Analysis and Evaluation of Machine Learning Models for Diabetes Risk Prediction

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Introduction

- Chronic Condition: Diabetes is a long-term health condition that impacts millions globally.
- Significance of Early Diagnosis: Early detection and prediction of diabetes are crucial for effective disease management and reducing healthcare expenses.
- Role of Classification Models: Classification models are widely used for predicting diabetes. They leverage health data to make predictions.
- Advantages Over Traditional Methods: These models can identify complex patterns in datasets. They are often more effective than traditional statistical techniques.

Project Goals

- Develop machine learning models to predict diabetes risk.
- Understand the impact of key features on predictions.
- Balance accuracy and computational efficiency.
- Ensure model interpretability for clinical use.

Literature Review

- Role of Machine Learning in Diabetes Prediction
- Leverages structured datasets (e.g., glucose levels, age, BMI, family history).
- Identifies complex patterns beyond human capability, enhancing diagnostic accuracy.
- 2. Classification Models Overview
- Logistic Regression: Simple and interpretable. Effective on datasets like PIMA Indian Diabetes Dataset (PIDD).
- Decision Trees: Highly interpretable, using branching logic. Perform well on small datasets but prone to overfitting.
- Ensemble Models (Random Forest Gradient Boosting): Combine multiple models for robust predictions. Random Forest reduces overfitting. Gradient Boosting (e.g., XGBoost) captures nonlinear interactions.
- . Data and Challenges
- Class Imbalance: Diabetes datasets often have fewer positive cases. Techniques like SMOTE address imbalance.
- Feature Selection: Key predictors: glucose levels, BMI, family history. Use techniques like PCA to remove redundant features.
- Evaluation Metrics: Accuracy, precision, recall, F1-score, AUC-ROC. Prioritize recall and F1-score in imbalanced datasets.

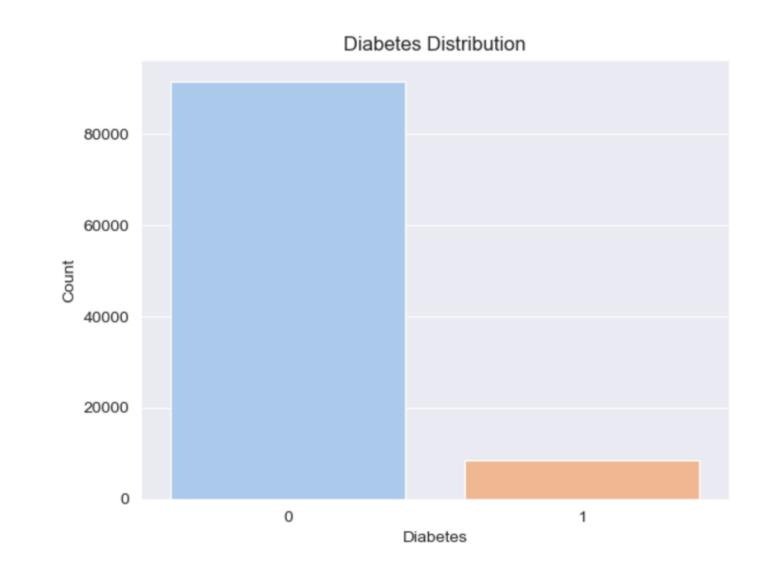


Figure 1. Class imbalance between people with diabetes (1) and people without (0)

Methodology

- Data Pre-processing:
- Data Cleaning: Removed entries with missing values and "Other" gender.
- Categorical Encoding: One-hot encoded "gen der" and "smoking history".
- Feature Scaling: Standardized continuous variables (age, BMI, HbA1c, blood glucose level).
- Model Implementation
 - The implemented models include Decision Tree, Random Forest, XGBoost, LightGBM, and CatBoost. The dataset was split into (80-20) percent testing sets using stratified sampling to ensure the class distribution remained consistent across both subsets.
- Model Evaluation
- Precision
- Recall
- F1-score
- ROC-AUC

