

Disease prediction system using machine learning

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Abstract- In contemporary times, the prevalence of diverse diseases has surged due to environmental conditions and lifestyle choices. Consequently, the early prediction of diseases has emerged as a crucial task to enhance healthcare outcomes. However, accurately predicting diseases solely based on symptoms poses significant challenges for medical practitioners. Overcoming this obstacle, data mining has surfaced as a pivotal tool to predict diseases more effectively. The field of medical science witnesses a substantial growth in data volume each year, presenting an opportunity for accurate analysis of medical data, thereby benefiting early patient care. By harnessing disease data, data mining uncovers hidden patterns within vast medical datasets. In this study, we propose a novel approach for general disease prediction, utilizing patient symptoms as predictive indicators. To achieve this, we employ the K-Nearest Neighbor (KNN) and Support Vector Machine (SVM) machine learning algorithms, known for their accuracy in disease prediction. The success of our predictive model hinges on the availability of a comprehensive dataset containing disease symptoms.

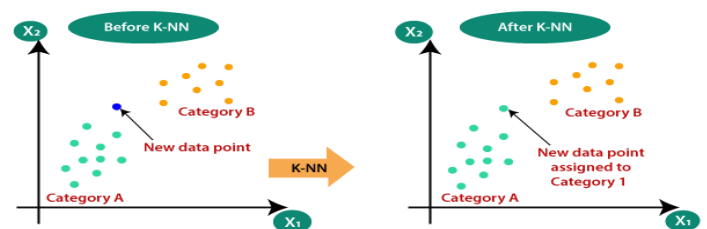
I. INTRODUCTION

Machine learning has played a pivotal role in enhancing computer intelligence, enabling computers to think and learn autonomously. Various machine learning techniques, including Unsupervised, Semi-Supervised, Supervised, Reinforcement, Evolutionary Learning, and Deep Learning, have proven valuable for classifying vast datasets swiftly. The field of general disease prediction has seen significant advancements, where living habits and checkup information are now considered for more accurate results. In this context, the CNN (Convolutional Neural Network) algorithm has demonstrated a remarkable accuracy of 83.5%, surpassing the KNN algorithm. Additionally, CNN proves to be more efficient in terms of time and memory requirements. Beyond predicting diseases, this system further provides risk assessment, categorizing individuals into lower or higher risk for general diseases. Machine learning classifiers have become indispensable tools for medical data analysis and disease prediction, partitioning datasets into multiple classes based on their distinct characteristics.

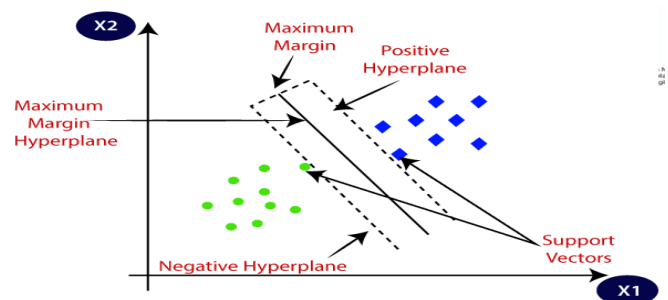
II. METHODOLOGY

1. **K Nearest Neighbour:** The K-Nearest Neighbor (KNN) algorithm is a simple Supervised Learning technique in Machine Learning. It operates on the principle of similarity, where a new data point is classified into a category based on its resemblance to the available data. KNN stores all the existing data and, when presented with new data, assigns it to a suitable category based on similarity. It can be used for both Regression and Classification tasks, but it is primarily

employed for Classification problems. As a non-parametric algorithm, KNN makes no assumptions about the underlying data distribution. Additionally, it is known as a lazy learner algorithm since it postpones learning from the training set and performs classification when new data is encountered.



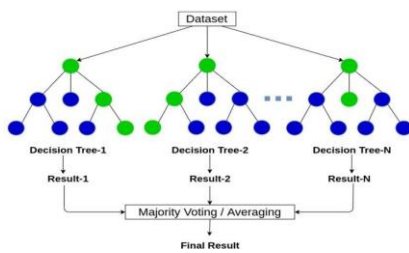
2. **Support Vector Machine:** Support Vector Machine (SVM), a widely used Supervised Learning algorithm for both Classification and Regression tasks. However, its primary application is in Classification problems within Machine Learning. The main objective of SVM is to establish an optimal decision boundary, known as a hyperplane, that effectively separates the n-dimensional space into distinct classes. By identifying the extreme points, also called support vectors, SVM constructs the hyperplane. The paragraph concludes by mentioning a diagram illustrating how SVM can classify two different categories using a decision boundary or hyperplane.



