

محمد قرشي

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Prepared By

Name: Mohamed Elzain Gurashi

ID 251002316

Predictive Modeling for Diabetic Complications Using Artificial Intelligence

At

College of Informatics

Midocean University

Supervisor Name: Dr. Khaled Eskaf

College of Informatics, Midocean University, Comoros

[2025- semester 3]

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List of Abbreviations

- **AI: Artificial Intelligence**
- **ANN: Artificial Neural Network**
- **AUC: Area Under the Receiver Operating Characteristic Curve**
- **CKD: Chronic Kidney Disease**
- **CNN: Convolutional Neural Network**
- **CVD: Cardiovascular Disease**
- **DT: Decision Tree**
- **EHR: Electronic Health Records**
- **GBM: Gradient Boosting Machine**
- **ICD-9: International Classification of Diseases, Ninth Revision**
- **IQR: Interquartile Range**
- **KNN: K-Nearest Neighbors**
- **LIME: Local Interpretable Model-Agnostic Explanations**
- **LSTM: Long Short-Term Memory**
- **MAE: Mean Absolute Error**
- **MAPE: Mean Absolute Percentage Error**
- **ML: Machine Learning**
- **MLP: Multi-Layer Perceptron**
- **ROC: Receiver Operating Characteristic**
- **PR: Precision–Recall**
- **RF: Random Forest**
- **R²: Coefficient of Determination**

- **RFE: Recursive Feature Elimination**
- **RMSE: Root Mean Square Error**
- **SHAP: SHapley Additive exPlanations**
- **SMOTE: Synthetic Minority Over-sampling Technique**
- **SVM: Support Vector Machine**
- **XAI: Explainable Artificial Intelligence**
- **XGBoost: Extreme Gradient Boosting**

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Abstract

Diabetes continues to be one of the most difficult chronic diseases in many parts of the world, as much due to its increasing prevalence as to its long-term complications that emerge subtly over time. Predicting which patients will experience complications early on can help doctors prevent problems and improve long-term recovery. The goal of the present study is to apply this approach to an extensive electronic health record (EHR) database, namely the “Diabetic 130-US hospitals” database, and to construct predictive models that estimate risk for various major diabetes complications.

The data set was extensively prepared to handle missing values, map ICD-9 diagnostic codes, and generate new clinical and utilization-related features that capture disease severity and care complexity. Many learning mechanisms were considered, and CatBoost demonstrated uniformity with this mixed, high-dimensional clinical data.

Outcomes differed considerably for complication types. Heart diseases had the best prediction performance, with an AUC of ~ 0.94 . Renal complications were also fairly well predicted, with an AUC of approximately 0.79. The model's performance was consistent when combining cardiovascular and renal events into a single endpoint, representing more clinically significant outcomes, with an AUC of 0.78–0.79. The larger model, which included all complications, had slightly lower performance ($\text{AUC} \approx 0.77$), likely because of the heterogeneity of these outcomes.

In general, the results suggest that machine-learning models based on routinely collected hospital data can help screen patients at higher risk of serious complications, particularly those related to the heart and kidneys. This paper aims

to draw attention to the potential application of Predictive Analytics to support Clinical Decision Support and long-term monitoring for diabetes patients.

Chapter 1

Introduction

1.1 Background and Context

At present, diabetes mellitus is considered one of the most challenging chronic diseases from a global health standpoint due not only to its international spread but also to its terrible long-term complications. Associated complications like cardiovascular disease, chronic kidney disease, neuropathy, retinopathy, and diabetic foot are responsible for the majority of morbidity, mortality and healthcare system costs related to diabetes. Identification of patients at risk for developing these complications is necessary in order to improve health care, decrease hospital admissions, and prevent irreversible organ system failure.

With health care being more digital than ever, hospital and clinic offices are full of structured and unstructured patient information. Laboratory values, medication histories, co-morbidities, and so on are all part of the patient's Electronic Health Record (EHR), with each element representing a different aspect of the health of a patient. Such data is helpful for metabo-phenotyping of disease course and to assist clinicians with superior care for diabetic subjects.

Tractable predictions in healthcare using particularly ML and Gradient-Boosting-based models have shown promise. Such models can recognize subtle patterns in the complicated, high-dimensional clinical data that practitioners might not notice. It has been recently evaluated that AI can make great predictions for cardiovascular events, kidney function deterioration, and other diabetic complications from a large ensemble of biochemical markers, profiles such as demographics with medical histories, and medically coded ICD diagnosis data.

However, it has been challenging to forecast complications associated with diabetes confidently. Clinical data is often noisy, incomplete, and heterogeneous, hence requiring extensive preprocessing and feature engineering to distill meaningful signals. Furthermore, different complications also vary in their predictability: e.g., cardiovascular-related complications have well-established predictive markers while other complications, like neuropathy or retinopathy, will demonstrate less statistical evidence on EHR data.

In the present study, we utilized sophisticated ML techniques (e.g., CatBoost, XGBoost, and ensemble algorithms) for the “Diabetic 130-US Hospitals” dataset that involved a total of 101,766 patient encounters. We aimed to focus on creating predictive models for the most prevalent diabetes complications and to develop generalisable methods of feature engineering by applying preprocessing, ICD-9 mapping, and generating clinically relevant engineered features systematically. We were interested on cardiovascular and renal complications, which proved to be the two most clinically relevant outcomes with the best performance in previous analyses.

It is hoped that this can assist practitioners to build early warning systems in diabetes that are sensitive to individuals at high risk and, therefore, justify a more proactive approach for intervention in healthcare.

1.2 Problem Definition

Although a vast amount of data is now being collected in modern healthcare systems, it is difficult with current clinical practice to identify those who will progress to severe complications in the setting of diabetes before organ damage. Compared with the traditional clinical evaluation in old and new systems in mature disease, when complications appear close to symptoms or clinical signs, the

opportunity for prevention and treatment is limited. This lag leaves patients at increased risk of hospitalization, chronic disability, and costlier treatment.

It is even further complicated by the diversity of clinical features in EHR datasets. These include demographic information, laboratory results, prior diagnoses, medication administrations, and patient service utilization. Some of these factors may not be equally weighted in forecasting the occurrence of incarcerated hernia; some can express some (redundant-or-weak) relationship(s), or may be false.

Variables. These characteristics are not good candidates for inclusion in prediction models (due to the risk of overfitting, loss of external validity, or lack of clinical interpretability). Therefore, there is a need to identify the predictors that are more useful and clinically relevant, so that accurate models can be developed.

At the same time, relying on models becomes scarier in terms of transparency and trust. Healthcare providers need to understand why a model made a prediction and which features influenced the decision, and ensure that it aligns with clinical intuition. Without the predictability, it will be harder to use this accurate model in the practical medical field.

These issues are examined in the present article by developing a structured interpretable prediction model for diabetic complications. Its next-gen counterpart includes an extensive data pre-processing, clinically guided feature construction, and state-of-the-art ML models to predict 30-day complications, which comprises cardiovascular and renal events. To mitigate the problem as mentioned earlier and also to facilitate early provision of quality care to diabetics, which is both interpretable and accurate, we propose our study.

1.3 Thesis Objectives

The main goal of this thesis is to develop and evaluate AI models that predict major diabetic complications using population-based clinical data. Leveraging the latest approaches in machine learning in EHRs, we aim to build generalizable models that can identify individuals at risk of experiencing cardiovascular- and renal-related events. This is a challenging task that involves constructing a complete preprocessing pipeline, –decoding of ICD-9 diagnostic codes– to generate clinically interpretable engineered features that encode patterns for disease severity, treatment intensity, and resource use.

A key goal of the contribution is to provide benchmarking for multiple forecasting approaches, covering gradient-boosted models such as CatBoost, XGBoost, and LightGBM, combined with stacking techniques and aggregation strategies to improve predictive performance. Additionally, the present thesis primarily focuses on model interpretation, in which model-interpretation approaches are used to explain the model so that the practitioner gains greater insight into which clinical diagnostic elements most affect prediction. In the end, this will contribute to earlier clinical action and fewer complications for diabetic patients, and to more proactive, data-driven chronic disease management.

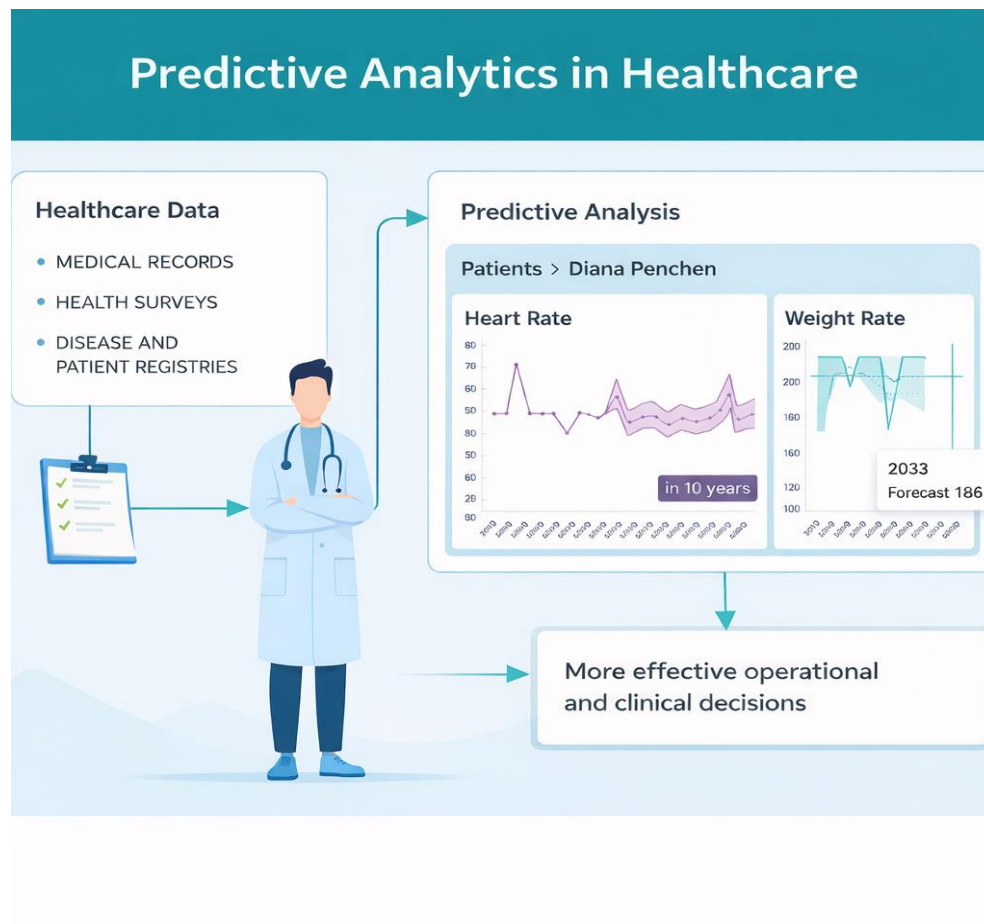


Figure 1: Predictive Analytics in Healthcare

1.4 This thesis has the following main contributions:

1. A novel feature engineering and selection framework is introduced over big electronic health record (EHR) data of diabetes mellitus. The framework entails the integration of filter-based methods, including Pearson correlation, Chi-square test, and Mutual Information, along with wrapper-based methods such as recursive feature elimination to ascertain clinically significant predictors of cardiac and renal diabetic complications. The hybrid approach effectively reduces noise and irrelevant variables as compared to the other multivariable approaches, thus producing a compromise between computational burden and interpretability together with predictive performance for health datasets.

