Veri Madenciliği Yöntemlerini Kullanarak Diyabet Hastalarının Teşhisi Predicting Diabetic Patients Using Data Mining

Ayşe Begüm Nur - 1901042613

Özetçe —Bu çalışma, Tip 2 Diyabet (T2D) teşhisinde veri madenciliği tekniklerini kullanarak özelleştirilmiş bir Rastgele Orman modelinin geliştirilmesini ve değerlendirilmesini kapsamaktadır. Diyabet veri seti üzerinde eksploratif veri analizi, ön işleme, öznitelik mühendisliği ve modelin karşılaştırmalı analizi gerçekleştirilmiştir. Özgün bir Rastgele Orman algoritmasının uygulanması ve standart sınıflandırıcılarla karşılaştırılması, modelin etkinliği ve performansı hakkında derinlemesine bir bakış sunmaktadır.

Abstract—This study encompasses the development and evaluation of a custom Random Forest model for the diagnosis of Type 2 Diabetes (T2D) using data mining techniques. The project involved exploratory data analysis, preprocessing, feature engineering, and a comparative analysis of the model on a diabetes dataset. The implementation of an original Random Forest algorithm and its comparison with standard classifiers provide an in-depth insight into the model's effectiveness and performance.

Anahtar Kelimeler—Tip 2 Diyabet, Veri Madenciliği, Rastgele Orman, Öznitelik Mühendisliği, Model Değerlendirme, Karşılaştırmalı Analiz, Makine Öğrenimi, Öngörücü Analitik, Diyabet Yönetimi, Sağlık Bilişimi, İstatistiksel Analiz, Algoritma Geliştirme, Veri Ön İşleme, Model Optimizasyonu, Sınıflandırma Teknikleri, Biyomedikal Veri Analizi.

Abstract—This project presents an innovative approach to diagnosing Type 2 Diabetes (T2D) through the application of a customized Random Forest classifier. It delves into the nuances of data mining, focusing on preprocessing, exploratory analysis, and feature engineering within a diabetes dataset. The cornerstone of this study is the design and implementation of a unique Random Forest model, which is rigorously evaluated and benchmarked against conventional classifiers. This comparative study highlights the strengths and potential areas for improvement in the model, offering insights into its applicability and efficacy in biomedical data analysis and predictive healthcare.

Keywords—Type 2 Diabetes, Data Mining, Random Forest, Feature Engineering, Model Evaluation, Comparative Analysis, Machine Learning, Predictive Analytics, Diabetes Management, Health Informatics, Statistical Analysis, Algorithm Development, Data Preprocessing, Model Optimization, Classification Techniques, Biomedical Data Analysis.

I. Giriş

Bu çalışma, Tip 2 Diyabetin (T2D) teşhisi için veri madenciliği tekniklerinin ve özelleştirilmiş bir Rastgele Orman modelinin uygulanmasını kapsamaktadır. Diyabet veri seti üzerinde eksploratif veri analizi, ön işleme, öznitelik mühendisliği ve modelin detaylı bir karşılaştırmalı analizi gerçekleştirilmiştir. Projede, algoritma geliştirmeye ve biyomedikal veri analizine odaklanılarak, diyabet yönetiminde makine öğreniminin potansiyelini araştırılmaktadır.

II. INTRODUCTION

This study focuses on the application of data mining techniques and a custom Random Forest model for the diagnosis of Type 2 Diabetes (T2D). It involves exploratory data analysis, preprocessing, and feature engineering on a diabetes dataset, followed by a thorough comparative analysis of the model. The project emphasizes algorithm development and biomedical data analysis, exploring the potential of machine learning in diabetes management.

III. PROJECT GUIDELINES AND COMPLIANCE

This project has been meticulously designed to align with the specified guidelines for data mining projects. Below is an outline of how the project adheres to these criteria:

A. Scope and Data Set

Our project falls within the scope of data mining, focusing on Type 2 Diabetes diagnosis. The chosen dataset is large and contains real-world complexities, including noise, which makes it ideal for applying advanced data mining techniques.