Confusion matrix

Cross validation accuracy and cross validation standard deviation

Models	Precisio n	Recal 1	F1 Score	ROC AUC	CV Accurac
Decision Tree	0.7030	0.743	0.722 7	0.857	0.9519
Random Forest	0.9430	0.690 6	0.797 3	0.963 2	0.9699
XGBoost	0.9562	0.692 9	0.803 5	0.978 3	0.9710
LightGB M	0.9655	0.691 2	0.805 6	0.979 2	0.9717
CatBoost	1.0	0.672 9	0.804 5	0.958 0	0.9718

Figure 2. Confusion Matrix

- Proposed Framework
- The framework consists of the following main steps:
- Data Source: The dataset containing various health features (age, gender, BMI, HbA1c level, etc.).
- Data Preprocessing: Includes cleaning the data, encoding categorical variables, handling missing values, and scaling continuous features.
- Model Training: Training multiple machine learning models such as Decision Tree, Random Forest, XGBoost, LightGBM, and CatBoost.
- Evaluation Metrics: Metrics like precision, recall, F1-score, ROC-AUC, and cross-validation are used to evaluate the models.

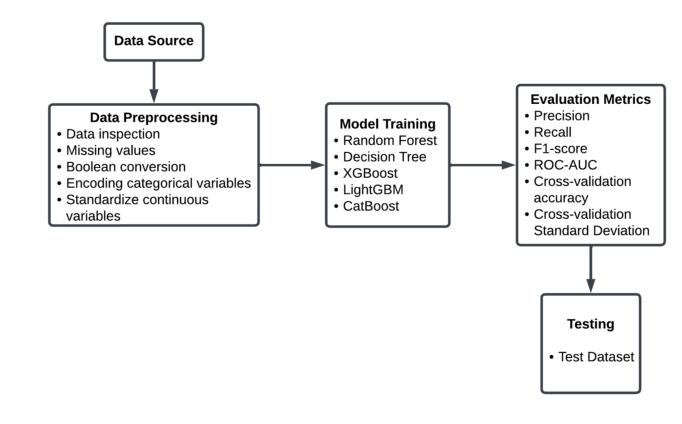


Figure 3. Proposed Framework

Evaluation Results

. Evaluation Metrics Used:

Models were assessed using precision, recall, F1-score, ROC-AUC, confusion matrix, cross-validation accuracy, and standard deviation.

- 2. Performance of Models:
- Decision Tree: Worst overall but had the highest recall.
- Random Forest XGBoost: Performed consistently well across all metrics.
- CatBoost: Excelled in precision and CV-accuracy but lower recall affected its F1-score.
- LightGBM: Best overall with the highest F1-score and ROC-AUC, crucial for classification evaluation.
- 3. Comparison of Top Models:
- LightGBM outperformed others but CatBoost was competitive with strong recall and similar F1-score.
- 4. Feature Importance Across Models: HbA1c level and blood glucose level were most important for Decision Tree, Random Forest, XGBoost, and CatBoost.

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Figure 4. Evaluation Result Table

Conclusion

- 1. LightGBM's Strength: LightGBM was the most effective model, showcasing the highest F1-score and ROC AUC, demonstrating its ability to balance precision and recall effectively for diabetes prediction.
- 2. CatBoost's Competitiveness: CatBoost closely followed LightGBM, excelling in precision and cross-validation accuracy, though its lower recall slightly impacted its overall performance.
- 3. Consistency of Random Forest and XGBoost: Both models provided reliable and consistent results across all evaluation metrics, making them strong alternatives in predictive modeling.
- 4. Decision Tree's Limitation: The Decision Tree model underperformed in most metrics, though its high recall highlighted its potential in prioritizing positive cases.
- 5. Feature Insights: HbA1c levels and blood glucose were identified as critical predictors across most models, emphasizing their importance in diabetes risk assessment.

Recommendations

- 1. Feature Engineering: Explore techniques like Principal Component Analysis (PCA) to uncover new predictive variables or derive more informative features from existing ones.
- 2. Address Class Imbalance: Use methods like SMOTE (Synthetic Minority Oversampling Technique) to improve model performance, particularly in recall and F1-score.
- 3. Hyperparameter Optimization: Apply advanced tuning techniques such as grid search or Bayesian optimization to achieve significant performance enhancements.
- 4. Enhance Generalization: Focus on strategies that ensure the models perform effectively across varied patient populations and healthcare settings.