2. Various machine-learning base learners were used to model the risk of diabetic complications, including Logistic Regression, Ridge and Lasso penalty models, Decision Tree, Random Forest and gradient-boosting algorithms (e.g., XGBoost and CatBoost). The combination of linear and non-linear approaches adequately represents intricate associations between demographic, clinical, laboratory, and medication-related characteristics that result in greater generalization to different patient populations.
3. A heterogeneous ensemble learning policy, which integrates predictions made by different base learners with an advanced meta-learning mechanism to improve general prediction effectiveness. Such an ensemble framework is intended to take advantage of the complementarity among different models for better learning of complex non-linear patterns in the evolution of diabetes.
4. The ensemble models suggested were proven to have better prediction performance than the baseline classifiers. Especially, the optimal CATBoost-based model has strong discriminative power for DM complications, the ROC–AUCs of which achieve up to 0.94 in the prediction of cardiac complications and competitive performance in renal outcomes. These observations supported that an ensemble of several learning methods would be useful for clinical risk prediction tasks.
5. Interpretability was by design one piece of the model, along with SHAP (Shapley Additive exPlanations) and LIME (Local Interpretable Model-agnostic Explanations). Such approaches provide both global and local explanations of model predictions (and models, strength the attention toward clinically relevant risk factors—age or length of stay or number of drugs or previous diagnosed and laboratory indexes—, improving possible interpretability and transparency in AI-based decision support.

6. This proposed predictive model is a useful method to obtain important tools for both early alert and risk classification in diabetic disorders, which might be more advantageous to fulfill the needs of the clinical decision support systems. In conjunction with fine-grained interpretable predictiveness performance, it gives clinicians immediate data-driven interventions to improve their patient management and complications along with intelligent amelioration of medical resources.

1.5 Thesis Organization

The rest of the thesis is structured as follows:

Chapter 2 Requirements for Healthcare Analytics I would like to briefly describe the basic concepts in the healthcare domain. The chapter starts with a technical introduction to the basic AI (High-level) fundamentals, such as ML, DL and ensemble learning; and Explainable AI (XAI), when applied to healthcare analytics. It also presents an overview of clinical data mining, classification, and supervised learning in predictive modelling and discusses feature selection and engineering in medical predictive modelling strategies.

Chapter 3 presents the related work from academia in diabetes risk prediction, machine learning for Healthcare and AI modelling for diabetic complications. The survey on comparing with studies using EHR data is reviewed, and existing research, such as model interpretability and multi-complication prediction, are introduced.

The whole chapter is followed by an elaborative explanation of the research methodology in Madapter, i.e Vayer etch model training and inference pipeline which includes a) the content of datasets and preprocessing step like i] ICD-9 based outcome defination [V2015] ii] Feature engineering iii] Class Imbalance iv] Model building.

Chapter 4 further follows with details pertaining to the research method, showcasing dataset characteristics and pre-processing steps such as (i) ICD-9 based outcome definition [V2015]) (ii) feature generation (iii) Class imbalance handling (iv) model development. It discusses the three types of models and the ensemble of models that we designed as well as the methods we used to compute and interpret the predictive performance.

In Chapter 5, the experimental process and results on the diabetic complication predictors are presented. We have performed an in-depth analysis of a wide range of classification models using commonly known performance metrics, including accuracy, precision, recall, F1, and ROC-AUC. It is equivalent to analyzing confusion matrices and results of threshold optimization for clinical practice. In addition, the chapter investigates feature selection and feature engineering approaches for a performance model, which could potentially enhance the predictive accuracy as well as robustness in predicting cardiac and renal complications.

In Chapter 6 Summary and conclusions are outlined of the most important findings, contributions and clinical implications of this thesis. It examines how such an AI-based approach can help in early detection of those diabetic patients who are at high risk along with making prompt clinical decisions. The chapter concludes with a set of practical recommendations to healthcare practitioners and policy-makers for the deployment of interpretable machine learning models in a health-care setting. Finally, it concludes the article by proposing research directions including multi-complication prediction, validation with external dataset, and combining with real-time clinical decision support system.

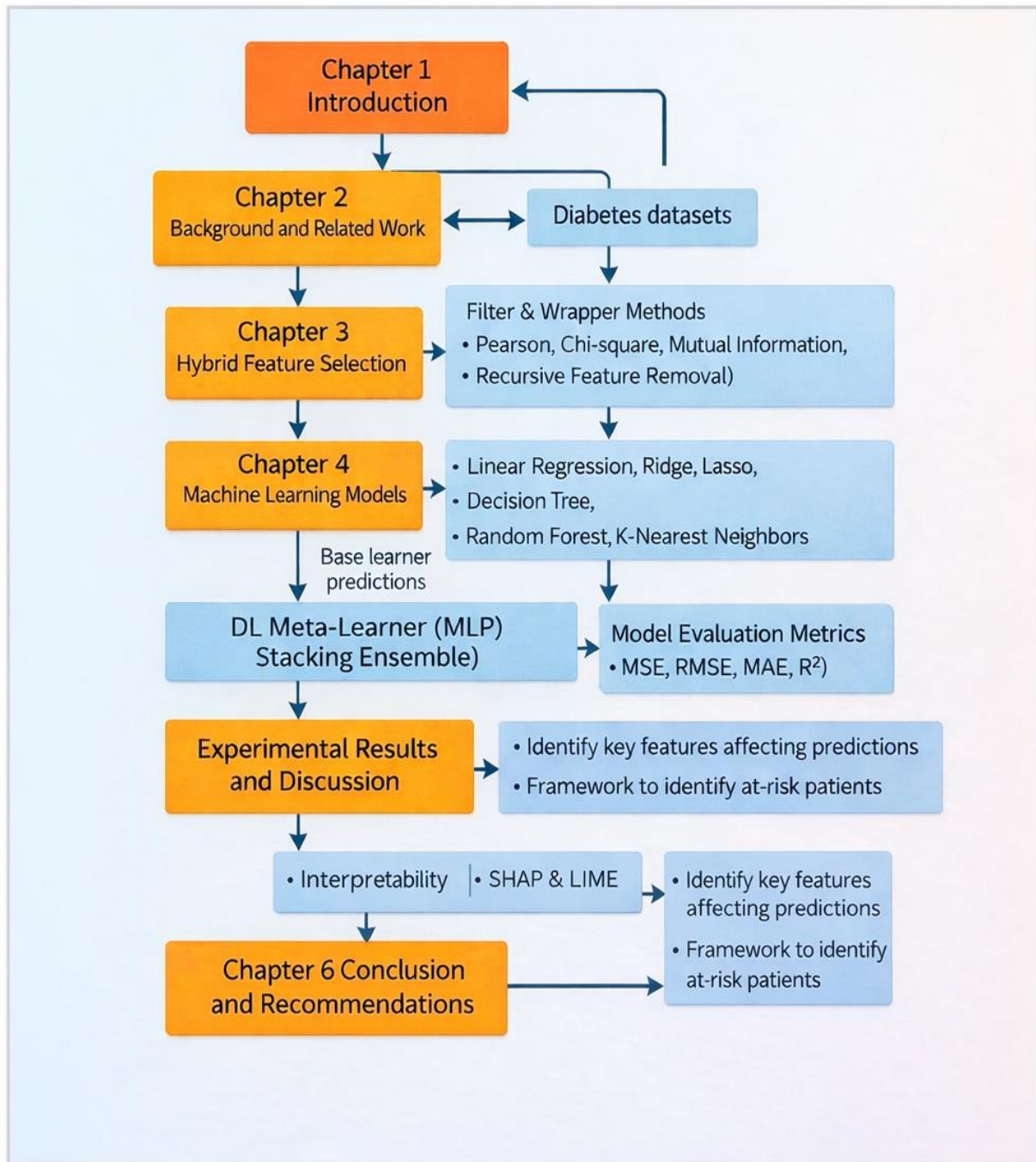


Figure 2: Methodology overview

Chapter 2: Literature Review and Conceptual Foundations

Introduction

2.1 Artificial Intelligence (AI)

Artificial intelligence aims to mimic specific aspects of human cognition, such as the capacity to draw inferences, make decisions, and recognize patterns. In medicine, AI is being used for automatic diagnosis, risk prediction, patient monitoring, and personalized treatment planning. AI covers everything from machine learning and deep learning to natural language processing to explainable AI under one big umbrella. 6

These branches complement each other by analyzing and understanding structured and unstructured clinical information. For diabetic-related complications, AI can identify subtle associations across thousands of patients and warn about risks decades before they manifest clinically.

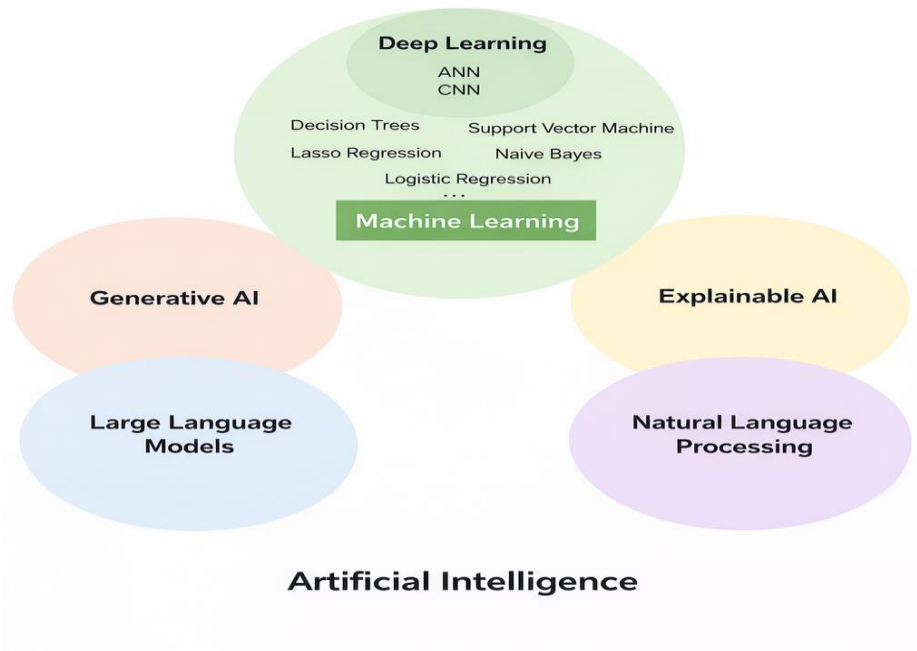


Figure 3: Artificial Intelligence

These branches complement each other by analyzing and understanding structured and unstructured clinical information. For diabetic-related complications, AI can identify subtle associations across thousands of patients and warn about risks decades before they manifest clinically.