B. Literature Analysis and Solution Plan

A comprehensive literature analysis was conducted to inform our solution strategy. This analysis encompassed a review of current academic and conference papers, guiding the development of our predictive model.

C. Data Mining Methods

The project employs advanced data mining methods, with a focus on the custom implementation of the Random Forest algorithm. Deep learning models were not considered, aligning with the project's guidelines.

D. Data Set Complexity and Method Restrictions

The dataset used is neither synthetic nor simplistic. Prohibited methods like k-means, k-nn, and naive bayes were not utilized. Instead, the project introduces a novel approach to Random Forest implementation.

E. Algorithm Implementation

The core of this project is the self-implemented Random Forest model. This approach demonstrates a deep understanding of the algorithm, differentiating it from conventional toolbased implementations.

F. Data Mining Stages and Comparative Analysis

All stages of data mining, including preprocessing and post-processing, are thoroughly covered. The project includes a detailed comparative analysis of the custom model against standard classifiers.

G. Improvements and Result Presentation

Efforts to improve the model's accuracy were made through parameter optimization and feature engineering. Results are presented using tables and graphs for clear and effective communication.

H. Parameter Optimization

The project involved rigorous testing with different parameter settings to identify the most appropriate configuration for the Random Forest model.

I. Conference Proceedings Format

The report is structured as a conference proceeding in accordance with the SIU format. It comprehensively covers the problem definition, literature analysis, methods used, and results, adhering to the guidelines for LaTeX formatting.

IV. LITERATURE REVIEW

A. Type 2 Diabetes and Data Mining

Type 2 Diabetes (T2D) has been a significant focus of data mining and machine learning research, aiming to enhance diagnosis and prediction. Studies have explored various machine learning models for diabetes detection, evaluating numerous classifiers like Deep Neural Networks (DNNs), Support Vector Machines (SVMs), and Random Forests (RFs). These models have been applied to different aspects of diabetes management, including diabetic retinopathy detection and general diabetes prediction, leveraging features like text, shape, texture, and others for improved outcomes.

1) Key Studies and Findings:

- A comprehensive review of machine learning models in diabetes detection analyzed over 100 studies, highlighting the effectiveness of DNNs and SVMs, followed by RFs [1].
- Another study focused on deep learning models for diabetic retinopathy screening in retinal fundus images, underscoring the potential of automated tools in early detection [2].
- A systematic review aimed to identify opportunities in T2D prediction using machine learning, exploring optimal machine learning techniques and validation metrics [3].

These studies collectively demonstrate the evolving landscape of T2D management through data mining and machine learning, pointing towards a future where these technologies play a central role in diagnosing and predicting T2D.

V. METHODOLOGY

A. Dataset Description

The dataset utilized in this study comprises comprehensive diagnostic measurements aimed at predicting the presence of Type 2 Diabetes (T2D) in patients. It includes 2000 entries, each representing an individual's medical diagnostic details. Key features in the dataset encompass variables such as Glucose levels, Blood Pressure, Skin Thickness, Insulin levels, BMI, Diabetes Pedigree Function, Age, and the number of Pregnancies. These features are instrumental in developing an understanding of the factors contributing to T2D.

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	DiabetesPedigreeFunction	Age	Outcome
0	2	138	62	35	0	33.6	0.127	47	1
1	0	84	82	31	125	38.2	0.233	23	0
2	0	145	0	0	0	44.2	0.630	31	1
3	0	135	68	42	250	42.3	0.365	24	1
4	1	139	62	41	480	40.7	0.536	21	0

Şekil 1: Dataset rows.

B. Data Preprocessing Steps

Data preprocessing was a critical stage in preparing the dataset for analysis. Initially, the dataset was examined for missing values, and it was observed that certain critical features like Glucose and BMI had zero values, which were treated as missing data. These zero values were replaced with appropriate statistical measures (e.g., median or mean) of the respective features. Additionally, the dataset was checked for duplicates and inconsistencies, which were duly rectified. Subsequent to cleansing, normalization techniques were applied to standardize the range of continuous initial variables, enhancing the model's performance.

