


Cardiovascular Risk Detection Through Big Data Analysis

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ABSTRACT

Cardiovascular diseases are the main cause of mortality in the world. As more people suffer from diabetes and hypertension, the risk of cardiovascular disease (CVD) increases. A sedentary lifestyle, an unhealthy diet, and stressful activities are behaviors that can be changed to prevent CVD. Taking measures to prevent CVD lowers the cost of treatments and reduces mortality. Data-driven plans generate more effective results and can be applied to groups with similar characteristics. Currently, there are several databases that can be used to extract information in real time and improve decision making. This article proposes a methodology for the detection of CVD and a web tool to analyze the data more effectively. The methodology for extracting, describing, and visualizing data from a state-level case study of CVD in Mexico is presented. The data is obtained from the databases of the National Institute of Statistics and Geography (INEGI) and the National Survey of Health and Nutrition (ENSANUT). A k-nearest neighbor (KNN) algorithm is proposed to predict missing data.

KEYWORDS

Big Data, Cardiovascular Disease, Data Science, KNN, Machine Learning, Pool Cohort Equation, Structured Data, Web Tool

INTRODUCTION

According to the World Health Organization (WHO, 2018), ischaemic heart disease and stroke, are the main cause of death in the world; the organization estimates that 15.2 million people died due to those diseases in 2016. The main factors of cardiovascular disease (CVD) are hypertension, dyslipidemia, abnormal glucose levels and diabetes (Anstiss & Passmore, 2020). Despite its high mortality, CVD can be prevented through behavioral changes as physical activity, healthy diet, limiting tobacco exposition, restricting alcohol consumption, and reducing stress (Hooker, 2013). Changing the lifestyle of people is a big challenge due to the increase in sedentary work and the consumption of high calorie foods, so dedicated attention and training for CVD prevention is required (Saeed et al., 2018). Plans and actions to reduce the risk of CVD can be improved by analyzing the data generated every day from the treatment of people with CVD cases, including their origin and evolution.

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Multiple databases are fed with CVD data every day around the world. Centers for Disease Control and Prevention (CDC, 2020), National Survey for Health and Nutrition (ENSANUT, 2020), National Cardiovascular Disease Database (NCVD, 2020), and World Health Organization (WHO, 2020) are some organizations that keep databases available online. Data stored in public databases can be supplemented with patient records, clinical analyzes, diagnoses, and data collected from mobile devices that people use to track their treatments. New correlations between data can be discovered through big data analysis tools. Big Data analysis can help identify common characteristics in people with CVD, and those results could help select actions of greater impact that could be carried out in countries where the lack of resources restricts the amount of clinical analysis that can be performed in the population.

The term Big Data was coined by Gartner Group in 2008 (Emmanuel & Stanier, 2016) for referring to the large amount (volume) of data that is generated continuously (velocity) from several sources (variety) every day (Edward & Sabharwal, 2015). To analyze the collected data, an exploration is performed, followed by a description, and then visualized to interactively explore the content; in this way, users can identify patterns and infer correlations to support decisions (Bikakis, Papastefanatos, & Papaemmanouil, 2019). Good results in the analysis phase can be guaranteed with quality of data. Quality is ensured by evaluating the accuracy, completeness, consistency, distinctness, precision, timeliness and volume of data (Cappiello, Samá, & Vitali, 2018). After data preparation, a set of analysis techniques can be applied to discover relevant information; most used techniques are: pattern matching, classification, clustering, and regression (Mogha, Ahlawat, & Singh, 2018). All the stages involved in big data processing allow the generation of new insights that could be used to take actions for solving the problem under study.

According to Hong et al. (2018) Big Data in healthcare can be divided in four categories: medicine and clinics, public health and behavior, medical experiment, and medical literature. In the first class the data is extracted from electronic health records (Hoffman, 2016), electronic medical records (Setiawan, Utami, Mengko, & Indrayanto, 2014), personal health records (Okore, Bakyarani, & Phil, 2015) and medical images (Tahmassebi et al., 2019). Data of second class is collected from electrocardiograms (Cipresso, Rundo, Conoci, & Parenti, 2019) and vital signs (Mohammad Forkan, Khalil, & Atiquzzaman, 2017). The third class obtains data from molecular biology (Cannataro, 2019), human body samples, and clinical trials (Mayo et al., 2017). Finally, medical literature is obtained from structured knowledge, journal and conference articles (Wang et al., 2018).