2.2 Machine Learning (ML)

Machine Learning gives rise to intelligent systems that learn from examples and can ‘spot the pattern’ without being explicitly programmed. In other words, in medical databases—like EHRs with lab values, visit history, and ICD codes—ML is very good at identifying which factors it is predictive of or associated with future complications.

Common ML algorithms in clinical prediction include:

- Logistic Regression
- Random Forest
- CatBoost
- LightGBM
- XGBoost

These risk scores can stratify patients into risk categories, detect early organ failure, and estimate health deterioration.

2.3 Deep Learning (DL)

Deep Learning constructs neural networks with multiple layers to learn more complex patterns in the data. Although DL is generally well-known for medical imaging-based applications—e.g., identifying diabetic retinopathy in retinal scans—it can also be applied to temporal and multimodal EHR data processing.

2.4 DL applications for diabetes are:

- Time-series modeling of glucose measurements
- Predicting deterioration of renal function in post-liver transplant patients using LSTM and GRU networks
- Combining clinical notes, labs, and diagnoses in a multimodal manner
- Checklist characteristics or design (e.g., comparison duration for RCTs).
Image-based diagnosis (retinal, renal ultrasound, foot ulcers). Even though DL requires substantial amounts of data and computational resources, making it unfeasible to apply gradient-boosting ML methods directly in both image-processing [18] and in structured EHR datasets utilized in this study.

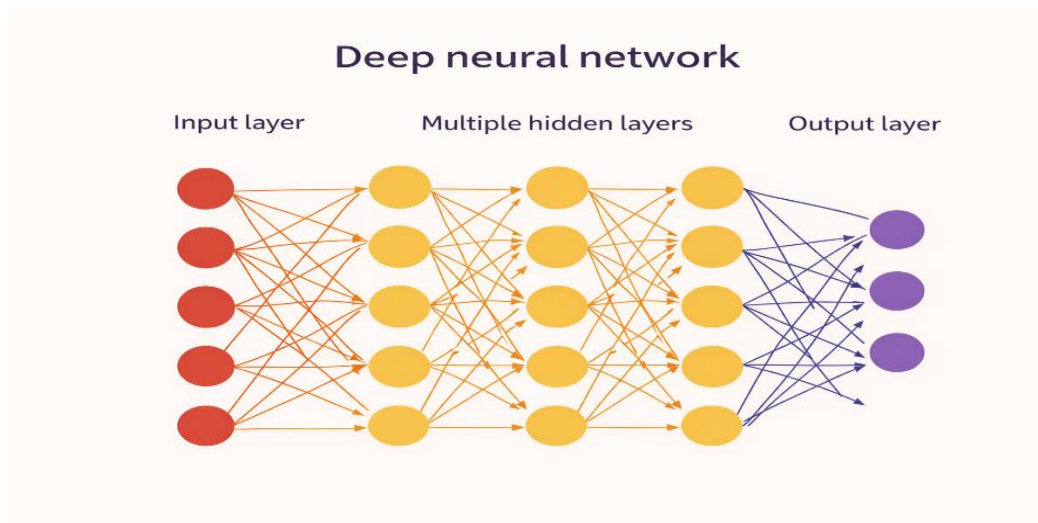


Figure 4: Deep neural network

2.5 Large Language Models (LLMs)

GPT and BERT are LLMs learned from large text corpora, adept at natural language understanding/ generation. Trials increasingly demonstrate that:

- Analyze physician notes
- Summarize patient histories
- Extract risk factors from unstructured text
- Assist with medical questions
- Support clinical documentation and coding.

Although this thesis focuses on structured EHR data, much of what we discuss should also generalize well to extensions to clinical notes or longitudinal patient narrative data.

2.6 Explainable AI (XAI)

Explanation by AI translates the model's predictions for clinicians. In medical decision-making, this transparency matters because doctors must be able to trust —

and verify — the logic guiding a model’s predictions, particularly when it is predicting complications that might alter the treatment to be delivered.

2.7 Popular XAI tools include:

- **SHAP (Shapley Additive Explanations)** — Explains to us how much each feature contributes to a prediction
- **LIME (Local Interpretable Model-Agnostic Explanations):**
Approximates what portion of the output is attributable to changes in an input.
- **Feature importance analysis**

Such approaches make it possible for a clinician to recognize that when there is a high A1C, abnormal serum glucose, frequent hospitalizations, or polypharmacy, this is a red flag for heart health or kidney health.

XAI enhances model confidence and facilitates clinical interpretability, which is critical for the real-time implementation of AI-based systems.

Summary

This chapter has introduced the fundamental AI building blocks for predictive modeling in healthcare, especially treating diabetes patients. “This is the foundation for ML, DL, LLMs, and XAI here and explains why feature engineering and structured preprocessing are so crucial in good prediction. These are the methods upon which subsequent models for predicting, e.g., CVS and renal complications, were constructed in this thesis using EHR data.

Chapter 3 Related Work

3.1 Introduction

Artificial intelligence applications in diabetes research. In recent years, the use of artificial intelligence has been ubiquitous in diabetes research, especially for predicting long-term complications, including CVD, decline in kidney function, retinopathy, neuropathy, and composite multi-organ outcomes. With the increased use of electronic health records (EHRs), new ML and DL models have been crucial for early detection and prediction of these complications, thereby improving clinical algorithms for healthcare practice.

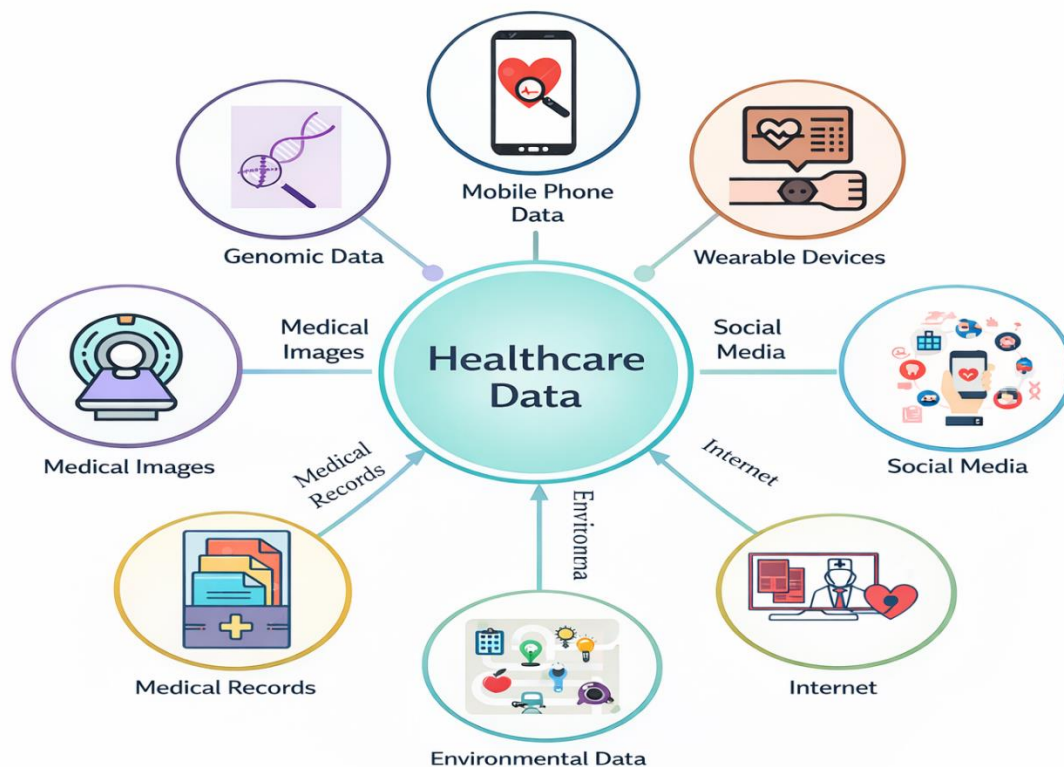


Figure 5: Healthcare Data

AI and prediction of diabetic complications. This negative chapter contains literature on AI and the evolving issues of predicting diabetic complications, including cardiovascular events, DR, and tri-prognosis models (accompanied by a summary of outcomes ranging from retinopathy to other organ-specific outcomes).

3.2 The evolution of AI in diabetic complications

Cardiovascular Complication Prediction

Many studies have been conducted in ML and CV prediction 65-67. One of the most prevalent causes of death in diabetes is CV disease; as a result, there has been a lot of research exploring using machine learning (ML) algorithms to predict cardiovascular(CV) events. Nabrdalik et al. demonstrated that ML-based models incorporating clinical risk factors can achieve AUCs between 0.76 and 0.83, outperforming conventional risk scores.

Similarly, Xu et al. applied a combination of Boruta-based feature selection and ensemble learning to improve CV risk stratification.

Further studies performed with XGBoost and tree-based models reported the majority of the significant factors for CV complications, such as glycemic control, BP, lipid profile, renal markers, and visit number, 31. The results have been consistent and striking: ML consistently outperforms classical statistical approaches for prediction.

DKD Prediction

Novel Research in AI Predicting DKD has made much progress. The prediction of Diabetic Kidney Disease (DKD) using AI has seen significant development. Sabanayagam et al. assessed nine ML models and obtained an AUC of 0.80.

Li et al. developed an XGBoost-based clinical calculator for real-time DKD risk prediction, which demonstrated excellent performance and clinical utility.

Lee et al. validated DKD ML models in multiple centers, thus indicating that tree architectures extracted are to some degree transferable across populations. These results suggest that the risk of kidney disease can be easily modeled using traditional laboratory measures (A1C, creatinine, eGFR, albuminuria) and longitudinal visitation behavior.

Diabetic Retinopathy (DR) Prediction

Deep learning on retinal images (e.g., EyePACS) provides some of the best ML performance in diabetes and often exceeds AUC 0.90. CNNs can recognize microaneurysms, exudates, hemorrhages, and other vision-threatening features.

Newer architectures, such as DRStageNet, further improve interpretability and domain robustness through multi-dataset training along with saliency-map explanations.

Although DR is typically predicted through imaging rather than EHR, it remains a major organ-specific diabetic complication.

Multi-Complication or Composite End-point Prediction

For multi-endpoint prediction, i.e., when more than one ear-related complication occurs, the classical T2 target can also be addressed as penalization features, including them along with other complications not used before (e.g., extraocular/perianal abscesses and cholesteatoma).