3. **Random Forest:** Random Forest is a versatile machine learning algorithm utilized for both classification and regression tasks. It falls under the category of ensemble learning methods, which combine multiple decision trees to create a more accurate and robust model. The algorithm constructs numerous decision trees, each trained on a

random subset of the data and features at each node, generating a diverse set of trees. The combination of their predictions through ensemble learning results in a more accurate and stable prediction. In classification, the most common prediction among the trees is chosen, while in regression, the average prediction is taken. Random Forest possesses several advantages, including its ability to handle large datasets with high dimensionality, manage missing values, and tolerate noisy data.

Random Forest



III. HARDWARE & SOFTWARE REQUIREMENT

1. Software:

- Android Studio is the official integrated development environment (IDE) for Android app development. It is built on JetBrains' IntelliJ IDEA and supports Android development across various devices.
- VS Code is a source-code editor developed by Microsoft, offering features like debugging, code completion, snippets, and Git integration. It is a streamlined editor that provides essential tools for quick code development.
- Sublime Text is a cross-platform source code editor with support for numerous programming languages. Its functionality can be extended through community-built plugins.
- Octave is a high-level programming language focused on numerical computing. It offers a versatile interpreter, graphical user interface, and supports matrix-based syntax, making it compatible with MATLAB.

Hardware:

- CPU: Intel Core 2 Quad CPU Q6600 @ 2.40GHz (4 CPUs) / AMD Phenom 9850 Quad-Core Processor (4 CPUs) @ 2.5GHz
- RAM: 4 GB
- OS: Windows 10 64 Bit, Windows 8.1 64 Bit, Windows 8 64 Bit, Windows 7 64 Bit

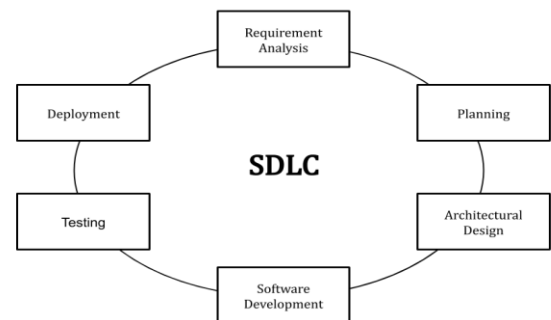
Service Pack 1, Windows Vista 64 Bit
Service Pack 2*

- VIDEO CARD: 32 MB Direct3D Video Card
- FREE DISK SPACE: 10GB
- DEDICATED VIDEO RAM: 1 GB

IV. SOFTWARE DEVELOPMENT LIFE CYCLE

The Software Development Life Cycle (SDLC) is a systematic process aimed at producing software of high quality, minimizing costs, and reducing development time. It offers a well-structured flow of phases that enable organizations to efficiently create thoroughly tested and production-ready software. SDLC ensures a methodical approach to software development, resulting in reliable and effective software products.

The SDLC involves six phases:



The various software development life cycle (SDLC) models are including the Waterfall model, Rapid Application Development (RAD) process, Spiral model, V-Model, Incremental model, Agile methodology, and Big Bang model. The Waterfall model divides the software development process into sequential phases. RAD is a time-sensitive adoption of the Waterfall model, focusing on quicker development through focus groups to gather requirements. The Spiral model combines rapid prototyping and concurrency, addressing objectives and risks in cycles. V-Model emphasizes parallel planning and testing phases. The Incremental model breaks requirements into groups and iteratively adds functionality. Agile methodology promotes continuous interaction between development and testing in incremental builds.

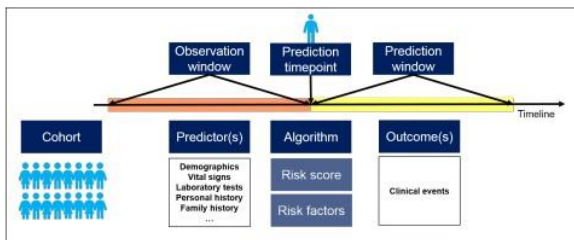
The SDLC model which we used is Prototype Model.

Prototype Model, a software development approach that starts with requirements gathering and involves creating a quick design followed by the development of a prototype. The loop continues with feedback from users leading to better versions of the prototype until the customer is satisfied. Once approved, the prototype is converted into the final system with considerations for quality and security. The

second part of the paragraph discusses the effectiveness of the Prototype Model in the context of disease prediction. It mentions using a disease dataset from the UCI machine learning website and performing preprocessing to clean the data. Feature extraction and selection are carried out, followed by classification using techniques like KNN and SVM for accurate disease prediction.

V. RISK ASSESSMENT

A risk prediction model is a set of equations that estimates the likelihood of developing a specific clinical outcome based on personal profiles. The key elements of such a model include the cohort, which represents the group of individuals likely to develop the outcome. Predictors are personal characteristics used in risk prediction, such as demographics, vital signs, laboratory results, and family disease history. For instance, predictors for type 2 diabetes may include age, family history of diabetes, body mass index, and waist circumference. Similarly, predictors for cardiovascular disease may include factors like smoking, diabetes, hyperlipidemia, and hypertension, among others. While including more predictors can improve prediction performance, it may also increase data collection burden. Hence, using easily accessible and less invasive predictors may encourage wider adoption of the model.