This work proposes a methodology for CVD risk detection at state-level making use of a web tool for processing data obtained from public databases. The paper is organized as follows: first, the related work is presented; then, a methodology for the detection of CVD through a big data tool is described; the methodology to extract, describe, and visualize data from a CVD case study at the state level in Mexico is presented below; then, a K-Nearest Neighbor (KNN) algorithm is proposed for the classification of missing data; finally, future work and challenges to be addressed are presented.

BACKGROUND

Alshraideh et al. (2014) developed an algorithm to detect arrhythmias in individuals by identifying 18 attributes of personal information and electrocardiogram (ECG) signals. The system is made up of an Android application that obtains the ECG signal wirelessly and sends the information to a web server where the algorithm is stored. If CVD is detected, an SMS is sent to the user to view their information through a web interface. Roses et al. (2017) conducted a review to identify the effectiveness of positron emission tomography supplemented with computed tomography (PET/CT) for the detection of CVD. PET/CT is used to determine the presence of ischemia in myocardial tissue. They conclude that there is currently a limitation of the indicators, especially in asymptomatic patients, which is why new protocols are required to improve the effectiveness of PET/CT for the detection of CVD. Hernesniemi et al. (2018) created a database to integrate data from hospital records and national registries, disease

phenotype, details of invasive operations, patient clinical information, biometric data physiological, anatomical measurements, and laboratory results obtained from 73,000 individuals treated in TAYS Heart Hospital; the next step will be to analyze the data with a convolutional neural network. Pfohl et al. (2019) developed an adversarial network to consider under-represented minorities in the risk model of atherosclerotic cardiovascular disease (ASCVD) by considering race, gender, and age. Manonmani and Balakrishnan (2020) proposed a semantic annotation process on healthcare data collected from sensors, handheld devices, and datasets to establish relationships between the characteristics identified in the data through fuzzy rule mining. Once the data is annotated, classification algorithms can be used to determine whether or not a patient has a specific disease. Leone et al. (2020) identified that the risk of CVD in patients with systemic lupus erythematosus (SLE) is underestimated using standard risk calculators; For this reason, they propose the use of echocardiography as a non-invasive detection tool to accurately assess the risk of CVD in patients with SLE.

Detecting CVD risk is a process which requires to analyze several factors on patients or healthy persons. The risks assessment can be performed through the identification of risk factors, imaging methods or methods like fibrinogen, homocysteine, and lipoprotein-associated phospholipase 2. According to the European Guidelines on cardiovascular disease prevention in clinical practice (Piepoli et al., 2016), for patients 20 to 79 years of age who are free from clinical atherosclerotic cardiovascular disease (ASCVD), the first step is to assess ASCVD risk factors: Age, gender, race/ethnicity, current smoker (yes/no), any family history of heart attack in first degree relative (parent/sibling/child) (yes/no), diabetes, total and HDL cholesterol, use of lipid lowering medication (yes/no), systolic blood pressure (mmHg), and use of antihypertensive medication. For patients outside this age range, providers should refer to applicable clinical practice guidelines and adult primary prevention guidelines.

The assessment of cardiovascular risk is not necessary for specific subgroups of asymptomatic individuals at unusually high risk, such as those with genetically determined extreme values of traditional risk factors, hypercholesterolemia, hypertension and others diseases like diabetes, chronic kidney disease, obstructive sleep apnoea, and erectile dysfunction (Arnett et al., 2019). These parameters are presented below:

1. Diabetes and age > 60 years;
2. Diabetes with microalbuminuria (>20 mcg/min or urinary albumin:creatinine ratio [UACR] >2.5 mg/mmol for males, >3.5 mg/mmol for females);
3. Moderate or severe chronic kidney disease (CKD) (persistent proteinuria or estimated glomerular filtration rate [eGFR] <45 mL/min/1.73 m²);
4. A previous diagnosis of familial hypercholesterolaemia;
5. Systolic blood pressure (SBP) ≥180 mmHg or diastolic blood pressure (DBP) ≥110 mmHg;
6. Serum total cholesterol (TC) >7.5 mmol/L.