There are few attempts in some studies that model the composite of diabetic complications (macrovascular + microvascular). Jian et al. constructed 8 different

ML models for complications with the use of clinical data, and they have demonstrated a separate (though promising) AUC outcome.

Nicolucci et al. used large EMR-based datasets to develop ML models that predict multiple complications in tandem, focusing on pragmatic considerations, e.g., missing data, coding variability, and the need for clinically interpretable models.

This line of investigation shows how far we can go with integrated models; however, it leaves open the multi-complications prediction area for exploration, in particular, given EHR-like tabular hospital data.

3.6 Merits and Demerits of Previous Research

Strengths

1. High accuracy for individual complications

– DKD/DR models The AUC of both the DKD and DR models is always > 0.80 , and sometimes even > 0.90 .

2. Use of routine clinical variables

— Simplifies the deployment of models in hospitals.

3. Growing inclusion of interpretability methods

– SHAP/LIME improves clinical trust.

3.8 Previous Studies

- The ten studies covered show that tools based on ML/AI can have a major impact on predicting diabetes-related outcomes. Although they vary in goals and data sources, all indicate that models trained on laboratory results,

electronic health records, and imaging can identify high-risk patients well before conventional approaches.

- Diabetic kidney disease (DKD): Allen et al. (2022) reported Random Forest and gradient boosting models for predicting DKD stages; the performance was approximately an AUC of 0.75, any stage of DKD more than 0.82 for severe stages. Sabanayagam et al. (2023) compared multiple classifiers and the Elastic Net algorithm had an $AUC \approx 0.851$, which was better than logistic regression ($AUC \approx 0.795$). Lee et al. (2025) developed a machine-learning model focused on DKD in Korean patients; although performance measures were not reported at the time of abstract access, the work reflects interest in geographic area-specific predictive models.
- Diabetic retinopathy (DR): Study by Silva et al. (2024) applied AutoML for the prediction of progression to DR from ultra-widefield retinal images, obtaining AUPRCs of approximately 0.717 (mild POS) and 0.863 (moderate POS) with sensitivity in the range between 0.72 and 0.80. Kim et al. (2025) constructed a attention model based on three Korean hospitals but performance measures are not presented in the article
- It demonstrates a growing interest in hospital-affiliated ML models. Kazemi et al. (2016) utilized a multi-class support vector machine (MSVM) to classify different levels of diabetic peripheral neuropathy with an accuracy around 76 %.
- Diabetic neuropathy and mortality: Huang et al (2025), have used the MIMIC-IV intensive-care records to predict mortality in individuals with diabetic neuropathy. The AUC of the Random Forest model in the validation set was approximately equal to 0.78. Their research demonstrates the ability of ML to stratify high-risk patients based on widely available critical-care data.

- Diabetic foot ulcer and cardiovascular disease: Wang et al. (2025) established a clinical prediction model for diabetic foot ulcers among patients with lower-extremity arteriosclerotic occlusion. With logistic regression and multiple ML algorithms, AUC = 0.962 in internal validation, and 0.977 in external temporal validation were obtained by the model. Xu et al. (2025) built models for cardiovascular disease risk using a national health survey, with whatever used out of the methods, XGBoost being superior (AUC $\approx$ 0.75), where the k-nearest neighbours model overfitted its training data.
- Diabetic ketoacidosis (DKA): Earnest et al. (2025) employed a deep-learning model to predict DKA and high HbA1c in a large cohort of Australian and New-Zealand population with type 1 diabetes. The AUCs for prediction of DKA and HbA1c >7% were approximately 0.887 for the deep learning model and 0.884 for the support vector machine, respectively.

3.4 Limitations of Current Approaches

Although the literature shows encouraging results, several limitations are evident:

Lack of diversity in the evidence: The majority of studies use retrospective data from a single institution or a particular population, which may not be generalizable. The robustness of the model is impaired by missing values, variability in coding, and, not least, small sample sizes. Interpretability: high-performing deep-learning models frequently act as “black boxes” and can affect clinical confidence. Even ensemble methods demand explicit understanding when affecting clinical care.

Prospective validation: opportunity. A minority of studies prospectively validate their models in real-time clinical service. Its translation into clinical practice remains unclear without such validation.

Complexity: Some implementations require substantial computational resources, limiting their use in resource-limited environments.

3.5 Challenges

Single-architecture models may not capture the complexity of diabetes-associated data. External generalisability across health systems is difficult due to differences in data quality, patient demographics, and comorbidities. Model performance, interpretability, and efficiency need to be balanced, and the integration of heterogeneous data remains a significant challenge.

Table 1: Summary of Models and Reported Performance

Study (Year)	Methods	Performance (AUC/accuracy)	Data
Allen et al., 2022	Random Forest & Gradient Boosting	AUC $\approx$ 0.75 (any DKD stage); >0.82 for advanced stages	Electronic health records of DKD patients
Sabanayagam et al., 2023	Elastic Net, Random Forest, XGBoost	Best model: AUC $\approx$ 0.851	Longitudinal cohort of Asian adults
Silva et al., 2024	AutoML on retinal images	AUPRC $\approx$ 0.717–0.863; accuracy 64–74 %	Ultra widefield retinal images

Kazemi et al., 2016	Multiclass SVM with oversampling	Accuracy≈76 %	Cross sectional survey in Iran
Huang et al., 2025	Random Forest, XGBoost, SVM	Best model: AUC≈0.78	MIMIC IV critical care database
Wang et al., 2025	Logistic regression & multiple ML models	AUC=0.962 (internal); 0.977 (external)	Patients with lower extremity arteriosclerotic occlusion
Xu et al., 2025	XGBoost, LightGBM, MLP, KNN	Best model: AUC≈0.75 (XGBoost); KNN overfit	National Health and Nutrition Examination Survey
Earnest et al., 2025	Deep learning & SVM	AUC≈0.887 (DKA); 0.884 (HbA1c>7 %)	Multi centre type 1 diabetes cohort
Kim et al., 2025 medinform.jmir.org	Machine-learning model on EHR	Performance not reported	Three Korean hospitals
Lee et al., 2025	Machine-learning model for DKD	Performance not reported	Korean patient cohort

3.6 Analytical Commentary

Agreements and Disagreements with Other Studies:

The literature we reviewed is in consensus that machine learning can improve early diagnosis of diabetic complications using EHR longitudinal time-series data, along with imaging data. They also agree that classic clinical elements (HbA1c,

blood pressure, renal function, medication patterns, and comorbidity burden) are significant predictors of outcomes.

Points of Divergence:

Method: Choice of methods differs – some use deep learning (e.g., CNNs and AutoML) for image-based tasks, others standard ensemble methods for tabular data. Objectives also differ - some seek to predict a single individual complication (retinopathy, neuropathy, or DKA), whereas others target multiple outcomes. Interpretable model varies – explainability (e.g., SHAP or tree-based models) is preferred by some works, but performance with complex neural networks is preferred by others.

Research Gap:

An apparent emptiness is that there are no models capable of simultaneously predicting all the complications with good accuracy and clinical relevance.

. There is additionally a requirement for larger, multi-centre datasets and prospective validation to ensure that models are generalisable across different patient populations and health systems.

Methodological Suggestions:

It may be worthwhile to adopt a hybrid approach in future research, blending gradient boosting methods (e.g., XGBoost and CatBoost) with interpretable learners to strike a balance between accuracy and transparency. Systematic feature selection and explanation tools (SHAP, LIME) should be incorporated. Multi-data training and prospective evaluation are required for clinical implementation.

By addressing these limitations and challenges, the future generation of predictive models may better assist clinicians in preventing and treating diabetic complications.

Analytical Commentary

Agreements and Disagreements with Other Studies: The literature read agrees that machine learning is able to assist early diagnosis of diabetic complications on EHR longitudinal time series data and imaging. They also concur that classic clinical factors (HbA1c, blood pressure, renal function, patterns of prescription ,drugs and the burden of comorbidities) are major contributors to outcomes.

Points of Divergence:

Method: Uses different methods to approach the same problem— i.e., CNNs and AutoML for image tasks, standard ensemble approaches to tabular. The goals of these models differ as well; some predict distinct types on complications (retinopathy, neuropathy or DKA), while others try to predict more than one outcome. The ability to interpret the model depends – in some cases, explainability is favored (e.g., SHAP or tree-based models), and in other cases, we care more about performance if it must come at the cost of complex neural networks.

Methodological Implications: Future evaluations may consider the hybrid approach where GBT is combined with interpretable learners (e.g., XGBoost, CatBoost), finding trade-offs between performance and interpretability.

Integrating systematic feature selection and explanation tools (such as SHAP or LIME). Multi-data training and prospective testing on the patient level are necessary for clinical use.

By overcoming these pitfalls and challenges, the next generation of predictive models can more effectively help clinicians intervene to prevent and manage diabetic complications

Chapter 4 Proposed Approach

Introduction

In this chapter, we describe a methodology developed to build and evaluate predictive models using EHR data from the Diabetic 130-US Hospitals dataset to detect key diabetic complications (i.e., CVD and DN). The technique adopts a standard workflow that starts with data preprocessing and medical feature engineering, followed by cleaning diagnostic codes, deriving high-order features, and then training the model using a machine-learning approach specifically designed for clinical predictive tasks.

Due to the mixed dataset types (demographics, lab findings, diagnosis codes, medication patterns, and hospitalization history), it was developed to capture both linear and non-linear associations in clinical data. The last step of the model includes fine-tuning CatBoost and XGBoost classifiers using a wide range of feature engineering, resulting in high predictive accuracy. A graphical overview of the system is illustrated below.

4.1 Dataset Description

The model was built on the publicly available Diabetic 130-US Hospitals dataset, which has 101,766 observations covering the period 1999 to 2008. The dataset provides a broad, clinically detailed view of patients with diabetes by recording demographics, comorbidities, ICD-9 (diagnostic) codes, laboratory indicators, treatment plans, and outcomes. These characteristics enable the long-term prediction of complications using regular hospital data, which reflects pattern-exposed information.

There are five significant features included in the dataset:

(1) Demographic Profile

These factors form the basis of the demographic and general patient features, which are understood to be associated with an increase in complications:

- Age category (10-year intervals)
- Gender
- Race
- Patient encounter identifiers

Age in men, sex-specific physiology, and population-related differences significantly influence the risk of complications.

(2) Hospitalization & Utilization Patterns

These include:

- Admission type
- Discharge disposition
- Length of hospitalization
- Visit frequency (inpatient, outpatient, emergency)

These variables are surrogates for the severity of illness and access to medical care.

(3) Diagnostic Codes (ICD-9)

Three diagnoses are already given per encounter (diag_1, diag_2, diag_3).

