



Non- invasive technique using breath analysis for detection and classification of diabetes

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Abstract

Diabetes, a metabolic disease that is characterized by high glucose level in the blood, is a major problem affecting millions of people today. This disease if left unchecked can create enormous implication on the health of the population. Among the various non-invasive methods of detection, breath analysis presents an easier, more accurate and viable method in providing comprehensive clinical care for the disease. This paper examines the concentration of acetone levels in breath for monitoring blood-glucose levels and thus predicting diabetes. The analysis uses the support vector mechanism to classify the response to healthy and diabetic samples. For the analysis, ten subject samples of acetone levels are taken into consideration and are classified according to three labels, which are healthy, type one diabetic and type two diabetic.

Keywords: Acetone Concentration; Breath; Blood Glucose Level; Diabetes; Supports Vector Classifier.

1. Introduction

Diabetes can be described as a group of metabolic diseases where the blood-glucose level in the body is higher than the normal prescribed parameter. When a person suffers from diabetes, it is seen that their body is either unable to secrete enough insulin or their body is not able to use the insulin produced by the liver. This causes sugar to build-up in the blood thus leading to diabetes. There are two major types of diabetes, which include type 1 and type 2. Type 1 diabetes is the result of the body's failure to produce enough insulin. While type 2 is a condition in which cells fail to respond to the insulin produced in the body properly. It is seen that the prescribed parameter of blood-glucose levels (BGL) in healthy subjects before meals is around 70 to 80 mg/dL. Sugar less than 100 mg/dL while fasting is considered normal by today's standards. Any BGL higher than normal is considered unhealthy.

The method of obtaining glucose levels in the body is by drawing blood samples using invasive techniques. A lancet device is used to draw the blood by pricking the finger or the forearm [1]. The droplet of blood obtained is then placed on a disposable strip which consists of a sensing element. A glucometer is then used to calculate the blood-glucose levels from these strips. Though this process is highly accurate, it is also painful and inconvenient, especially when multiple readings are required to be taken in a day. Thus there is an essential need for a non-invasive technique for monitoring BGL.

Investigations show that urine, sweat, saliva; tears and breath contain traces of glucose in them, and these traces vary with the levels of glucose in the blood. Therefore, these human serums have recently gained recognition as feasible alternatives to using blood for glucose measurement. Extensive research conducted in this area concludes that human breath is a good alternative to monitor and diagnose glucose levels as acetone in the breath has shown a good correlation to BGL. Using breath as a deduction technique allows deduction of blood-glucose levels by just exhaling into the monitoring device. It is seen that in the human breath, there are numbers of chemical compounds that relate to different diseases [2], [3]. Traces of acetone in the breath are used for the detection of diabetes.

According to the studies, it is seen that patients who have diabetes have body cells that are unable to absorb the glucose in blood. In such cases, when the liver breaks down fat for energy, there occurs an abnormal increase in ketone bodies in the patient's blood. Acetone is one of the three kinds of ketone bodies, which are volatile, and the body exhales the acetone thus formed. Therefore, higher concentration of acetone is found in the exhaled air of a diabetic patient [4].

Initially complicated techniques such as gas chromatography mass spectroscopy, selected ion flow tube mass spectroscopy and cavity ring down spectroscopy were used to determine the concentrations of acetone in breath. Gas chromatography- mass spectroscopy [5] works on the principle that the difference in the chemical properties between the different molecules in the gaseous mixture and their relative affinity for the stationary phase of the column promotes the separation of the molecules. In the Selected ion flow tube-mass spectroscopy method it was seen that when the neutral analyte molecules of a sample vapour met the precursor ions, they may undergo chemical ionization, which depends on their chemical properties, such as their proton affinity or ionization energy [6]. Cavity Ringdown Spectroscopy [7] is an optical spectroscopic technique that measures the absolute extinction by samples that absorb or scatter light. These techniques remain unsuitable for clinical application due to their low portability, complex mechanisms and high cost. The electronic nose or e-nose model [8], [9] overcomes these drawbacks significantly and is emerging as a good alternative suited to clinical applications. Research conducted with e-nose provides substantial data on acetone concentrations in breath [10].

This paper classifies the concentration and uses support vector mechanism (SVM) classifier to analyze the data signals. Different acetone concentration levels are classified as healthy breath, type one diabetic or type two diabetic which is then displayed as the output. Initially, the acetone levels were classified using SVM classifier but the accuracy levels remained inadequate. To overcome this issue and increase the level of accuracy SVM classification with margin sampling has been adopted.

