

# Diabetic Patients Readmission Project

## Overview

This project utilizes a dataset from a US hospital containing over 100,000 records. While the dataset provides ample data points for training a classical machine learning (ML) model, it suffers from two key challenges:

1. **Heavy class imbalance:** which requires careful handling.
2. **Missing values:** across several columns, necessitating preprocessing.

Additionally, model explainability is a critical requirement for medical applications, which must be addressed.

## Data

### Missing Values

First, I identified seven columns with missing values. To handle these:

- Columns with missing values exceeding **35% of all rows** were dropped. This threshold is a tunable hyperparameter and can be adjusted for better results.
- For the remaining missing values:
  - **Numerical features:** were filled with the median.
  - **Categorical features:** were filled with a new category labeled "**Unknown.**"

While these are simple imputation strategies, more sophisticated methods could be explored for improved performance.

### Imbalanced Dataset

As noted earlier, the dataset is highly imbalanced, which biases ML models toward the majority class. To address this, two balancing techniques were considered:

1. **Undersampling** (reducing the majority class).
2. **Upsampling** (e.g., SMOTE for synthetic minority class generation).

Given the project's demo nature and computational constraints, only **undersampling** was implemented, while upsampling (SMOTE) was left for future exploration as an option implemented in code.

## Normalization

Normalization is essential in ML to prevent models from overemphasizing features with larger scales. Here, **standard scaling** was applied to all numerical features.

## Models

Five ML models were evaluated for their readmission prediction performance:

1. **Random Forest**
2. **XGBoost**
3. **LightGBM**
4. **SVM**
5. **MLP**

All models were implemented using either **scikit-learn** or their respective specialized libraries (e.g., **xgboost**, **lightgbm**). They were trained and tested on the same dataset, with results documented in the notebook.

## Evaluation Metric

Standard classification metrics (e.g., accuracy, F1-score) were reported, but **recall for the <30 days readmission class** was prioritized. This is because:

- **False negatives** (missed early readmissions) pose a greater risk in healthcare.
- Misclassifying patients who either **will not be readmitted** or will be readmitted **after 30 days** is less critical and carries less risk.

Among all models, **SVM achieved the highest recall** for the "<30 days" class. However, its performance may still be insufficient for real-world medical deployment, necessitating further tuning.

## Explainability

Explainability is crucial in medical AI applications. While deep learning models often lack interpretability, classical ML models (like those used here) offer better transparency.

For this project, **SHAP (SHapley Additive exPlanations)** was employed to interpret model decisions. One key visualization—the **feature importance plot** for LightGBM—is included in the **images** directory.

## Future Considerations

To further enhance model performance, the following strategies could be explored:

### 1. Advanced Modeling Techniques

- Experimenting with more sophisticated models (e.g., ensemble methods, deep learning architectures) may yield better predictive accuracy.

### 2. Hyperparameter Optimization

- Conducting **grid search or brute force** for key hyperparameters (e.g., learning rate, tree depth, regularization) could fine-tune existing models.

### 3. Improved Class Imbalance Handling

- Implementing **upsampling techniques (e.g., SMOTE)** instead of undersampling may better address data imbalance and improve recall for minority classes.

### 4. Enhanced Explainability

- While SHAP's **feature importance plot** was used here, deeper interpretability analysis (e.g., **dependency plots, interaction effects, or decision plots**) could provide richer insights into model behavior.

Note: See test\_run.ipyn for a sample running of code

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