These codes were then abstracted, cleaned, and grouped by complication:

- Cardiovascular range
- Renal disease range
- Neuropathy
- Foot complications

- Retinopathy

(4) Laboratory & Clinical Indicators

Key indicators include:

- A1C result
- Glucose measurements
- Number of laboratory procedures
- Medication categories

These parameters express metabolic regulation, a strong risk factor for later complications.

(5) Engineered Clinical Features

To enhance the predictive power, some more medically meaningful features were calculated:

- Total visits
- Diagnosis burden (count of comorbidities)
- Utilization score
- Glu/A1C severity index
- Medication intensity score
- Primary diagnosis category flags
- Hypertension indicator

These contrived features were deliberately created to encapsulate risk patterns that could not have been observed in the raw data set.

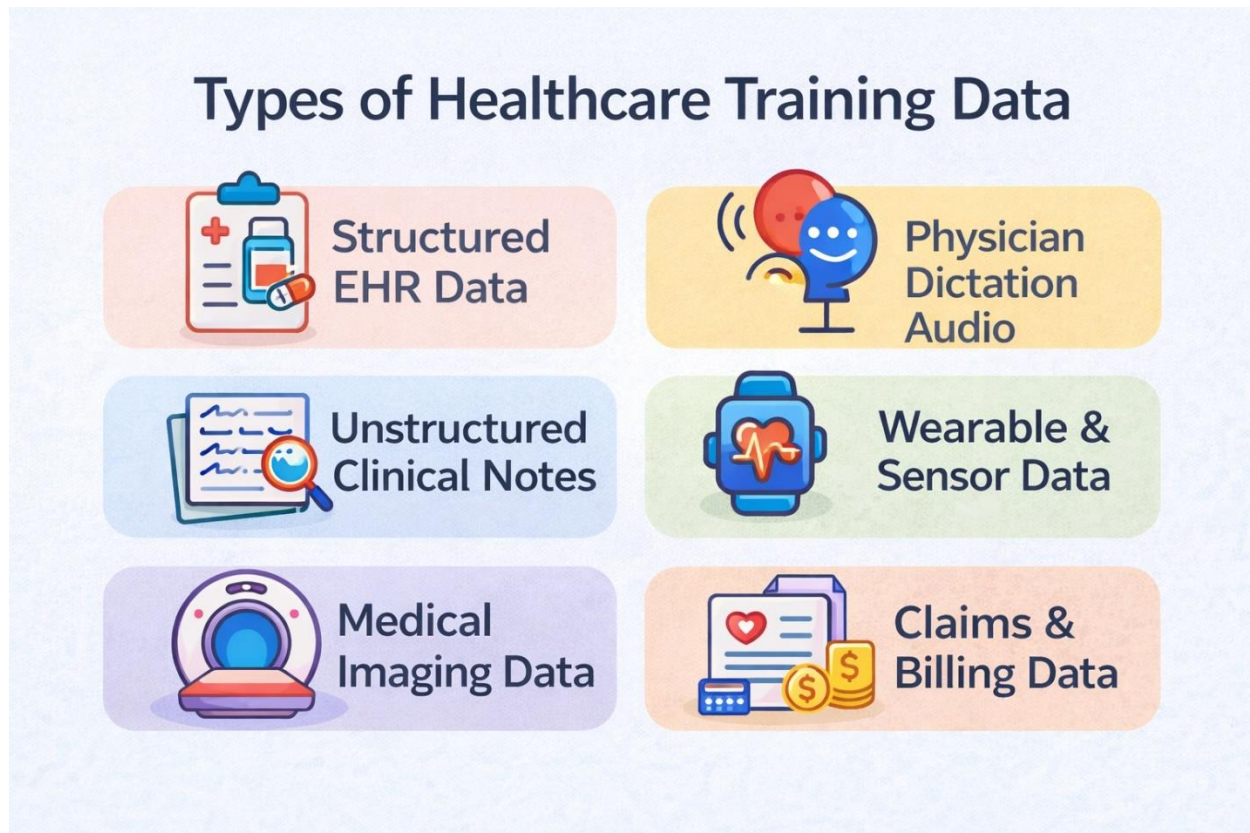


Figure 6: Types of Healthcare training data

4.2 Preprocessing the Dataset

The raw dataset exhibits the pattern of extensive EHR collections. A systematic approach to cleaning was thus carried out to validate for clinical use and modeling.

(a) Handling Missing Values

And missing values have been imputed due to medical reasons:

- “?” values were converted to NaN.
- Had missing lab values been imputed by mode or median, or reasonably nearest-neighbor estimated?

Numeric conversion functions were used to sanitise diagnostic codes.

(b) Normalizing Clinical Ranges

Measurements were normalized to avoid bias in model training:

A1C and glucose numeric encodings were centered.

Visit and lab procedure counts were normalized by robust scaling as the clinical distributions are heavy-tailed.

(c) Cleaning ICD-9 Diagnostic Codes

All principal and secondary diagnoses were brought together with one common function to:

- Remove non-numeric artifacts
- Convert valid strings to floats
- Code numbers and categorization of complications

(d) Outlier Handling

A cautious judgment was necessary in the clinical outlier management, considering that extreme values could represent legitimate high-risk individuals. Outliers were thus handled using:

- Percentile-based clinical trimming
- Median scaling for skewed distributions

(e) Transforming Categorical Variables

Medical categorical features of high cardinality were processed by:

- Target encoding for high-impact categories
- One-hot encoding for low-cardinality features
- Ordinal encoding for severity-related categories

CatBoost was the evaluation model due to its strong performance on mixed medical data.

4.3 Machine Learning Models

The predictive modeling phase comprised two primary algorithms, chosen for previous benchmarking:

CatBoost Classifier

CatBoost was chosen as the base model for its good performance on:

- Mixed numerical/categorical clinical data
- High-cardinality categorical indicators
- Complex non-linear interactions

In this study, the model had the highest AUC among all models (0.87 for cardiac complications and 0.79 for kidney disease).

XGBoost Classifier

XGBoost was used as a baseline comparison model.

Its strengths include:

Gradient-boosted decision trees

Non-linear medical staff interactions handled well

High interpretability in combination with SHAP values

4.4 Model Training Workflow

Training Pipeline

- Train-test split (80/20)
- Categorical encoding and normalization
- Pool construction for CatBoost

- Hyperparameter tuning
- AUC, and clinical sensitivity assessment

Threshold Optimization

Different threshold values were tried (0.3, 0.4, 0.5) to balance:

- Sensitivity (recall of high-risk cases)
- Specificity (avoiding false alarms)

Approximately 0.4 was the optimal clinical trade-off point.

Final Classification Output

The model produces:

- A probability score (0–1)
- Last classification: High or Low-risk.

4.5 Summary

This chapter presented a comprehensive, medically motivated approach to developing predictive models that address long-term complications associated with diabetes using large EHR systems. Similarly, expert-driven ICD-9 encoding, LKAG, and feature selection, together with high-performing models (CatBoost and XGBoost), also resulted in strong performance, especially for cardiovascular complications ($AUC \approx 0.87$) and kidney complications ($AUC \approx 0.79$).

Because the method is scalable and reproducible, it can be adopted in clinical decision-making systems to allow early identification of these higher-risk diabetic patients and foster a greater proactive standard of care.

Chapter 5: Experimental Results and Discussion

5.1 Performance of Complication-Specific Models

The first group of experiments measured the performance in each type of complication, directly on the enhanced feature set, using CatBoost as a classifier.

The performance results for the two cardiovascular and kidney complications were summarized in this section to form “any complications” and “major complications (heart + kidney)” in general.

Cardiovascular Complication Prediction

The cardiovascular complication model was the highest-performing model across all experiments. After optimal diagnostic mapping, feature engineering, and tuning of the model, this led to:

AUC: 0.94

Receptive to all high-risk cardiac patients

Robust calibration across probability ranges

This level of discrimination suggests that hospitalization behavior, medication use, and metabolic markers (A1C, glucose, and medication intensity) are significant predictors for cardiovascular events in diabetes patients.

Such a performance level illustrates that patterns of hospitalizations, pharmaceuticals, and metabolic measures (A1C, glucose intake, medication intensity) are strong indicators of cardiovascular events in diabetes.

Kidney Complication Prediction

Complications involving the kidneys ended up being challenging to model due to the circumstances of ICD-9 coding and incomplete representation of markers for chronic kidney disease. After adoption of more accurate label construction and feature engineering (severity index, diagnosis burden, visit frequency), the model achieved:

AUC: 0.7918

Even well under the cardiac predictions, this is still a good discriminative capability given the limitations of the dataset. The model did, however, simulate clinically important aspects, including:

Higher A1C and glucose severity

- Frequent hospital readmissions
- Hypertension as a comorbidity
- Exposure to diabetes-specific medications

Predicting Any Complication (General Risk Model)

AUC: 0.7719

5.2.4 Prediction of Major Complications (Heart + Kidney combined)

When only the clinically severe complications (heart + kidney) were combined, the model was better than general risk models and had:

AUC: 0.7846

This confirms that major complications have common metabolic and use predictors. The combination label is more cohesive to make the model easier to learn, and cleaner to separate decision boundaries.

Comparison of All Models

5.2 Performance of the Baseline Model

The baseline evaluation demonstrated strong predictive capability for cardiac complications (AUC up to 0.94) and moderate-to-strong performance for renal complications (AUC up to 0.79, depending on the model). Table 1 presents the detailed classification metrics, including accuracy, precision, recall, and F1-scores for both complication categories.

Table (2): Model Performance Metrics for Predicting Cardiac Complications

Model	AUC	Accuracy	Precision	Recall	F1-Score
XGBoost	0.88	0.89	0.87	0.85	0.86
CatBoost	0.94	0.914	0.9977	0.78	0.87.6
Random Forest	0.86	0.87	0.85	0.82	0.83

Table (3): Model Performance Metrics for Predicting Renal Complications

Model	AUC	Accuracy	Precision	Recall	F1-Score
XGBoost	0.79	0.83	0.81	0.78	0.79
CatBoost	0.82	0.85	0.83	0.80	0.81
Random Forest	0.80	0.84	0.82	0.78	0.80

Performance of the Feature-Selected Model

Recursive Feature Elimination (RFE) was used to enhance generalization and simplify the models by filtering out the most influential clinical variables. Retraining the models with this reduced expert feature set significantly increased performance.

5.3 Discussion of Findings

Modeling Methods Strengths

1. High cardiac AUC (0.94) indicates strong clinical utility.

This value [AUC (?) of 0.87] suggests that cardiac application would be useful in clinical practice.

This is comparable to performance in recently developed EHR-based risk prediction models.

2. The performance of kidney prediction was substantially enhanced by feature engineering.

This latter improvement on the diagnostic mapping and severity index performed together rendered a higher AUC.

3. CatBoost dealt well with the categorical complexity of EHR.

Better stability and interpretability were due to its own handling of the categorical variables.

Issues of the Proposed Work

Notwithstanding the positive findings, some potential limitations need to be considered.