2. Breath analysis procedure a step before the final submission

2.1. Proposed analysis

The proposed analysis system here distinguishes the concentration of acetone levels in the breath as healthy or diabetic. The basic model of the proposed system is described by the following flow chart given in Figure 1.

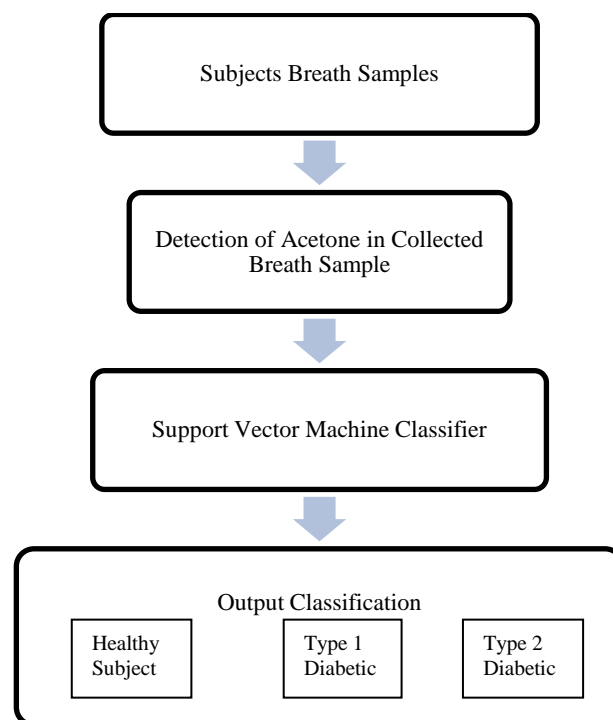


Fig. 1: The Flow Chart of the Proposed System.

The operation of the system happens in four stages. Initially, the subject's breath samples are used for the acetone concentration prediction. The acetone concentration data from various samples are detected using metal oxide semiconductor sensor and are taken individually as input in the next stage. The SVM classification model then classifies the acetone concentration levels based on the thresholds described in the next section. In the final stage, the output is

displayed under any one of the three classifiers considered for this system. Thus the diabetic and the healthy breath samples are separated in this analysis model.

2.2. Acetone level concentrations

Human breath consists of many biomarkers which are used for the detection of many diseases. It is explained previously how acetone in the breath is used as the biomarker for the prediction of diabetes. Wang et al. [11] carried out extensive studies and found a linear correlation between the mean concentration of breath acetone and the mean blood-glucose levels of each group. It was also observed that the breath acetone declined linearly with blood-glucose levels [12]. According to [12], [13], it is seen that the concentration of acetone levels in the breath varies for a healthy and a diabetic subject. It is seen that in all healthy samples, acetone levels are less than 0.76ppmv and in all samples with diabetes acetone levels showed levels higher than 1.71ppmv. The Table I shows the variation of the concentration of acetone levels.

Table 1: Variation of Acetone Levels

Classification	Samples	Acetone Concentration [ppm]
1.	Healthy Subject	0.22 to 0.80 ppm
2.	Type 2 Diabetic Subject	1.76 to 3.73 ppm
3.	Type 1 Diabetic Subject	As high as 21 ppm

PPM- parts per million.

For this analysis, ten different concentrations of acetone levels are considered and are used for the classification of the data. The acetone concentrations in the breath of 10 subjects are taken in this paper. These data samples are further given to the classifier. The acetone concentration that is detected from the breath for this analysis is in the range of parts per million.

2.3. Classification

In this paper, a support vector machine (SVM) classifier is used to classify the data samples into three classes. SVM [14] is a supervised learning model that is used to analyze a given data set and help recognize patterns that is used for the classification of the samples. These models construct hyper planes that are used for the classification and the regression of the data [15]. Here the SVM divides the acetone concentration data which is the input into three different classes.

The Figure 2 shows the graphical representation of the acetone concentrations of the subjects considered for this analysis. The bar graph that is plotted clearly shows the variation of the acetone levels of a healthy breath and a diabetic patient. Hence the above thresholds can easily segregate the samples as healthy, type 1 or type two diabetes.

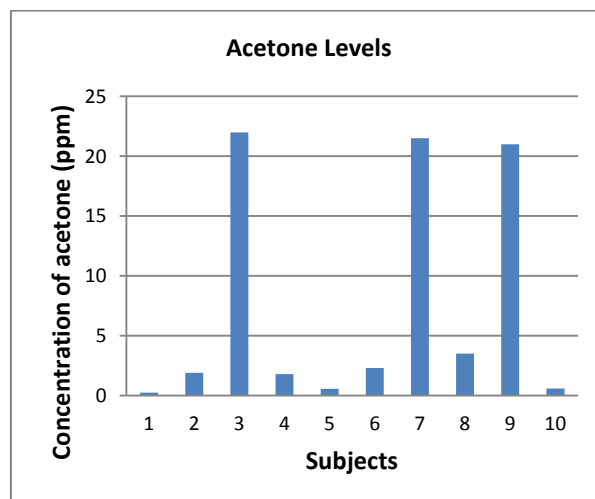


Fig. 2: Distribution of Acetone Concentration Levels of the 10 Samples.