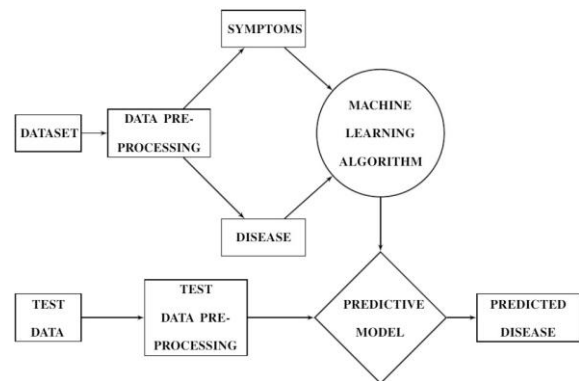


Evaluation metrics, feature engineering, pre-processing, feature construction, feature selection, and the risk equation are crucial components in developing effective risk prediction models.

Classification is a common type of risk prediction problem, particularly in binary classification, where the model predicts between two outcomes. Model performance can be assessed using various metrics, including discrimination (the model's ability to differentiate between positive and negative cases), calibration (how close the predicted risk aligns with the actual probability), and interpretability (how easily the model can be understood by humans). Classification models are evaluated based on discrimination, calibration, and interpretability, each providing valuable insights into the model's performance and understandability.

VI. SYSTEM DESIGN / WIREFRAME

1. Machine Learning Model



2. User Interface

The project involves the development of a UI using React and React Native, integrated with an ML model created through data mining techniques and ML algorithms. Several essential modules are included in the project:

1. Creation of a Linux machine Elastic Compute Cloud (EC2) instance on AWS.
2. Establishment of a postgres RDS instance for the database on AWS.
3. Downloading the project on the EC2 instance and setting up a suitable Python3 environment with Django installed.
4. Installation of pgadmin and accessing the remote RDS database on the local machine.
5. Creation of migrations and launching the project.
6. Accessing the project on a web browser using the public IP of the EC2 instance.
7. Implementation of user account creation and disease prediction functionality.

Each module plays a vital role in ensuring the proper functioning and seamless execution of the project. Missing any of these modules could lead to issues, as they collectively enable access to the UI and utilize the ML model for disease prediction.

VII. TESTING, TRAINING AND EVALUATION

Unit testing : Unit tests focus on verifying that correct changes occur in the program's state when a transaction is processed. Business logic in transaction processor functions is thoroughly tested to achieve high code coverage and detect typos or logic errors. Each algorithm is individually tested, and pre-processing is adjusted accordingly to enhance accuracy.

System testing: System Testing involves evaluating the complete and integrated software to assess its compliance with specified requirements. This level of testing encompasses various types, including Usability Testing to check user-friendliness, Load Testing to assess performance under real-life loads, Regression Testing to ensure no new bugs were introduced during development, Recovery Testing



to demonstrate reliability, and Migration Testing to assess smooth transitions between system infrastructures.

Functional test: Functional Testing, also known as functional completeness testing, involves identifying any possible missing functions in the program. It evaluates use-case scenarios and related business processes, ensuring the correct behavior of components like chatbots and smart contracts.

VIII. FUTURE SCOPE OF PROJECT

The future of disease prediction systems using machine learning is promising, driven by the increasing availability of healthcare data, improved algorithms, and better computing power. These systems hold immense potential for healthcare professionals and patients alike.

The primary advantage of disease prediction systems is their capability to detect diseases at an early stage. Early identification is crucial for successful treatment, and machine learning algorithms excel at identifying hidden patterns and risk factors that may elude human clinicians. With the integration of wearable technology and IoT devices, continuous monitoring becomes possible, facilitating early detection and timely interventions. Machine learning algorithms also play a key role in identifying risk factors and predicting disease outcomes for individual patients. By analyzing vast amounts of data from various sources, these systems can offer personalized recommendations for prevention and treatment, leading to improved patient outcomes and reduced healthcare costs.

As the technology continues to evolve, disease prediction systems using machine learning may integrate with emerging technologies like blockchain and decentralized databases. These advancements can enhance the security and privacy of patient data while facilitating seamless data sharing between healthcare providers and researchers.

IX .CONCLUSION

Disease prediction systems using machine learning hold immense potential to revolutionize healthcare by analyzing extensive data, identifying risk factors, and providing personalized recommendations. They enable early disease detection, leading to better treatment outcomes, and aid in developing effective public health interventions. However, careful implementation is essential, addressing ethical, privacy, and reliability concerns through rigorous testing. The future is promising, with advancements expected in accuracy and personalized treatments. Integrating technologies like blockchain can enhance data security and sharing. Overall, disease prediction systems using machine learning can transform healthcare and contribute to a healthier and equitable future.

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