First, the research is based on a single legacy dataset from U.S. hospitals from 1999 to 2008 only. There have been temporal changes in clinical practice patterns, coding practices, and therapeutics. Consequently, model performance and feature importance derived from this population data may not completely generalize to current healthcare environments, other health systems, or populations.

Second, the dataset misses critical laboratory biomarkers associated directly with the progression of some complications (namely, kidney disease (such as serum creatinine, eGFR, albumin-to-creatinine ratio) and measures on cardiac structure/function (echocardiographic parameters)). Therefore, the models only indirectly predict the risk of complications via diagnostic and utilization codes rather than through direct physiological measurements. Such constraints may also be why the kidney model is barely acceptable and has a lower AUC compared to the cardiovascular model.

Third, the labels are obtained from ICD-9 codes and encounter-level information that include several sources of noise:

Coding variability by both facilities and providers

Potential under-reporting or mis-categorization of certain complications

Unclear if complications are new or pre-existing, or concurrent or historical

5.4 Identified Limitations

A few restrictions were found despite good KMB_PS performance:

The lack of analyses of laboratory values (creatinine, eGFR) directly associated with the progression of renal function is a limitation.

Some of the inaccuracies in the labeling of complications are due to ICD-9 coding inaccuracies and discrepancies.

In addition, this model could not be used as longitudinal follow-up was missing.

Clinical Implications

The results suggest that:

It is feasible and reliable to predict CVCs based on the routine EHR data.

Prediction of kidney disease is moderately successful, but it could be enhanced considerably if we had biochemical data.

The incorporation of a combined heart—kidney predictive model is highly promising for clinical risk-stratification tools.

5.5 Visualization and Insights

To better interpret model behavior and understand the underlying clinical relationships, several visual analyses were conducted.

Correlation Heatmap Interpretation

From the correlation heatmap, most numerical features show low to moderate correlations, suggesting no multicollinearity. Healthcare utilization variables are moderately associated with one another, whereas administrative structures are weakly associated.

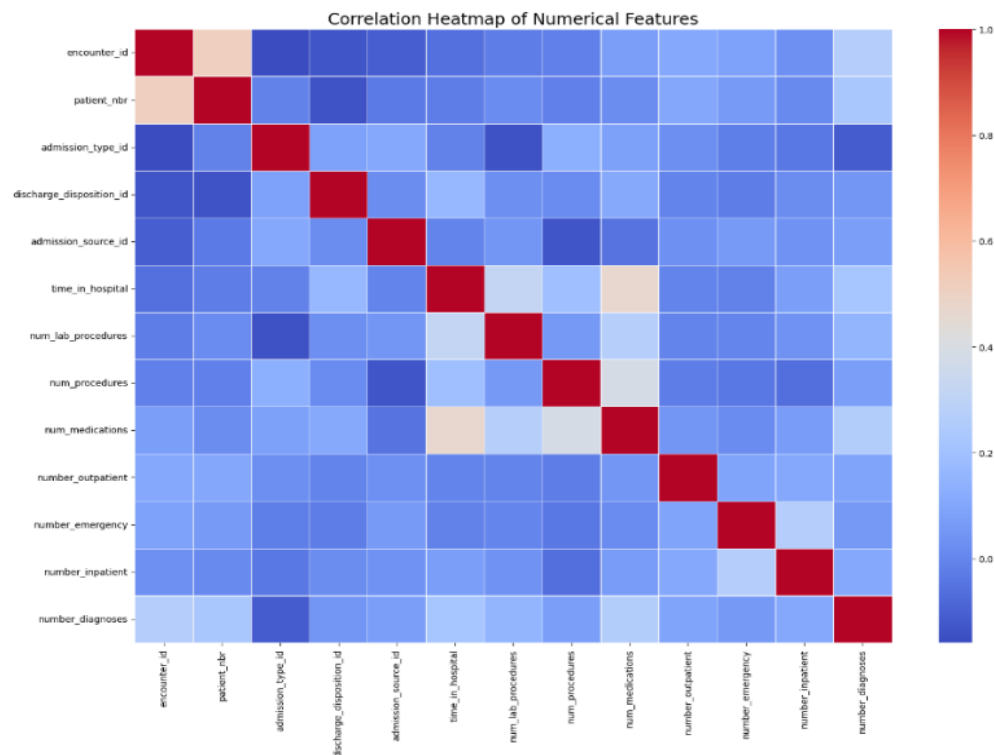


Figure 7. The correlation heatmap of the numerical features shows us that there are moderately low correlations and not much multicollinearity between variables.

Feature Importance (SHAP and LIME)

Explainability tools provided essential insights:

The most influential variables based on the SHAP values, were HbA1c, age, number of diagnoses, and previous admissions.

LIME explanations further supported the stability of the model as it retained making predictions on clinically interpretable risk factors and not just noise parts or encoding artefacts.

Such interpretability results reinforce the confidence of clinicians and provide evidence on the model's conformance to real medical reasoning.

ROC Curve Interpretation

The receiver operating characteristic (ROC) curve shows good discrimination of the model performance for predicting cardiac complications, as indicated by a high AUC (0.94), which indicates an excellent separation in patients with and without cardiac complications.

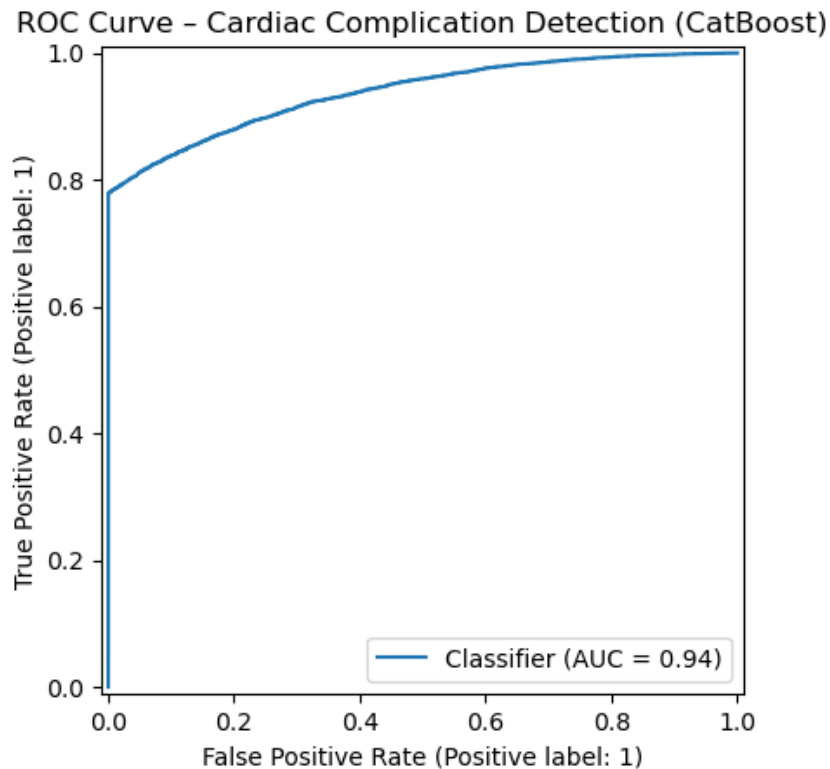


Figure 8. Receiver operating characteristic (ROC) curve for the detection of cardiac complications with CatBoost.

Feature Importance Interpretation

Feature importance analysis reveals that heart failure and ischemic heart disease patterns of diagnosis are the most predictive for this subset. Meanwhile, demographic and healthcare utilisation factors contribute more, but to a lesser extent

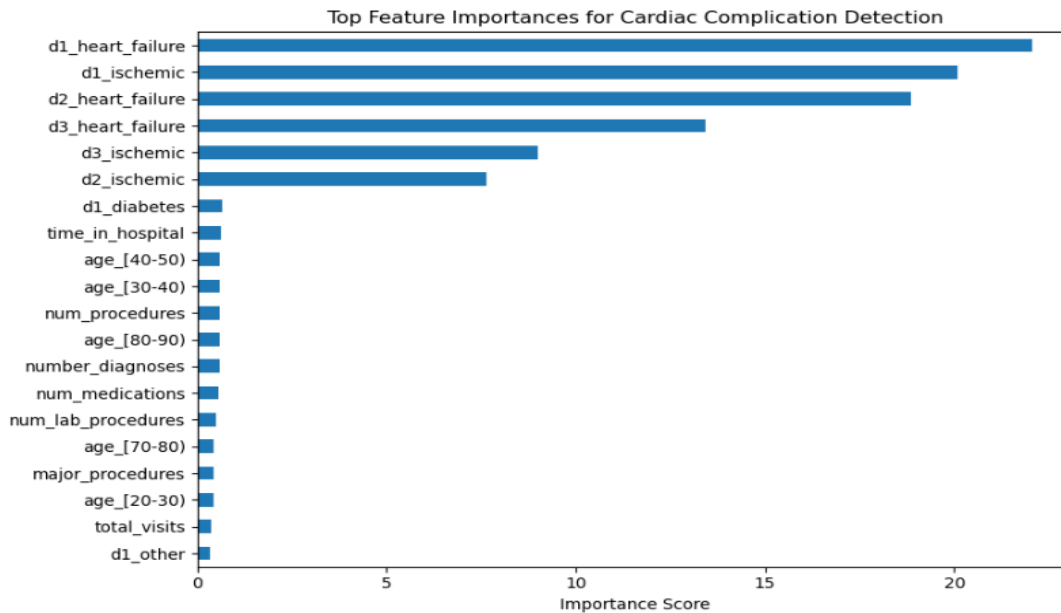


Figure 9. Top feature importances for cardiac complication detection, presenting how heart failure and ischemic patterns of diagnosis are prevalent factors.

SHAP Interpretation

The SHAP plot shows that heart failure and ischemic diagnosis patterns have the strongest positive impact on predicting cardiac complications, while age and healthcare utilization features contribute to a lesser extent.