Others methods for diagnosis that are used as a complement to the risk assessment are:

1. **Imaging Studies:** Measurement of carotid intima-media thickness (Ravani et al., 2015) and/or screening for atherosclerotic plaques by carotid artery scanning (Romanens, Sudano, Adams & Schober, 2019), measurement of ankle-brachial index (Bendermacher et al., 2012), computed tomography for coronary calcium (AlJaroudi et al., 2019), and exercise electrocardiography with microalbuminuria (Al-Saffar, Nassir, Mitchell & Philipp, 2015);
2. **Measurement of Inflammatory and Thrombotic Biomarkers:** High-sensitivity C-reactive protein (CRP) and homocysteine (Cheng, Tu, Shen, Zhang & Ji, 2018), fibrinogen (Kaur, Singh, Indu, & Singla, 2012), homocysteine and lipoprotein-associated phospholipase 2 (Lp-PLA₂) (Nicholls, 2018);

3. **Genetic Testing:** Deoxyribonucleic acid (DNA)-based tests for common genetic polymorphisms (Dinter et al., 2019);
4. **Assessment of Psychosocial Risk Factors:** Low socioeconomic status, lack of social support, stress at work and in family life, depression, anxiety, hostility, and the type D personality (Sood & Gidding, 2016).

The application of the methods described above will depend of the level of risk:

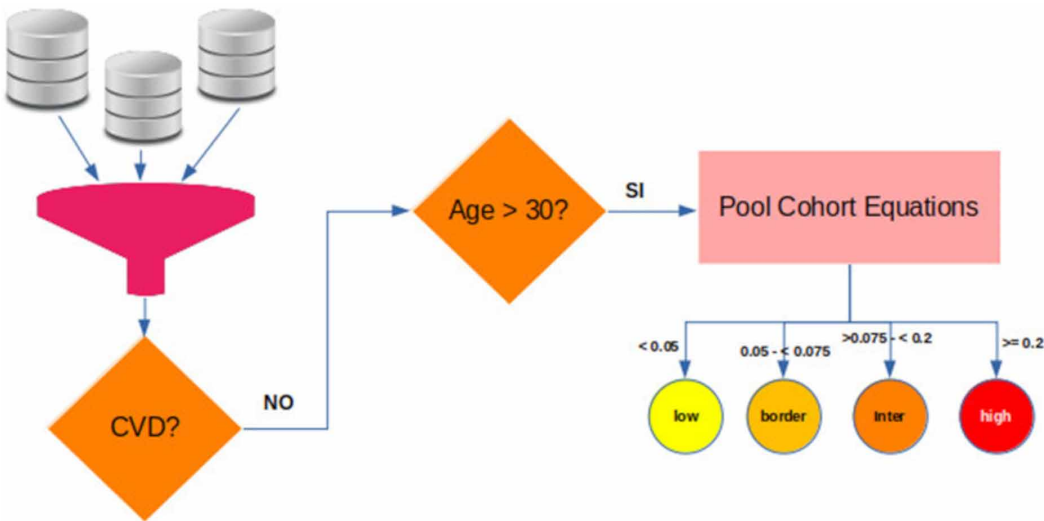
- **Moderate Risk:** High-sensitivity-C-reactive protein (CPR) and homocysteine (Deebukkhum, Pingmuangkaew, Tangvarasittichai & Tangvarasittichai, 2012), measurement of ankle-brachial index (Nishimura et al., 2017), computed tomography for coronary calcium (AlJaroudi et al., 2019), and exercise electrocardiography (Van de Sande, Breuer & Kemps, 2016);
- **High Risk of a Recurrent Acute Atherothrombotic Event:** Lipoprotein-associated phospholipase 2 (Lp-PLA₂) (Rallidis et al., 2012).

The genetic testing is not recommended, the testing do not add significantly to diagnosis, risk prediction, or patient management but the psychosocial risk factors can be assessed through data mining to improve treatment adherence.