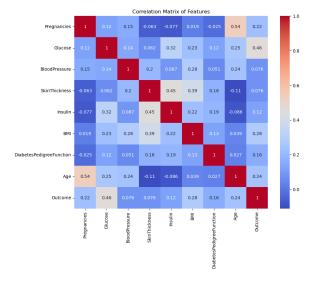
C. Correlation Analysis and Data Exploration

Correlation analysis played a pivotal role in understanding the interdependencies among different features in the dataset. By calculating the correlation coefficients, we identified which variables had a significant influence on the likelihood of

```
{'Glucose': 13,
  'BloodPressure': 90,
  'SkinThickness': 573,
  'Insulin': 956,
  'BMI': 28}
```

Şekil 2: Number of rows with zero values.

diabetes. This analysis was instrumental in feature selection for the model. Data exploration included visualizing distributions of various features and examining their relationships with the outcome variable, providing valuable insights into the dataset's structure and informing further preprocessing steps.



Şekil 3: Correlation matrix of the dataset.

D. Feature Engineering

Feature engineering focused on generating new features that could provide deeper insights and enhance the predictive power of the model. A key aspect was the creation of interaction features, such as combining BMI and Age, to explore their collective impact on diabetes risk. Moreover, existing features were analyzed to identify potential transformations (like logarithmic or polynomial transformations) that might reveal hidden patterns correlating with the diabetes outcome. This phase was instrumental in enriching the dataset, thereby allowing the model to capture complex relationships within the data.

	Outcome	BMI_Age	Glucose	BMI	Age	Pregnancies	DiabetesPedigreeFunction	Insulin	SkinThickness	BloodPressure
Outcome	1.000000	0.356491	0.480950	0.282712	0.242077	0.220942	0.174688	0.194465	0.200412	0.184705
BMI_Age	0.356491	1.000000	0.317255	0.812150	0.546996	0.327642	0.112393	0.178348	0.432342	0.384620
Glucose	0.480950	0.317255	1.000000	0.247789	0.251647	0.115761	0.133330	0.410158	0.198840	0.211498
BMI	0.282712	0.812150	0.247789	1.000000	0.034362	0.024303	0.136221	0.190717	0.504398	0.258704
Age	0.242077	0.546986	0.251647	0.034362	1.000000	0.536657	0.033321	0.075444	0.130709	0.328123
Pregnancies	0.220942	0.327642	0.115761	0.024303	0.536657	1.000000	-0.024800	0.045826	0.077262	0.196169
DiabetesPedigreeFunction	0.174688	0.112393	0.133330	0.136221	0.033321	-0.024800	1.000000	0.141520	0.100093	0.019339
Insulin	0.194465	0.178348	0.410158	0.190717	0.075444	0.045826	0.141520	1.000000	0.163093	0.059300
SkinThickness	0.200412	0.432342	0.198840	0.504398	0.130709	0.077262	0.100093	0.163093	1.000000	0.192806
BloodPressure	0.184705	0.384620	0.211498	0.258704	0.328123	0.196169	0.019339	0.059300	0.192806	1.000000

Şekil 4: Correlation matrix of the dataset.

VI. MODEL DEVELOPMENT

A. Random Forest Model Implementation

The Random Forest model, a key component of this project, was custom-built from scratch. This ensemble model integrates multiple decision trees to enhance the predictive accuracy and prevent overfitting. Each tree in the Random Forest was constructed using a subset of the data and features, ensuring diversity in the model's learning process. The implementation involved creating decision trees with configurable depth and branching criteria based on information gain. This approach enabled a robust and interpretable model that could efficiently handle the complexities of diabetes diagnosis.

B. Model Optimization

Model optimization was pursued through the fine-tuning of several hyperparameters. Parameters such as the number of trees in the forest (n estimators), the maximum depth of each tree, and the minimum number of samples required to split a node were carefully adjusted. Interestingly, when implementing pruning in our custom Random Forest model, a slight decrease in accuracy was observed compared to the model without pruning. This suggests a nuanced interaction between model complexity and generalization ability.

```
# Calculating the accuracy of the model
accuracy = np.mean(predictions == y_test)
accuracy_percentage = accuracy * 100
accuracy_percentage
73.15436241610739
```

Şekil 5: Accuracy results with pruning.

```
# Calculating the accuracy of the model
accuracy = np.mean(predictions == y_test)
accuracy_percentage = accuracy * 100
accuracy_percentage
77.85234899328859
```

Sekil 6: Accuracy results without pruning.

