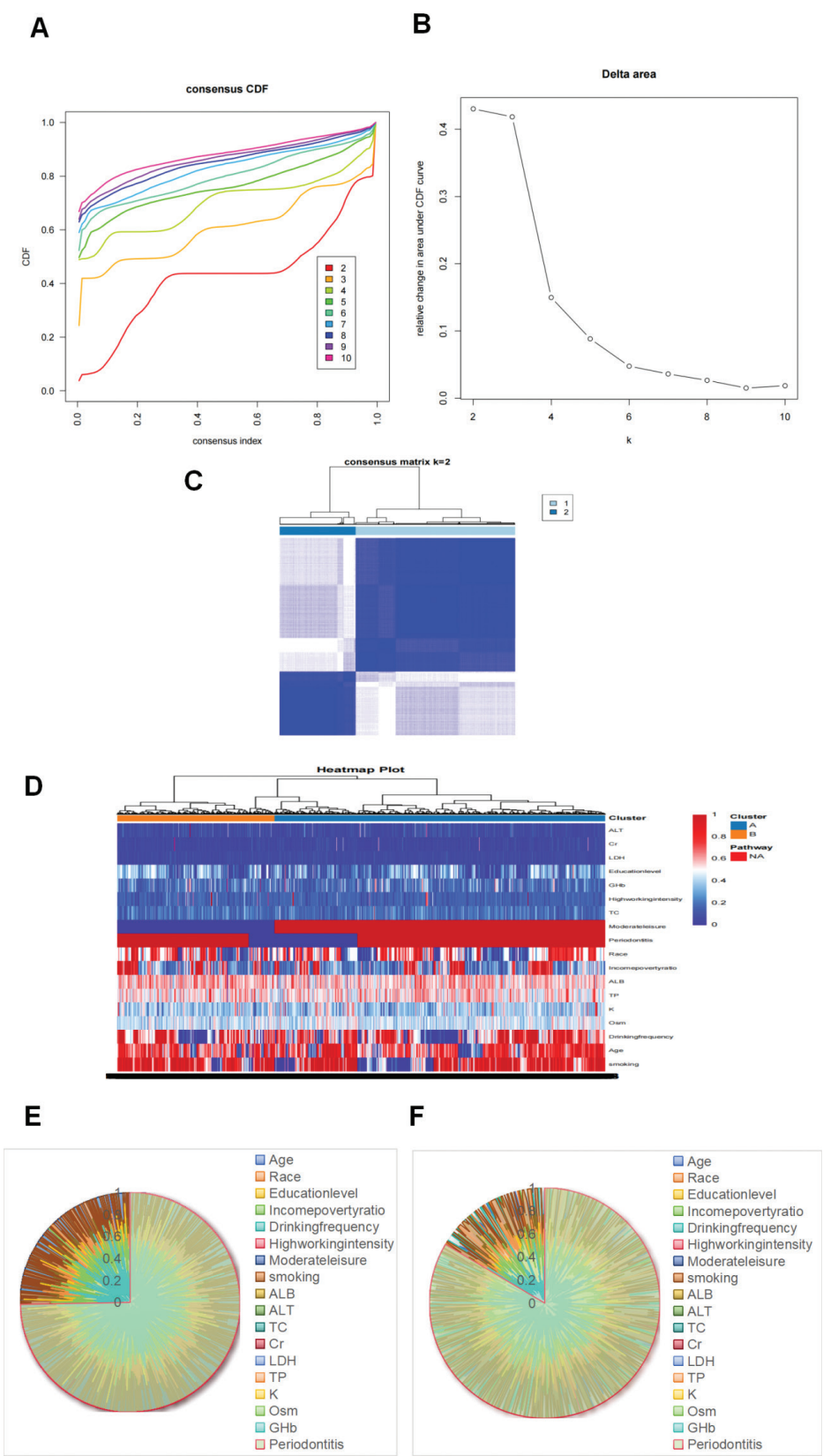


Figure 6. Consensus clustered cumulative distribution function (CDF) for (A) $k = 2-10$; relative change in the area under the CDF curve for (B) $k = 2-10$; heatmap of the cooccurrence matrix of the sample of diabetic patients for (C) $K = 2$. Heat map of cluster analysis results for (D). Diabetes subtype family 1 radar map for (E). Diabetes subtype family 2 radar map for (F).

Table 3. Cluster analysis table of diabetes subtypes based on periodontitis.

