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## Detection of variables for the diagnosis of overweight and obesity in young Chileans using machine learning techniques.

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### Abstract

Overweight and obesity are considered epidemic problems. The number of factors involved in developing extra body fat makes harder the detection of this problem. Therefore, among the several variables and their levels presented in overweight and obese people, there is a need to improve the classification of people with these conditions. To this aim, in this paper, we conducted a variable analysis from biochemical and lipid profiles in young Chileans with normal weight, overweight, and obesity using machine learning techniques. XGBoost library was selected as the classifier. 21 variables (13 from biochemical and 8 from lipid profiles) were chosen as features. 100 iterations were conducted, and an 80% cross-validation was obtained. The variables with greater relevance in the classification task were total cholesterol, glycemia, LDH enzyme, bilirubin, and VLDL cholesterol. All of these, except bilirubin, are consistent with previous research in which these features have been used to assess risk factors for developing overweight or obesity. Then, further research must include a deep study regarding bilirubin's influence over these conditions.

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## 1. Introduction

Overweight and obesity represent a public health concern worldwide. According to the WHO (World Health Organization), overweight and obese people experience several risks to their health because of the extra body fat [1]. The prevalence of these conditions in children and youth has risen special attention from government and public organizations [2]. For instance, the available data show that a third of US children and adolescents are estimated to be overweight or obese [3]. Then, most developed countries consider overweight and obesity in youth as epidemic problems [4]. Therefore, to decrease the incidence of these phenomena, public policies have been focused on the promotion of acquiring healthier lifestyles including physical activities at an early age [5].

In Latin America, overweight and obesity rates have an increasing behavior in recent years [6; 7; 8]. In this zone, some factors influence childhood overweight and obesity, such as socio-economic conditions and even genetics [7; 8]. Particularly, Chile is in the 2<sup>nd</sup> place among the Organization for Economic Cooperation and Development (OECD) members in obesity prevalence [9]. The country has followed the same pattern as global performance, increasing its overweight and obesity rates in the last few years [10]. Despite the latent need to prioritize this situation, there are limited interventions for obesity prevention in Latin America. However, there is a consensus that these actions need to take into consideration multidimensional factors. Overweight and obesity used to be defined based on the body mass index (BMI), using boundaries of 25 kg/m<sup>2</sup> and 30 kg/m<sup>2</sup> as cut-off values, respectively. However, there are other indicators that, together with BMI, have improved the detection of overweight and obese people.

One of these is the biochemical and lipid profile analysis. The relationship between lipid profiles and overweight or obesity has been proven before [11; 12]. Techniques such as bivariate and multivariate analyses have been used to study this association [13]. But there is evidence arguing that traditional approaches are limited in identifying predictors efficiently [14].

Thus, in this article, we try to fill this gap by finding the critical variables from biochemical and lipid profiles to classify people with normal weight and altered BMI (overweight and obesity) in young people using machine learning (ML) techniques. Machine learning has been used extensively in the healthcare sector [15]. Nevertheless, its use for detecting obesity and overweight in youth is under-explored.

The remainder of the paper is organized as follows. A brief literature review is included in Section 2. Then, we describe the machine learning tool and data selected for conducting our research in Section 3. This is followed by the main results and implications in Section 4. Finally, the last section includes conclusions and further research opportunities.

## 2. Literature Review

The detection of overweight and obesity has been studied before in literature. Some methods, such as anthropometric assessment has been used to determine nutritional screening indirectly [16]. Moreover, machine learning models, including linear regression, Support Vector Machine (SVM), Random Forest (RF), and Artificial Neural Networks (ANN) have been used for this purpose. Among these models for predicting body weight, ANN and RF have shown better performance [17]. Also, machine learning-based clinical decision support systems have been developed to predict people at risk of some diseases where overweight and obesity have a critical influence [18].

Random Forest (RF), with other techniques such as Decision Trees (DT) and XG Boost, have been studied to predict heart diseases [19]. Using XG Boost has improved the models' performance and accuracy [20]. eXtreme Gradient Boosting (XG Boost) was first introduced as a scalable end-to-end tree-boosting system [21]. XG Boost has shown excellent performance in improving the diagnosis of primary lesions related to metastatic tumors [22], for the prediction of breast cancer [23; 24], and as a classifier for stunting among children in Zambia [25].