METHODOLOGY FOR CARDIOVASCULAR RISK DETECTION (MCRD)

There exist several databases where individual characteristics are stored and can be used to perform and automated analysis in order to identify accurately CVD risk levels. For example, the Dartnet Institute provides a dataset containing records of blood pressure, LDL, Total Cholesterol, HDL, among others, obtained from patients of several institutions in the United States. In the United Kingdom the same data can be obtained from the Consumer Data Research Centre (CDRC). In Europe data is collected by the European Genome-phenome Archive and the European Society of Cardiology. Others datasets can be obtained from diverse institutions around the world, which can be used for identifying CVD risks levels by regions following the next methodology (Figure 1):

Figure 1. CVD detection from big data



1. **Datasets Merging:** Collect all data available which include most factors as possible of those described above. A more accurate assessment is obtained when multiple risk factors are analyzed. More data available increases accuracy;
2. **Identify People With Established CVD and Exclude:** When data is collected from several sources, data about persons which are currently under CVD treatment could be included. So, exclude them and continue working with the remaining registries;
3. **Age classification:** Separate men and women older than 30. Include those registries in the high level risk and continue working with the remaining registries;
4. **Risk Assessment and Risk Stratification:** For classifying the remaining registries apply the Pooled Cohort Equations (Yadlowsky et al., 2018) and classify according to the Table 1.

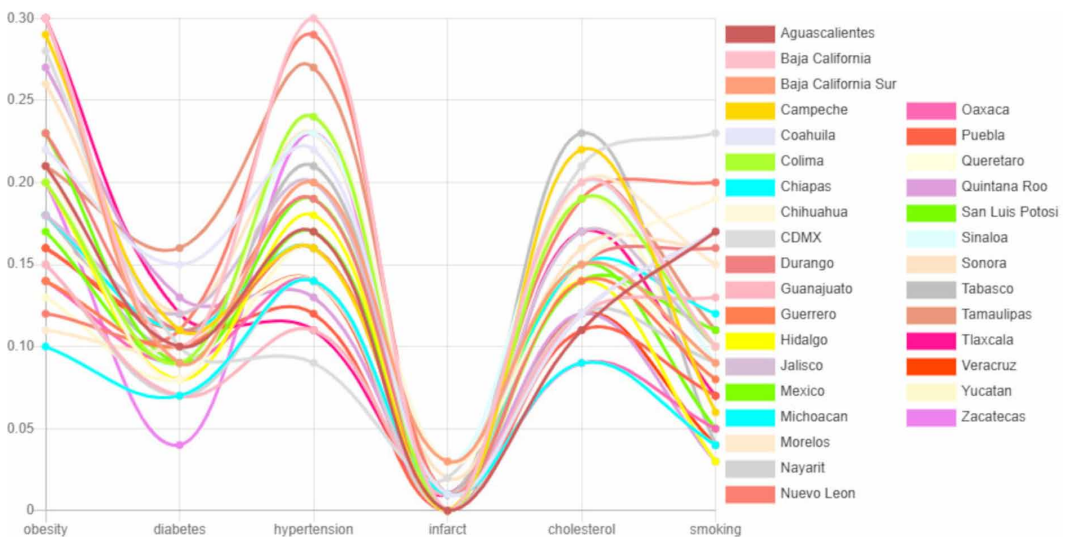
Table 1. Absolute cardiovascular risk (Arnett et al., 2019)

Categories	Score
Low Risk	< 5%
Borderline Risk	5% - < 7.5%
Intermediate Risk	>= 7.5% - < 20%
High risk	>=20%

RESULTS AT STATE LEVEL IN MEXICO

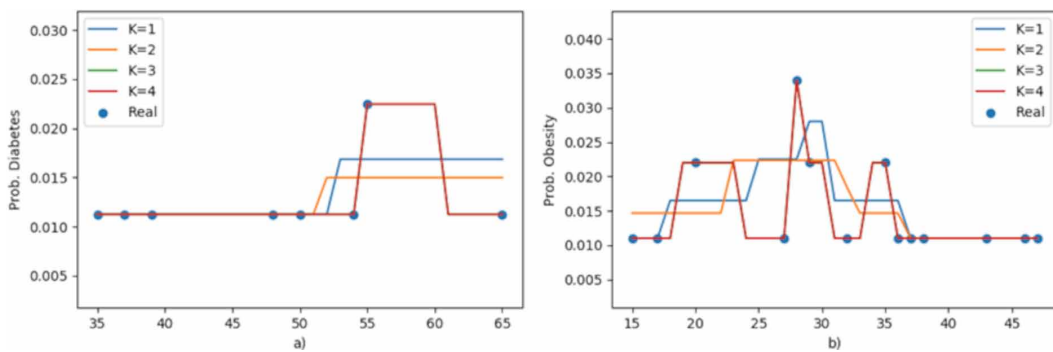
A web tool was designed to evaluate the proposed methodology. The first version of the tool allows the analysis of information at the state level. First, data is collected from public databases of ENSANUT and INEGI in Mexico; then, the variables to consider (obesity, diabetes, hypertension, infarct, cholesterol, and smoking) are extracted from the data (Figure 2).