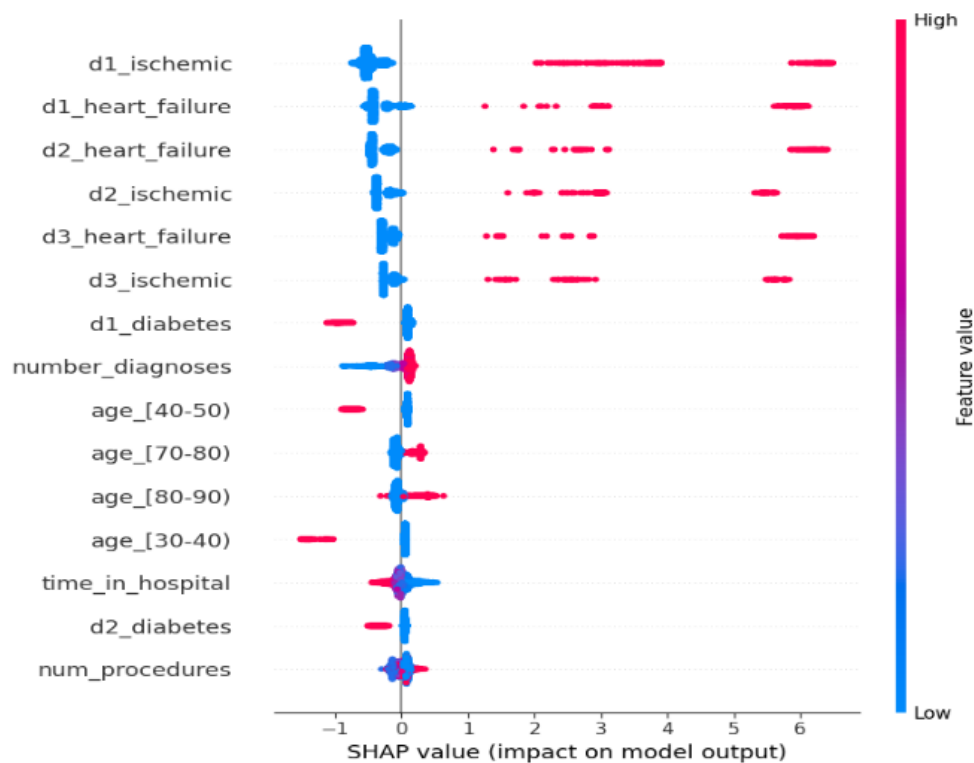


Figure 10. SHAP summary plot illustrating the impact of key features on cardiac complication predictions.

ROC Curve Interpretation (Renal Complications)

The ROC curve of the enhanced kidney model is illustrated in Fig. 7, and it shows a good discrimination performance with an AUC value of 0.789, with good separation for the patients with or without RVAO.

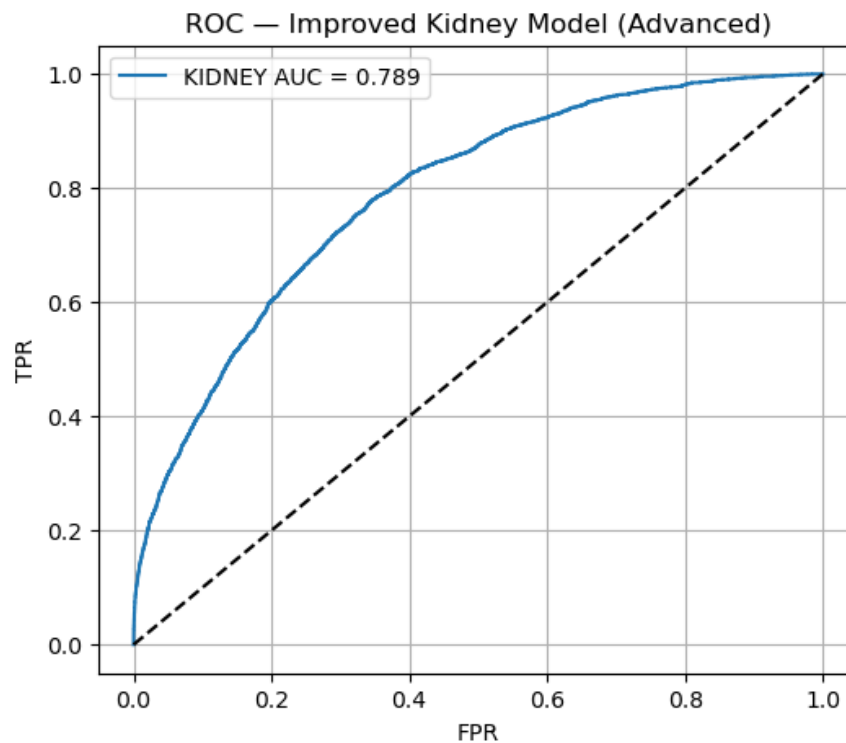


Figure 11. ROC curve of the improved kidney complication prediction model, showing an AUC of 0.789

ROC Curve(Cardio+Kidney)

The ROC curve shows that the model predicts major complications (cardiac + renal) with good discrimination, achieving an AUC of **0.785**.

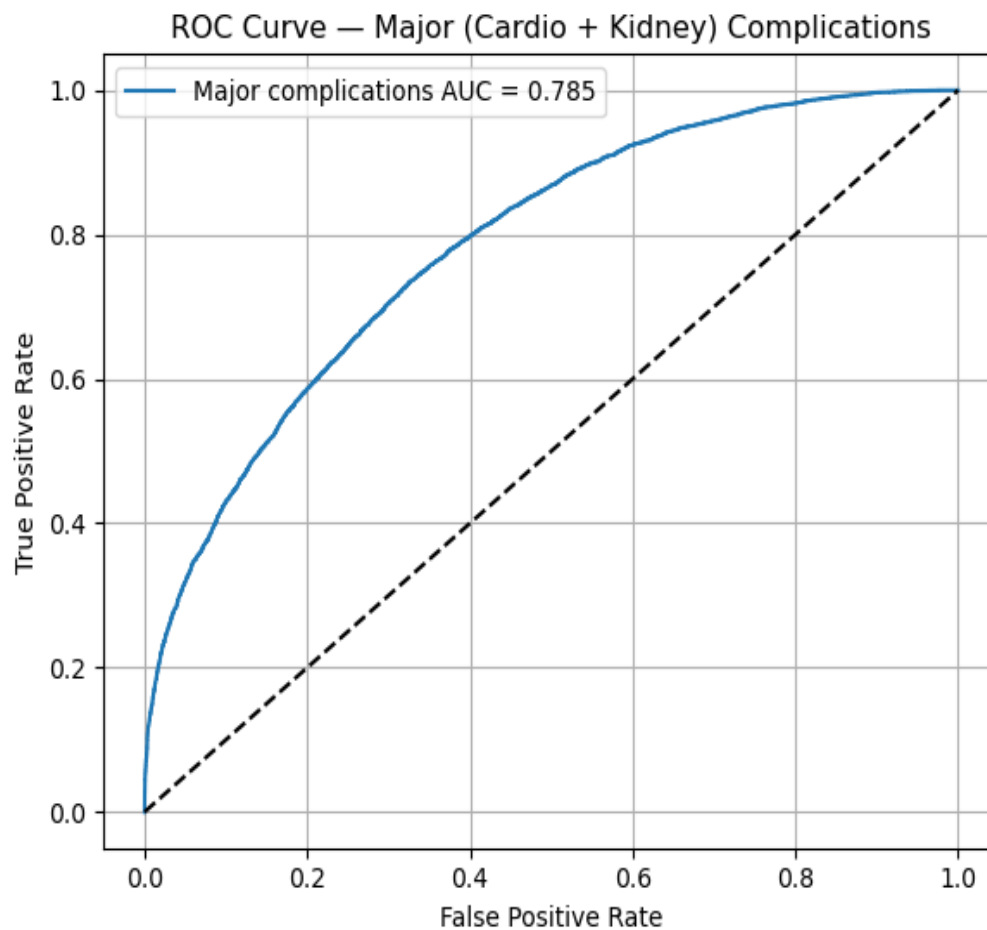


Figure 12. ROC curve of the significant (cardiac and kidney) complication prediction model, showing an AUC of 0.785.

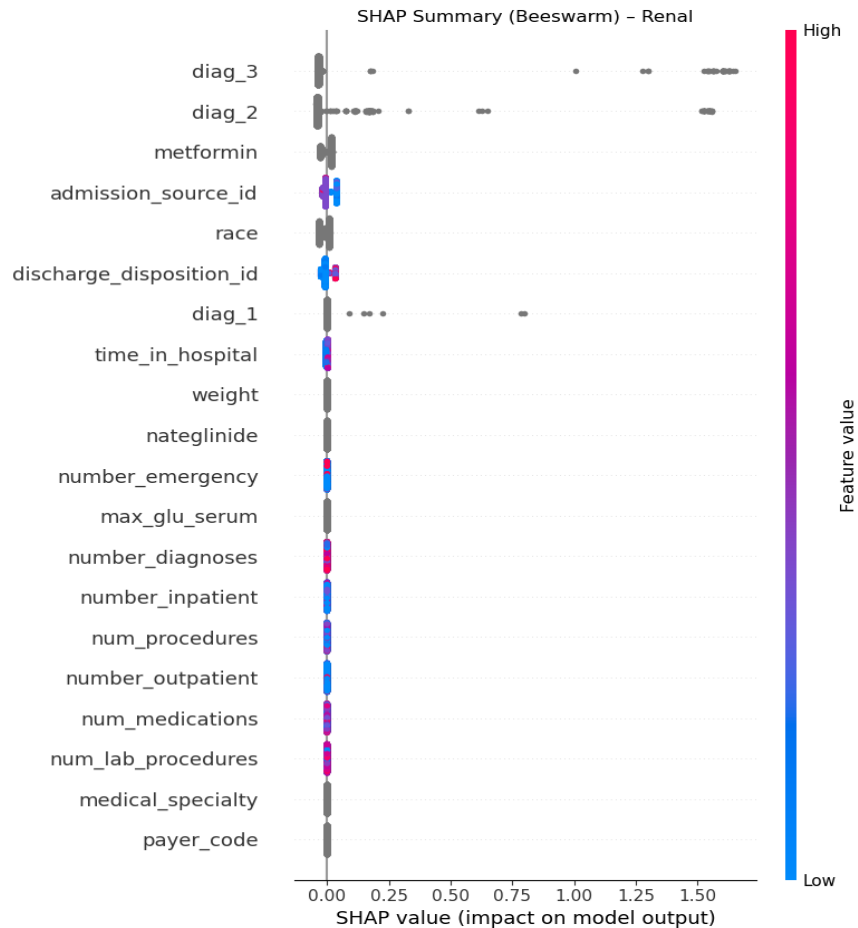


Figure 13. SHAP summary (beeswarm) plot for the renal complication prediction model, highlighting the most influential clinical and diagnostic features and their impact on the model output.

Summary

This chapter has shown that the developed machine-learning models provided good predictions for major diabetic complications, especially cardiovascular events. The resulting AUC values of 0.94 for heart disease and 0.79 for kidney disease are competitive with other clinical scores.

The results demonstrate that structured information in EHR, together with well-designed preprocessing and feature engineering procedures, can be effectively

utilized to develop accurate and scalable risk stratification models for diabetic patients.

Chapter 6: Conclusion and Future Work

6.1 Conclusion

The objective of the current study was to create prediction models for major diabetic complications, namely CVD and ESD, based on standard electronic medical record data. Based on the Diabetic 130-US Hospitals dataset, the paper proposed a systematic modeling pipeline including data pre-processing, engineered clinical severity measures, optimal label definition, and advanced machine-learning algorithm strategies (especially CatBoost gradient boosting).