Despite the advantages of the XG Boost technique, their use for predicting body weight is negligible in the literature. Recently, Santisteban Quiroz [26] studied the performance of this technique in classifying the incidence of obesity based on dietary habits. Therefore, this paper presents an application of the XG Boost method to predict normal weight and altered BMI (overweight and obesity).

### 3. Method

#### 3.1. Data.

A matrix with 21 variables was used for the lipid (L) and biochemical (B) profiles of 40 Chilean university students with ages 20 – 30 years. The sample selection criteria were their BMI. Also, people who follow any medical treatment for losing weight or athletes were excluded. Table 1 includes the chosen set of variables.

Table 1. Set of variables for lipid and biochemical profiles.

| Feature | Variable                                   | Profile type |
|---------|--|--------------|
| F0      | Basal insulin                              | Biochemical  |
| F1      | Total Cholesterol                          | Lipid        |
| F2      | Total proteins                             | Biochemical  |
| F3      | Albumin                                    | Biochemical  |
| F4      | Uric acid                                  | Biochemical  |
| F5      | Bilirubin                                  | Biochemical  |
| F6      | Phosphorus                                 | Biochemical  |
| F7      | Calcium                                    | Biochemical  |
| F8      | Urea nitrogen                              | Biochemical  |
| F9      | Glycemia                                   | Biochemical  |
| F10     | Alkaline phosphatase                       | Biochemical  |
| F11     | GOT transaminase                           | Biochemical  |
| F12     | LDH U/L (Lactate dehydrogenase)            | Biochemical  |
| F13     | Alpha 1                                    | Biochemical  |
| F14     | HDL (high-density lipoprotein) Cholesterol | Lipid        |
| F15     | LDL (low-density lipoprotein) Cholesterol  | Lipid        |
| F16     | VLDL (very-low-density lipoprotein)        | Lipid        |
| F17     | Triglycerides                              | Lipid        |
| F18     | Total cholesterol/HDL                      | Lipid        |
| F19     | LF/HF                                      | Lipid        |
| F20     | SD1  | Lipid        |

#### 3.2. Classification procedure and validation:

XGBoost was selected as the classifier tool. XGBoost is more accurate than other machine-learning techniques in medical classification [27].

The rest of the procedure was structured according to the phases included in Fig. 1.

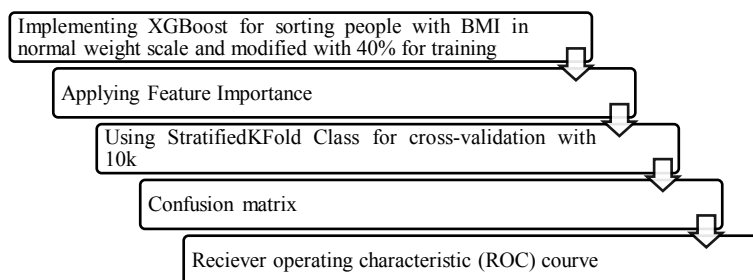


Fig. 1. Classifier implementation procedure.

### 4. Results

To conduct the analysis, Spyder 4.1.4 – Scientific Python and the toolkit of sci-kit-learn with XGBoost were used in an Intel® Core i5, 8GB RAM, Macbook Pro 13. After applying XGBoost a cross-validation of 80% was obtained.

In figure 2 (a) the best iteration is included with an 87.5% ROC curve and an area under the curve (AUC) of 0.818. Also, figure 2 (b) shows the confusion matrix for the best iteration where the classifier made two false negatives (type

II error), which means two observations where the actual classification was positive, and the predicted classification was negative.

