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Software for Disease Prediction

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Abstract: Diseases are one of the most serious challenges in both developing and developed countries. According to the International Diabetes Federation, there are 285 million ill people worldwide. This total is expected to rise to 380 million within 20 years.

Due to its importance, a design of classifier for the detection of symptoms of disease with optimal cost and better performance is the need of the age.

The Pima Indian database at the UCI machine learning laboratory has become a standard for testing data mining algorithms to see their prediction accuracy in data classification. The proposed method uses Support Vector Machine (SVM), a machine learning method as the classifier for diagnosis of disease. The machine learning method focus on classifying disease from high dimensional medical data set. The experimental results obtained show that support vector machine can be successfully used for diagnosing disease.

I. INTRODUCTION

A. Disease Prediction Using Machine learning Software Project Report

Diabetes is one of the common and rapidly increasing diseases in the world. It is a major health problem in most of the countries. Diabetes is a condition in which your body is unable to produce the required amount of insulin needed to regulate the amount of sugar in the body.

This leads to various diseases including heart disease, kidney disease, blindness, nerve damage and blood vessels damage. There are two general reasons for diabetes: The pancreas does not make enough insulin or the body does not produce enough insulin. Only 5-10 % of people with this have this form of the disease (Type-1). Cells do not respond to the insulin that is produced (Type-2). Insulin is the principle hormone that regulates uptake of glucose from the blood into most cells (muscle and fat cells). If the amount of insulin available is insufficient, then glucose will not have its usual effect so that glucose will not be absorbed by the body cells that require it.

Disease being one of the major contributors to the mortality rate. Detection and diagnosis of diabetes at an early stage is the need of the day. Disease diagnosis and interpretation of the data is an important classification problem. A classifier is required and to be designed that is cost efficient, convenient and accurate. Artificial intelligence and Soft Computing Techniques provide a great deal of human ideologies and are involved in human related fields of application. These systems find a place in the medical diagnosis. A medical diagnosis is a classification process. A physician has to analyze lot of factors before diagnosing the diabetes which makes physician''s job difficult.

In recent times, machine learning and data mining techniques have been considered to design automatic diagnosis system for diseases. Recently, there are many methods and algorithms used to mine biomedical datasets for hidden information including Neural networks (NNs).

These algorithms decrease the time spent for processing symptoms and producing diagnoses, making them more precise at the same time. There is a great variety of methods related to diagnosis and classification of diabetes disease in the literature. Polat etal used principal component analysis and neuro fuzzy inference for diabetes data classification. Deng and kasabov obtained 78.4% classification accuracy with 10-fold cross-validation (FC) using ESOM. Yu et al. combined Quantum Particle Swarm Optimization (QPSO) and Weighted Least Square (WLS) Support Vector Machine to diagnose Type-2 diabetes. Smith et al. proposed a neural network ADAP algorithm to build associative models. 576 randomly selected data are used for training and 192 test cases showed an accuracy of 76%. Quinlan applied c4.5 algorithm and the classification accuracy was 77.1%. Sahan et al used Attribute Weighted Artificial Immune System with 10- fold cross validation method. The Support Vector Machine (SVM) is a novel learning machine introduced first by Vapnik and has been applied in several financial applications recently, mainly in the area of time series prediction and classification.



II. WORKING MODULES

Fig. 1: Welcome page

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| Heart disease is one of the biggest cause for morbidity and mortality among the population of the world. Prediction of cardiovascular disease is regarded as one of the most important subject in the section of | Diabetes is considered as one of the deadliest and chronic diseases which causes an increase in blood sugar. Many complications occur if diabetes remains untreated and unidentified. The tedious | Breast cancer is one of the most common cancers among women worldwide, representing the majority of new cancer cases and cancer-related deaths according to global statistics, making it a significant |
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Fig. 2: Select option to check health.



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III. DATA-SET

The <u>Pima</u> are a group of Native Americans living in Arizona. A genetic predisposition allowed this group to survive normally to a diet poor of carbohydrates for years. In the recent years, because of a sudden shift from traditional agricultural crops to processed foods, together with a decline in physical activity, made them develop the highest prevalence of disease and for this reason they have been subject of many studies.

A. Dataset

The dataset includes data from 768 women with 8 characteristics, in particular:

- *1)* Number of times pregnant (opction in this project)
- 2) Plasma glucose concentration a 2 hours in an oral glucose tolerance test
- *3)* Blood pressure (mm Hg)
- 4) Triceps skin fold thickness (mm)
- 5) 2-Hour serum insulin (mu U/ml)
- 6) Body mass index (weight in kg/(height in m)^2)
- 7) Diabetes pedigree function
- 8) Age (years)

The last column of the dataset indicates if the person has been diagnosed with diabetes (1) or not (0) Source: The original dataset is available at UCI Machine Learning Repository and can be downloaded from this address: <u>http://archive.ics.uci.edu/ml/datasets/Pima+Indians+Diabetes</u>

B. The Problem

The type of dataset and problem is a classic supervised binary classification. Given a number of elements all with certain characteristics (features), we want to build a machine learning model to identify people affected by type 2 diabetes.

To solve the problem we will have to analyse the data, do any required transformation and normalisation, apply a machine learning algorithm, train a model, check the performance of the trained model and iterate with other algorithms until we find the most performant for our type of dataset.

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Fig 3: Symptopms for diabetes



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Fig 4: Symptoms for breast cancer

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Fig 4: Symptoms for Heart Disease



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IV. CONCLUSION

The development of software or website includes so many people like user system developer, user of system and the management, it is important to identify the system requirements by properly collecting required data to interact with supplier and customer of the system. Proper design builds upon this foundation to give a blue print, which is actually implemented by the developers. On realizing the importance of systematic documentation all the processes are implemented using a software engineering approach. Working in a live environment enables one to appreciate the intricacies involved in the System Development Life Cycle (SDLC). We have gained a lot of practical knowledge from this project, which we think, shall make us stand in a good state in the future.

V. ACKNOWLEDGEMENT

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