Figure 2. Data collected from ENSANUT and INEGI in 2018



Once the variables are extracted from the databases, a classification algorithm (k-closest neighbors) is trained to fill in the missing values from the databases for each variable and each state. Figure 3a illustrates the probability of diabetes in the state of Aguascalientes, Mexico; the data available in the databases is represented by blue dots and the predictions obtained after training the algorithm with 1, 2, 3 and 4 neighbors are illustrated with different colored lines. It is possible to observe that good results are obtained with 3 neighbors. The precision of the trained set is 0.24. For better accuracy more data is required. These data can be complemented with the new databases published by the National Institutes in the years in which the population evaluation is carried out. Figure 3b illustrates the results on the obesity variable obtaining a precision of 0.30.

Figure 3. a) KNN Prediction Algorithm for Diabetes in Aguascalientes, Mexico; b) KNN Prediction Algorithm for Obesity in Aguascalientes, Mexico



After completing all the data with the KNN algorithm for each state, the grouping cohort equation (Arnett et al., 2019) is applied to obtain the risk level category. Percentage of population older than 30 years (Figure 4), with obesity (Figure 5), and diabetes (Figure 6) are visualized in an interactive map.

CONCLUSION

The availability of more data sources opens up the possibility of finding patterns that improve decision-making; however, analyzing all the data generated is a challenging activity. Big Data tools help process large amounts of data and discover patterns that can be visualized through interactive interfaces that allow the analysis of information. In this work, a state-level analysis of CVD in Mexico was carried out through a web tool following the proposed methodology. This information can be used to establish policies that encourage people to change their habits to reduce the risk of CVD. As future work, the application will be improved to allow the analysis of information at the municipal level and to visualize the behavior of several years in real time. To make better predictions more data have to be collected and the machine learning algorithm improved.