Variant	Subcategory	Overview (n = 1,804)	Cluster A (n = 1,222)	Cluster B (n = 582)	p
Periodontitis, n (%)	0	403 (22.339)	308 (25.205)	95 (16.323)	<0.001
	1	1401 (77.661)	914 (74.795)	487 (83.677)	
Race, n (%)	1	314 (17.406)	219 (17.921)	95 (16.323)	0.125
	2	178 (9.867)	112 (9.165)	66 (11.340)	
	3	627 (34.756)	438 (35.843)	189 (32.474)	
	4	473 (26.220)	322 (26.350)	151 (25.945)	
	5	212 (11.752)	131 (10.720)	81 (13.918)	
Education level, n (%)	1.0	350 (20.278)	249 (21.392)	101 (17.972)	<0.001
	2.0	293 (16.976)	218 (18.729)	75 (13.345)	
	3.0	381 (22.074)	267 (22.938)	114 (20.285)	
	4.0	434 (25.145)	278 (23.883)	156 (27.758)	
	5.0	263 (15.238)	148 (12.715)	115 (20.463)	
Drinking frequency, n (%)	9.0	5 (0.290)	4 (0.344)	1 (0.178)	0.611
	1.0	295 (33.258)	189 (33.275)	106 (33.229)	
	2.0	207 (23.337)	127 (22.359)	80 (25.078)	
	3.0	385 (43.405)	252 (44.366)	133 (41.693)	
	4.0	171 (13.266)	110 (12.500)	61 (14.914)	
High working intensity, n (%)	2.0	1071 (83.088)	742 (84.318)	329 (80.440)	0.421
	3.0	11 (0.853)	7 (0.795)	4 (0.978)	
	4.0	4 (0.310)	3 (0.341)	1 (0.244)	
	5.0	17 (1.319)	10 (1.136)	7 (1.711)	
	6.0	6 (0.465)	2 (0.227)	4 (0.978)	
Moderate leisure, n (%)	7.0	9 (0.698)	6 (0.682)	3 (0.733)	nan
	1	582 (32.262)	0 (0.000)	582 (100.000)	
Smoking, n (%)	2	1222 (67.738)	1222 (100.000)	0 (0.000)	<0.001
	1.0	253 (27.772)	207 (31.846)	46 (17.625)	
	2.0	39 (4.281)	20 (3.077)	19 (7.280)	
Age, median [IQR]	3.0	619 (67.947)	423 (65.077)	196 (75.096)	0.373
	nan	59.000 [43.000,68.000]	60.000 [43.000,69.000]	57.000 [44.000,68.000]	
	nan	1.670 [0.950,3.310]	1.590 [0.920,3.140]	1.880 [1.050,3.580]	
Income poverty ratio median [IQR]	nan				0.002
ALB, median [IQR]	nan	41.000 [39.000,43.000]	41.000 [39.000,43.000]	42.000 [40.000,44.000]	<0.001
ALT, median [IQR]	nan	22.000 [17.000,30.000]	21.000 [16.000,30.000]	23.000 [18.000,31.000]	0.002
TC, median [IQR]	nan	4.577 [3.827,5.456]	4.603 [3.827,5.482]	4.500 [3.827,5.353]	0.228
Cr, median [IQR]	nan	79.560 [65.420,99.890]	79.560 [64.530,100.780]	77.790 [66.300,97.240]	0.172
LDH, median [IQR]	nan	129.000 [112.000,147.000]	129.000 [113.000,148.000]	127.000 [112.000,145.000]	0.096
TP, median [IQR]	nan	71.000 [68.000,75.000]	71.000 [68.000,75.000]	72.000 [68.000,75.000]	0.886
K, median [IQR]	nan	4.100 [3.800,4.300]	4.100 [3.800,4.300]	4.000 [3.800,4.300]	0.616
Osm, median [IQR]	nan	282.000 [278.000,285.000]	282.000 [278.000,285.000]	281.000 [278.000,285.000]	0.099
GHb, median [IQR]	nan	6.900 [6.400,8.000]	6.900 [6.400,8.000]	6.900 [6.400,8.000]	0.642

Periodontitis (1: yes; 0: no), Race (1: Mexican American; 2: Other Hispanic; 3: Non-Hispanic White; 4: Non-Hispanic Black; 5: Other Race), Education Level (1: College Degree or Higher; 2: College Degree; 3: High School Graduate/GED or Equivalent; 4: 9th-11th grade; 5: Less than 9th grade; 9: Don't Know), Drinking frequency (1: Week; 2: Month; 3: Year), High working intensity: days of a week, Moderate leisure (1: yes; 2: no), Smoking (1: daily; 2: occasional; 3: no). ALB: Albumin (g/L); ALT: Alanine aminotransferase (U/L); TC: Total cholesterol (mmol/L); Cr: Creatinine (umol/L); LDH: Lactate dehydrogenase (U/L); TP: Total protein (g/L); K: Potassium (mmol/L); Osm: Osmolality (mmol/Kg); GHb: glycated hemoglobin (%).

shows excellent performance and can be used in the clinic for individualized prediction of the risk of periodontitis development in patients with diabetes. The consistent consensus Cluster A analysis method can effectively reveal the relationship between clusters of patients with diabetes and the risk of developing periodontitis, providing more support and guidance for personalized medical treatment and precision therapy of

periodontal inflammatory disease in patients with diabetes. The accompanying online risk calculator serves as an easy-to-use tool for physicians to develop precise prevention and treatment strategies. In the future, our aim is to integrate more case data with imaging, molecular, and genetic data to further improve the performance of our risk assessment model and validate it in additional studies covering a wider population.

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Author contributions

All the authors have made substantial contributions to the conception and design of the study. ZA, YR, and CY were involved in the conception or design of the study. CT and RX were involved in the data collection and data analysis. ZA and LZ interpreted the data, drafted the manuscript, revised the manuscript critically and provided final approval of the version to be published.

Availability of data and materials

The datasets supporting the conclusions of this study are available in the NHANES database (<https://www.cdc.gov/nchs/nhanes>).

Declarations

Ethics approval and consent to participate: Not applicable.

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