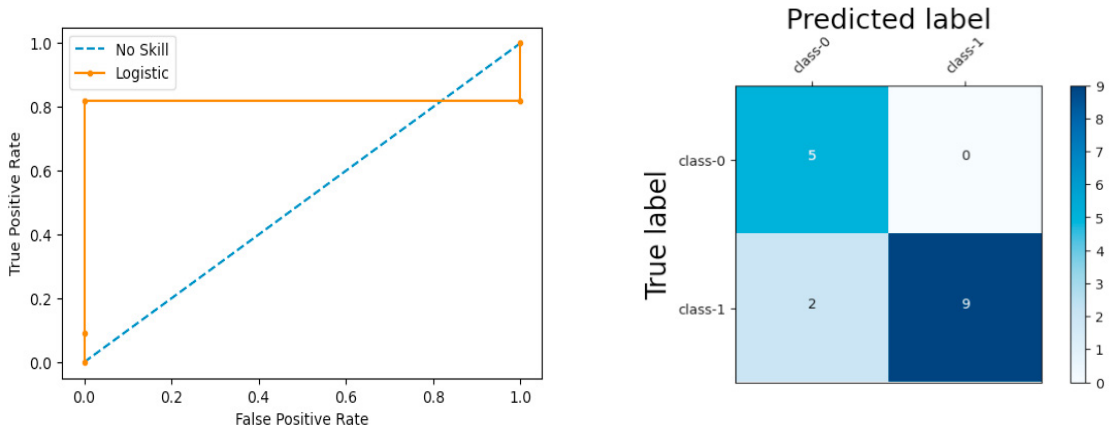


Fig. 2. (a) ROC for the best iteration. (b) Confusion matrix

Table 2. Set of sorted variables according to relevance after 100 iterations.

| Feature   | F1 | F9 | F12 | F5 | F16 | F18 | F0 | F4 | F6 | F20 | F10 | F14 | F3 | F15 | F19 | F2 | F7 | F17 | F11 | F8 |
|-----------|----|----|-----|----|-----|-----|----|----|----|-----|-----|-----|----|-----|-----|----|----|-----|-----|----|
| Relevance | 61 | 47 | 37  | 33 | 32  | 29  | 26 | 25 | 23 | 22  | 17  | 16  | 15 | 15  | 14  | 12 | 10 | 10  | 8   | 4  |

Table 2 shows that the variables with greater impact on the classification process were: Total cholesterol (F1), glycemia (F9), LDH (F12), bilirubin (F5), and VLDL (F16). The priority level of each variable is congruent with previous research in which these variables have been used before to model overweight and obesity patterns.

First, the total cholesterol relevance found in our study agreed with the suggestions issued by the AHA (American Heart Association). The AHA recommends doing a total cholesterol assessment for cardiovascular disease prevention, where obesity plays a critical role [28]. However, using total cholesterol to identify people with obesity is still controversial since this variable considers as much HDL as LDL [29].

Moreover, our results suggest that glycemia (F9) has second place in importance. This is a common variable used in obesity studies due to its relationship with energy imbalance and comorbidities such as diabetes and insulin resistance [30]. For its part, LDH enzyme levels change in presence of obesity [31]. Therefore, it explains its 3<sup>rd</sup> place in importance in our analysis.

The last two variables: bilirubin and VLDL, are key indicators for overweight and/or obesity measurement. Besides, bilirubin has an inverse relationship with overweight and obesity [32]. It means that the higher the body weight, the lower the bilirubin level, and vice versa. Finally, VLDL has been set that contributes 55% to the total content of triglycerides in the blood circulation process [33]. Hence, its level influences weight measurements.

## 5. Conclusions

This paper has presented an application of the XGBoost tool as a machine-learning technique for youths' classification in two categories: normal weight and altered BMI (overweight and obesity). For this purpose, a set of 40 Chilean university students was selected, and 21 variables were chosen from lipid and biochemical profile tests. The classifier showed good performance with an 80% cross-validation and a confusion matrix suggesting just two false negatives.

The main variables affecting the presence of overweight were: Total cholesterol, glycemia, LDH (Lactate dehydrogenase), bilirubin, and VLDL (very low-density lipoprotein). Then, three variables from the biochemical profile and two from the lipid profile can be used to classify young people in the above-mentioned categories.

Four of these five variables are congruent with previous research. Therefore, further research opportunities include increasing the sample size and being focused on studying bilirubin levels for regulating body weight since the evidence related is limited.

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