Figure 4. Population older than 30 years

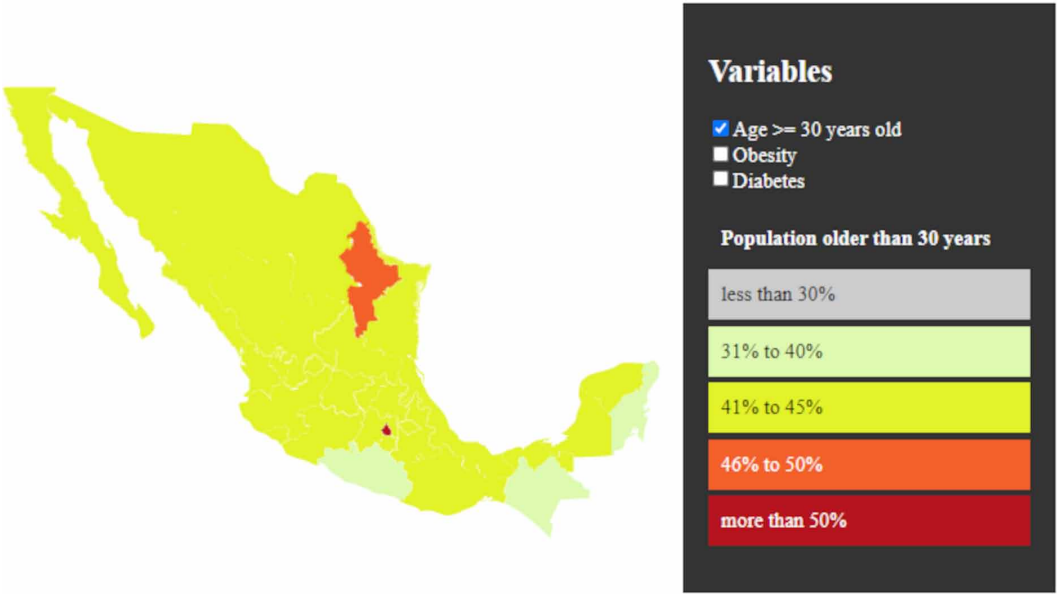


Figure 5. Percentage of the sample with obesity

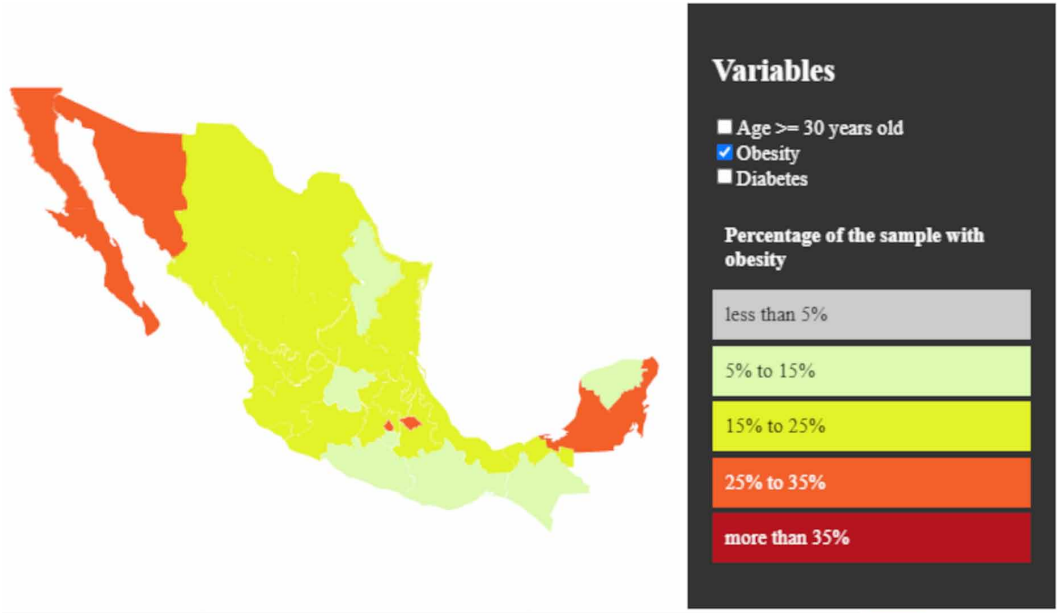
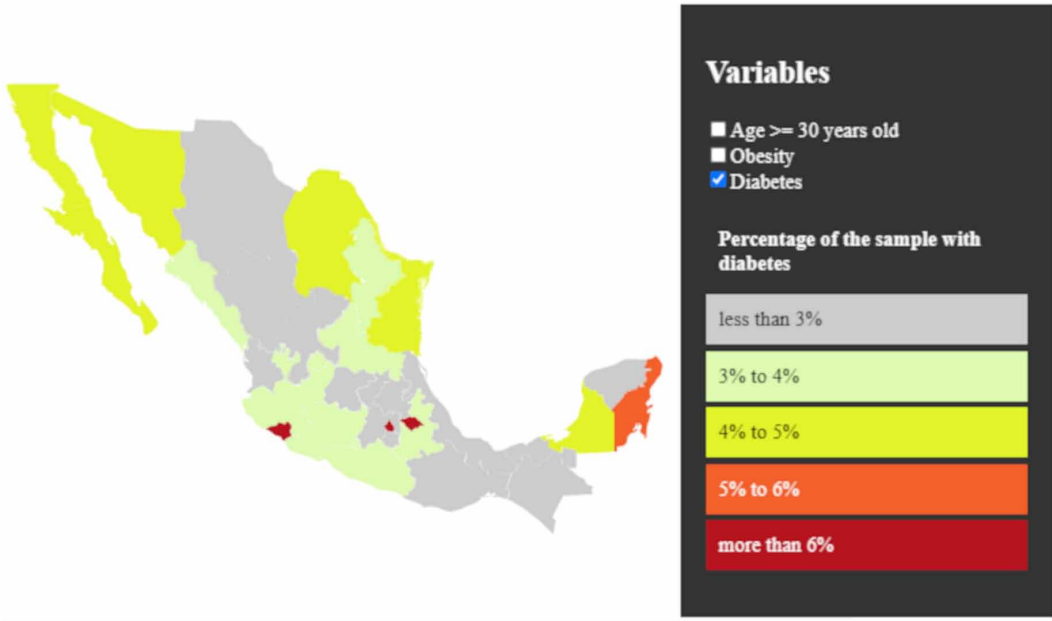


Figure 6. Percentage of the sample with diabetes



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