Bayesian Analysis Approach to Diagnose Diabetes Type-2

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Abstract: Diabetes is powerlessness of body to deal with the levels of sugar in the blood. It is being a standout amongst the most persistent infections around the world causes around 3.8 million deaths each year. The primary point of this paper will be to create a methodology (Bayesian Analysis Approach) that after investigation of certain parameters can foresee that whether an individual will be Type-2 diabetic or not. Bayesian Analysis has significant impact in decision making in numerous fields extending from keeping money industry, to travel industry, to correspondence industry, what's more, to automated industry. Bayesian analysis approach has been becoming normal and is being utilized in determining diseases like tumors, hepatitis, lung ailments and many other dreadful diseases. Authors have distinguished 8 parameters that assume a critical part in diabetes and arranged a rich database of preparing information which served as the spine of the forecast calculation. At the point when the parameters of the test information will be sustained to the framework, it envisions & characterizes the test information into one of the two classifications viz diabetic & not diabetic. The execution of Bayesian Analysis in the medicinal judgment framework was discovered to be 95% accurate.

Keywords: Bayesian Analysis, Diabetes, Contingent Probability

1. Introduction

Diabetes, regularly alluded to by specialists as diabetes mellitus, depicts a gathering of metabolic ailments in which the individual has high blood (glucose), either in light of the fact that insulin generation is lacking, or on the grounds that the body's phones don't react legitimately to insulin, or both. Patients with high glucose will regularly encounter polyuria (frequent urination), they will get to be progressively parched (polydipsia) and hungry (polyphagia). In 2013 it was assessed that more than 382 million individuals all through the world had diabetes. [2]

Types of Diabetes

Type 1 Diabetes - the body does not create insulin. More or less 10% of all diabetes cases are Type 1.

Type 2 Diabetes - the body does not create enough insulin for fitting capacity. More or less 90% of all instances of diabetes worldwide are of this sort.

Gestational Diabetes - this sort influences females amid pregnancy. [2]

What is Type 2 Diabetes?

We all need to eat sustenance, it issues us the vitality we have to do our every day exercises. Our bodies separate the sustenance we eat into sugar (glucose), and that sugar goes into our blood. In place for our bodies to utilize the sugar for vitality, it needs to go from the blood to our body cells. [1][2]
Figure 3: This image indicates how the body without diabetes functions, with Type 2 diabetes, the pancreas is making less insulin, and/or the body experiences issues utilizing that insulin.

Figure 4: This image demonstrates how what is occurring in the body with Type 2 diabetes and/or pre-diabetes.

What is Pre-Diabetes?

Pre-diabetes implies that the pancreas is making less insulin than it was in the recent past, and/or that your body is getting to be impervious to insulin (not able to utilize the insulin that is accessible). An individual with pre-diabetes has a higher than typical level of sugar in their blood, yet those levels are not yet sufficiently high for a finding of diabetes (hindered fasting glucose, or disabled glucose resilience).

A conclusion of pre-diabetes may be seen as a "reminder" to start to settle on changes in regular choices so that a determination of diabetes may be evaded. The Important thing to know is that Type 2 diabetes can be counteracted by eating healthy, being physically dynamic most days of the week, and losing even a little measure of weight. [2]

Bayesian Analysis Approach

Bayesian Network:

A Bayesian Network consists of the following:

- A set of variables and a set of directed edges between variables.
- Each variable has a finite set of mutually exclusive states.
- The variables together with directed edges form a directed acyclic graph (DAG).
- To each variable D with parents V1, ..., Vn, there is attached a conditional probability table that is P(D|V1, ..., Vn). [3]

D-Separation

Two distinct variables A and B are d-separated, if for all paths between A and B there is an intermediate variable V, (distinct from A and B) such that, the connection is serial or diverging is instantiated and the connection is converging, and neither V nor any of V’s descendants have received evidence. If A and B are d-separated, then we call them as d-connected. [3][4]

The Chain Rule or Factoring:

We can always write

\[ P(a, b, c, ..., z) = P(a | b, c, ..., z) P(b, c, ..., z) \]

(by definition of joint probability)

Repeatedly applying this idea, we can write

\[ P(a, b, c, ..., z) = P(a | b, c, ..., z) P(b | c, ..., z) P(c | ..., z) P(z) \]

This factorization holds for any ordering of the variables. This is the chain rule for probabilities.