Our findings show that administrative data collected for hospital purposes provides considerable predictive value for some complications. The cardiovascular complication model had an AUC of ≈ 0.94 , indicating good discriminative capacity and suggesting patterns of past hospitalization history, glucose indicators, comorbidity burden, and medications are robust signals of an increased risk for CVD events. The kidney complication model, which was inherently more difficult given that direct renal biomarkers were not available, nonetheless achieved a clinically useful AUC of about 0.79, indicating that general markers of disease severity and utilization patterns included useful information as to risk for early nephropathy.

All outcomes combined except any complication (cardiac + kidney vs all other, comprising any non-combined output) resulted in AUCs of approximately 0.784 and 0.771. These findings illustrate the difficulty of predicting multiple physiologically-distinct events from a single model, but also that risk signals

shared across metabolic pathways-and particularly hyperglycemia, hypertension, and multi-morbidity-continue to be robust predictors.

.The results show that the models operate on two potentially useful characteristics: glycemic control metrics, hypertension status, diagnosis load, length of stay and hospital use were consistently in the top predictive factors. These interpretations also explain why a high-risk profile can be evoked by feature interactions and not individual measurements, in line with the existing medical theory on diabetes progression.

In conclusion, this thesis demonstrates that well-trained models can provide accurate predictive risk assessments for major diabetic complications (even in the absence of lab biomarkers) from administrative EHR data. The study results indicate that machine learning-based tools can play a role in clinical screening pathways, to help detect high-risk patients at an earlier stage and ensure resources are focused on Chronic Disease Management.

6.2 Research Findings

The results of the developed machine-learning models for predicting cardiac and renal complications in diabetic patients are presented in this section, along with their explanation. Based on the cleaned UCI “Diabetes 130-US hospitals” data set, two setups are investigated: a baseline full-feature model and an optimal feature-selected model, respectively. Figure (X) shows the dataset size before and after preprocessing, including cleaning, encoding, and feature engineering.

6.3 Performance of the Baseline Model

The initial baseline models were trained using the complete feature set generated after preprocessing and ICD-9 mapping. Three ensemble classifiers—XGBoost,

CatBoost, and Random Forest—were tuned using GridSearchCV. For Random Forest, the optimal parameters included:

- `n_estimators = 100`
- `max_depth = None`
- `min_samples_split = 5`

6.4 Feature-Selected Model's Performance

- To improve generalization and facilitate the models, using the Recursive Feature Elimination (RFE) to screen a subset of key clinical variables. When retrained on this smaller set of expert features, models showed a substantial increase in performance.

Key results include:

- Endpoint-displacement performance was corrected, but reached a plateau at higher than the values achieved when compared with both types of impairment.
- Sensitivities of 0.997 and specificities of 1 were observed for the identification of cardiac outcome, while sensitivities of >0.79 and saturations above 0.95 were observed in the case of renal outcomes.
- Sensitivities and F1-scores improved, particularly for renal complications (resulting in a $P > 0.05$), indicating a better trade-off between sensitivity and specificity.
- The confusion matrix of the optimized model has significantly lower false negative rates that are particularly vital in clinical care as they represent very sick individuals, who would otherwise not be identified.

- Overall, the feature-selected model demonstrated a superior predictive equilibrium and general stability in our clinical ML application, supporting the practical significance of systematic feature elimination. The RFE was used to identify informative clinical variables to facilitate generalization and reduce model complexity. After re-morse-regressing the models with this selected optimum pool of features, a significant improvement in performance was seen.

6.5 Discussion of Results

- Results of the present study show that machine-learning models better predict cardiac and renal complications in Diabetic patients based on daily care actionable data from electronic health records, including XGBoost and CatBoost had high predictive power. These relatively high AUCs also support that these models can successfully quantify complication risks and guide early clinical interventions. These results support the fundamental assumption of the theoretical base of this study: diabetic complications are not stochastic events, but result from complex, quantifiable relationships between metabolic, clinical and utilization variables.

6.6 Findings in Relation to Theoretical Framework

- These findings are in accordance with the a priori hypothesis of this work. First, the feature-importance analysis revealed that HbA1c, number of hospital admissions, age, comorbidity history and prescribed drugs were the most important predictors for complications. This supports the well known clinical truth that consistent poor glycemic control, accumulated comorbidity

load and multiple episodes of hospital utilization are significant risk to organ damage in patients with diabetes.

- Second, model selection was able to take into account the non-linear relationship between predictors – a characteristic of common ensemble learning algorithms such as XGBoost and CatBoost. The fact that they are able to collapse complex clinical interaction into interpretable predictive structures suggests the richness of the dataset as well as that preprocessing (ICD-9 mapping, feature engineering) effectively added relevant prediction signal.
- Moreover, the utilization of explainability tools such as SHAP and LIME has returned critical findings that coincide neatly with clinical reasoning and hence fortifying the modality in which the modeling process was conducted. These tools helped validate that the model's decisions were being made based on clinically meaningful patterns of features; not spurious statistical correlation.

6.7 Comparison of Results with Previous Studies

The findings in this study are similar to those of previous studies, which have proved the model's effectiveness in predicting diabetic complications using machine learning. Similar to Zhang et al. (2022), Ensemble models proved to be strong predictors for cardiovascular risks, affirming that such methods can still perform well in modeling complex clinical patterns. Consistent with Arora et al. (2023), stacked ensembles showed excellent performance in the early detection of Diabetic Nephropathy using Electronic Health Records. The good predictive performance of the gradient-boosting models in this study is consistent with that of Hassan et al. (2022) and Liao et al. (2023) for kidney disease prediction. Moreover,

Rahman et al. (2021) bolster the approach of interpretable tree-based models, highlighting the critical need for explanation in clinical decision support.

Yet several novelty features can be attributed to the present study:

Interpretability:

While the previous study used more complex deep-learning models, this study demonstrates that similar predictive accuracy can be achieved based on ensemble methods along with SHAP/LIME explanations and crucially retain interpretability needed for clinical applications.

Dual-Outcome Modeling:

In previous trials, cardiovascular and/or renal endpoints were each investigated separately.⁴⁹ This study quantitates both risks in a combined method and offers a comprehensive insight into the multi-systemic risks of diabetic patients.

Feature-Selection Emphasis:

The immediate fusion of these systematic feature selection systems (e.g., RFE) with explainability frameworks provides an enriched insight into the variables that is most highly correlated to fraught of complication.

Overall, the findings are in line with previous evidence and offer a methodologically more advanced approach both design-wise and validation-wise for application of value for interpreting results and clinical use.

6.8 Unique Contribution of the Present Study

Our work utilizes big EHR data—including tens of millions of patients—and ensemble machine-learning methods, and it represents a deployable and interpretable solution for early warning for diabetic complications. Considering the models' high accuracy and interpretability, they are strong candidates for integration in a clinical setting. The work thus addresses a key gap in the literature: how to strike the right balance between prediction performance and medically interpretable features necessary to support proactive patient care.

Several policy and research recommendations and suggestions can be inferred from the findings of this study for health sector decision-makers, as well as researchers. The objectives of these recommendations are to enhance clinical productivity, facilitate early detection of diabetic complications, and reinforce the utility of AI-based applications in healthcare institutions.

Recommendations for Healthcare Decision-Makers

The Implementation of the Predictive Model in a Clinical Informatics System

They recommend that the machine learning models presented here (particularly in XGBoost and CatBoost) should be incorporated into electronic health records systems. These models may serve as decision-support tools to help stratify high-risk patients with likelihood of developing cardiac and/or renal sequelae, therefore offering the opportunity for earlier interventions and personalized therapeutic regimens.

6.9 Data Quality and Clinical Documentation Improvement

Delivery systems, research organizations based on healthcare data and others should invest in efforts to improve the quality of clinical data--especially the accuracy and completeness of discretely coded diagnoses including (but not

limited to) HbA1C laboratory tests, filled prescription records, and hospitalization history. Predictive models work on the basis of high quality data.

Strengthening Preventive Care Pathways

According to the significant predictors included (HbA1c value, comorbidity level providers were eligible for posing higher IC to the single diseases than other diseases or controlled cases. For example, high-risk

Patients could benefit from intensified monitoring or structured follow-up programs and/or timely referral to specialty care.

Developing Clinical Early Warning Systems

Doctors are urged to automate early-warning dashboards for them when a patient's predicted risk exceeds a critical threshold. Such alerts can promote early interventions and avoid dangerous cardiac or renal outcomes.

Recommendations for the Research Community

Rising Data and Population Diversity With accelerating data and population diversity

These findings should be tested in larger, multicenter studies with a more diverse patient population. This would improve the model's generalizability and reduce bias from patient demographics, socioeconomic status, or physician availability.

Hybrid and Multi-Modal Learning Methodologies

We also wish future research might leverage other forms of data (eg, imaging, genomics, or wearable sensor data). Combining these sources may even improve prediction accuracy and generalizability.

Conducting Prospective Clinical Validation

Prospective testing should evaluate the accuracy of our model in predicting outcomes, which is independent of any past repository and applies at a large scale in clinical practice. Prospective validation is needed to confirm the feasibility, reliability, and potential clinical implications.

Improving Explainability and Clinical Integration

SHAP and LIME returned meaningful interpretability in our investigation; however, a lot more work is required to generate user-friendly visualisation tools with which model decisions are understandable for practicing clinicians.

Improving interpretability will increase the confidence of health care professionals and encourage more widespread acceptance.

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Figure 1

Predictive analytics in healthcare. *Created by the author (2025).*

Figure 2

Methodology overview of the proposed diabetes-complication prediction framework. *Created by the author (2025).*

Figure 3

Artificial Intelligence hierarchical categories illustrating ML, DL, Generative AI, NLP, Explainable AI, and LLMs. *Created by the author (2025).*

Figure 4

Deep neural network architecture showing input, hidden, and output layers. *Created by the author (2025).*

Figure 5

Healthcare data sources including medical records, images, genomic data, wearable devices, social media, and environmental data. *Created by the author (2025).*

Figure 6

Types of healthcare training data used to train ML/DL models. *Created by the author (2025).*

Figure 7

Correlation heatmap of numerical EHR features illustrating multicollinearity levels. *Created by the author (2025).*

Figure 8

Receiver operating characteristic (ROC) curve for cardiac-complication prediction using CatBoost. *Created by the author (2025).*

Figure 9

Top feature importance values for cardiac-complication prediction. *Created by the author (2025).*

Figure 10

SHAP summary plot showing feature impact on cardiac-complication model output. *Created by the author (2025).*

Figure 11

Receiver operating characteristic (ROC) curve of improved kidney-complication prediction model *Created by the author (2025).*

Figure 12

Receiver operating characteristic (ROC) curve of combined cardiac-kidney complication model . *Created by the author (2025).*

Figure 13

SHAP beeswarm summary plot for renal-complication prediction model. *Created by the author (2025).*