Based on the above-mentioned range of acetone concentrations the SVM classifier is used to distinguish them. The acetone levels are classified into their labels, which are H for healthy breath, T1 for type one diabetes and T2 for type two diabetes. From the threshold range mentioned in the previous section, it can be seen that there is an order in the levels of acetone concentrations, which do not overlap each other, and they conform to the guideline of $T1 > T2 > H$. Here 10 various acetone concentrations are considered for the classification. The input sample vector is taken as these concentrations and the trained samples are the classified data. A set of hyper planes are used here so as to classify the

ten samples into the three different classes. The classifier chooses the hyper planes such that the nearest trained data point has the largest distance so that a good segregation of the data points are achieved.

The scatter plot that distinguishes the acetone levels are represented in the Figure 3. This figure gives the details of the classification of the acetone levels for a diabetic type 1 and type 2 as well as marks the healthy samples of the ten acetone concentrations which are considered in this paper. It is seen that for the type 2 and type one diabetic subjects the levels of acetone in their breath is high compared to the normal breath acetone levels.

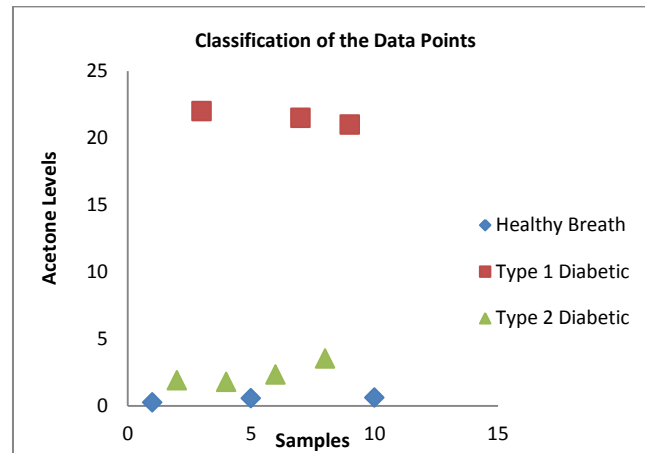


Fig. 3: Scatter Plot That Distinguishes the Trained Data Points into Three Classes Labeled 'Healthy Breath', 'Type 1 Diabetic' and 'Type 2 Diabetic'.

3. Results and discussions

The classified acetone concentrations of the ten subjects are done using the SVM classifier and the Table 2 shows the output.

Table 2: Variation of Acetone Levels

Subject	Concentration of Acetone (ppm)	Healthy	Type 2	Type 1	Labels
Patient 1	0.24	Yes	No	No	H
Patient 2	1.89	No	Yes	No	T2
Patient 3	22	No	No	Yes	T1
Patient 4	1.78	No	Yes	No	T2
Patient 5	0.56	Yes	No	No	H
Patient 6	2.31	No	Yes	No	T2
Patient 7	21.5	No	No	Yes	T1
Patient 8	3.51	No	Yes	No	T2
Patient 9	21	No	No	Yes	T1
Patient 10	0.60	Yes	No	No	H

H- Healthy Breath, T1- Type 1 Diabetic Breath, T2- Type 2 Diabetic Breath.

It is seen that based on the threshold parameters described in the previous section, the classification and the labeling of the data have been performed. It is clearly seen that the acetone concentrations in the breath of a diabetic patient (Type 1 and Type 2) is greater than the normal breath standards.

When the ten trained samples were classified, it was found that a higher level of accuracy can be obtained using an SVM classifier with margin sampling. This approach has given the data a clear margin over which they can be segregated. The ten concentrations were labeled into the three categories in which the samples were classified.

4. Conclusion

This paper investigates the concentration of acetone levels for classification of breath samples in monitoring diabetes. Acetone concentrations were collected and classified for predicting diabetes. The samples were classified using the SVM classifier and were divided into three groups labeled as healthy, type 1 and type 2 diabetes. In this paper, an attempt has been made to refine the existing classification approach (healthy and diabetic) further for a more accurate evaluation (healthy, type 1 and type 2 diabetic).

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