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Methodology

The authors selected the Bayesian analysis approach to diagnose type 2 diabetes in patients. For this analysis a dataset of size 50 has been collected from the Abbasi Shaheed Hospital, Karachi , Pakistan. In this survey, an endeavor is being made to give a review in regards to the applications of Bayes' hypothesis and clinical choice investigation in touching base at a conclusion. With a specific end goal to comprehend demonstrative thinking, it is vital to comprehend the essential numerical dialect of probability and Bayes' hypothesis as connected to clinical medication. [3][4]

Probabilistic reasoning in Clinical Diagnosis:
Probability as connected to clinical indicative thinking may be viewed as a measure of one’s quality of conviction that an occasion will happen and range from 0.0 to 1.0. In factual documentation, probability of an occasion A is composed as P[A].[3][6]

**Law of Total Probability**

The summation rule states that the entirety of probabilities of every single conceivable result of a chance occasion rises to 1.0. On the off chance that there are four conceivable results as K,L,M and N then 


**Joint Probability**

The associative event of any number of occasions will be characterized as joint likelihood of those occasions. In factual documentation, the joint likelihood of two occasions An and B is composed as P[A,B]. [3][5][6]

**Contingent Probability**

The probability that an occasion A happens, given that the occasion B will be known to happen will be characterized as the contingent likelihood of occasion A given occasion B alternately P[A/B]. The relationship in the middle of joint and contingent probabilities is given by the equation P[A,B] = P[A/B] * P[B]. Authors have used eight parameters/factors/symptoms as part of the diagnosis of diabetes.[3][4]

### Table 1: Factors/symptoms of diabetes

<table>
<thead>
<tr>
<th>Symptoms((V))</th>
<th>State of facts</th>
<th>Permitted data values</th>
</tr>
</thead>
</table>
| Age(V\(_1\))    | Age of the Person:  
\[ \begin{align*} 
& \text{Age is divided into three sub parameters, young} 
& \leq 30, \text{middle-age}\ 30 < V_1 \leq 50, \text{old age} > 50. 
\end{align*} \] | Discrete integer values |
| Family History(V\(_2\)) | Either any predecessor or successor is / was suffering type II diabetes. | Yes or No |
| Alcohol/Smoking (V\(_3\)) | Either the person does or does not drinking/Smoking. | Yes or No |
| Weakness(V\(_4\)) | Does a person get weakened in doing a little effort. | Yes or No |
| Frequency of maturation/Urination(V\(_5\)) | Number of times the person passes urine in a day | Discrete integer value. |
| Pancreatic disease (V\(_6\)) | | Yes or No |
| Pregnancy (V\(_7\)) | Either the female subject is pregnant or not | Yes or No |
| Overweight (V\(_8\)) | | Yes or No |

Each symptom or parameter in Table 1 has independent contribution to the prediction of the final result. Mathematically the probability model for a classifier is a conditional model \( p(D/V_1, V_2, ..., V_n) \), over a dependent class variable with a small number of outcomes or classes, conditional on several feature variables \( V_i \) to \( V_n \). Baye’s Theorem can be demonstrated as:

\[
p(D/V_1, V_2, ..., V_n) = \frac{p(D)p(V_1, V_2, ..., V_n)}{p(V_1, V_2, ..., V_n)} = \frac{p(D)p(V_1/D)p(V_2/D, V_1)p(V_3/V_2, V_1, D, V_3) \cdots p(V_n/V_{n-1}, V_2, V_1, D, V_n)}{p(V_1/D, V_2, ..., V_n)}.
\]

Now the “naive” conditional independence assumptions applied as: presuming that individual symptom for diabetes is conditionally independent of every other symptom for diabetes \( V_j \) for \( i \) not equal to \( j \). This means that:

\[
p(V_i/D, V_j) = p(V_i/D)
\]

For \( i \) not equal to \( j \) and so the joint probability can be expressed as:

**Figure 6: Main Symptoms of Diabetes**
\[ p(D, V_1, V_2, \ldots, V_n) = p(D) p(V_1 / D) p(V_2 / D) p(V_n / D) \ldots = p(D) \prod_{i} p(V_i / D) \]

<table>
<thead>
<tr>
<th>( V_i ) (young)</th>
<th>( V_i ) (middle)</th>
<th>( V_i ) (old)</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.0001</td>
<td>0.0015</td>
<td>0.002</td>
</tr>
<tr>
<td>0.002</td>
<td>0.005</td>
<td>0.01</td>
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<tr>
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<td>0.01</td>
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<td>0.0042</td>
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<td>0.001</td>
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<td>0.00252</td>
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</tr>
<tr>
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<td>0.01</td>
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<tr>
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</tr>
<tr>
<td>0.82</td>
<td>0.18</td>
<td>1</td>
</tr>
</tbody>
</table>

3. Conclusion

In the previous decade, there have been colossal advances in the utilization of Bayesian procedure for examination of epidemiologic information, and there are currently numerous useful focal points to the Bayesian approach. Bayesian models can undoubtedly suit in secret variables, for example, a singular's actual illness status in the vicinity of symptomatic slip. The utilization of earlier likelihood circulations speaks to a capable component for fusing data from past studies and for controlling jumbling. Apparatuses are currently accessible that permit disease transmission experts to exploit this capable way to deal with evaluation of presentation sickness relations.

References