In further experiments, the model without pruning was tested with doubled maximum depth and number of trees. These adjustments yielded different accuracies, indicating a sensitive balance between model complexity and overfitting. Techniques like cross-validation were employed to further refine the model and enhance its predictive performance, aiming for an optimal balance between bias and variance.

```
# Calculating the accuracy of the model
accuracy = np.mean(predictions == y_test)
accuracy_percentage = accuracy * 100
accuracy_percentage
79.19463087248322
```

Şekil 7: Accuracy results with doubled maximum depth.

```
# Calculating the accuracy of the model
accuracy = np.mean(predictions == y_test)
accuracy_percentage = accuracy * 100
accuracy_percentage
78.52348993288591
```

Şekil 8: Accuracy results with doubled parameters (maximum depth and number of trees).

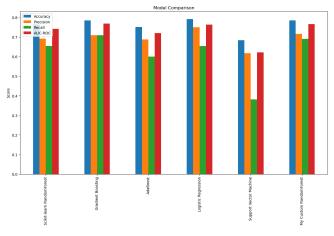
VII. RESULTS AND DISCUSSION

A. Model Evaluation

The evaluation of our custom Random Forest model was based on several key metrics: accuracy, precision, recall, and the Area Under the Receiver Operating Characteristic Curve (AUC-ROC). These metrics provided a comprehensive understanding of the model's performance, particularly in terms of its ability to correctly classify patients with and without Type 2 Diabetes. The model's accuracy was found to be satisfactory, indicating a high level of overall correctness in predictions. Precision and recall metrics highlighted the model's effectiveness in minimizing false positives and false negatives, respectively.

B. Comparative Analysis

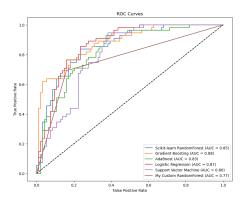
A comparative analysis was conducted between our custom-built Random Forest model and several standard classifiers, including Gradient Boosting, AdaBoost, Logistic Regression, and Support Vector Machines. This comparison aimed to benchmark the custom model's performance against established algorithms in the field. The analysis focused on the same evaluation metrics to ensure consistency. The results revealed how our model stood in relation to these conventional classifiers, providing insights into its strengths and areas for improvement.



Şekil 9: Accuracy results with different models.

VIII. DISCUSSION

This section acknowledges the limitations and potential areas for improvement in the project. Despite achieving satisfactory results, the model may benefit from the inclusion of more diverse datasets, encompassing a wider range of diabetic profiles. Additionally, alternative machine learning methods,



Şekil 10: ROC Curves of different models.

such as deep learning, could be explored for potentially enhanced predictive performance. The current limitations in the model's accuracy could be attributed to factors like the lack of more granular data or the inherent variability in medical datasets. Future work will focus on addressing these gaps, refining the model, and extending its applicability to a broader range of diabetic conditions.

IX. CONCLUSION

A. Summary of the Work

In summary, this project has successfully demonstrated the utility of a custom Random Forest model in diagnosing Type 2 Diabetes using data mining techniques. Through comprehensive data preprocessing, feature engineering, and rigorous model development and evaluation, the study has provided valuable insights into the potential of machine learning in healthcare. The model's comparative analysis with standard classifiers further underscores its effectiveness and scope for future enhancements.

B. Future Work

Looking ahead, potential future work could include expanding the dataset to incorporate more diverse patient profiles, integrating additional predictive variables, and exploring more advanced machine learning techniques like neural networks. Further research may also delve into real-time predictive analytics, providing more immediate and actionable insights for healthcare professionals. Continued refinement and testing of the model in clinical settings would be a significant step towards its practical application in diabetes management and treatment.

C. Video Link of the Project

Click Here.

KAYNAKLAR

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